MECHANISMS AND NOVEL THERAPIES IN GRAVES' ORBITOPATHY: CURRENT UPDATE

EDITED BY: Kelvin Kam-Lung Chong, Ilaria Muller, Huifang Zhou and

Marian Elizabeth Ludgate

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MECHANISMS AND NOVEL THERAPIES IN GRAVES' ORBITOPATHY: CURRENT UPDATE

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Editorial: Mechanisms and Novel Therapies in Graves' Orbitopathy: **Current Update**

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Graves' orbitopathy (GO) is the main extra-thyroidal manifestation in patients with Graves' disease (GD), with a prevalence of up to 50%, although in most cases GO is mild (1). GO has a variety of clinical manifestations including eyelid retraction, exophthalmos, restrictive strabismus, exposure keratitis, and dysthyroid optic neuropathy. It is the commonest orbital disease causing blindness, orbital deformity and visual disability and exerts a profound negative impact on patients' quality of life (2).

This Research Topic provides a timely update on different aspects of GO, in 20 excellent contributions, ranging from epidemiology, pathogenesis, disease evaluation, to comments on existing therapies and novel treatment strategies. Bartalena et al. reviewed GO incidence and prevalence, then highlighted risk factors such as age and smoking, and how they might be regulated or avoided to optimize clinical management. Clinically, hyperthyroidism and orbitopathy often develop simultaneously or within a few months of each other (2). Both thyroid epithelia and orbital fibroblasts (OF) express the thyrotropin receptor (TSHR), suggesting that these two conditions may evolve from common underlying systemic processes (1). Uncontrolled hyperthyroidism, which is caused by autoantibodies (TRAB) to the TSHR in GD, is a major GO risk factor with high TRAB levels long known to correlate with GO incidence and severity (3). In this context, George et al. provided a comprehensive overview introducing thyroid-stimulating antibody (TSAb) as a biomarker for GO whilst Kelada et al. investigated polyautoimmunity as a risk factor of GO activity and severity from their retrospective cohort of 267 GO patients.

The pathogenesis of GO is complex and not fully understood although some consensus has emerged in recent years (4). Most would regard the OFs as target cells contributing to the tissue remodeling which leads to expansion of the orbital contents. Draman et al. summarized OF signaling cascades and how they affect two of these processes, adipogenesis and hyaluronan production. To further our understanding of OFs, Virakul et al. compared the proteome of DNA methylation of OFs from GO patients and healthy controls. They found over-expression of genes implicated in inflammation and proliferation, together with subtle differences between active and inactive GO profiles. A similar microarray approach was applied to the GO lacrimal gland by Tu et al.

GO is an autoimmune disease and orbital inflammation is initiated by the loss of self-tolerance to the putative, shared antigens between thyroid glands and orbits, notably the TSHR (5). Antigen-presenting cells (APC) recognize and present TSHR to T helper (Th) cells for unknown reasons, and the existence of a soluble TSHR may be implicated (6). Upon antigen activation, T cells interact physically with APCs and facilitate B cell proliferation and differentiation, ultimately leading to autoantibody production. It is widely accepted that both cellular and humoral immunities contribute to the GO pathogenesis (3, 7). Some studies have suggested the importance of Th1 cells in the early active phase and Th2 cells in the chronic fibrotic phase of GO (7). An increased number of Th17 cells have been detected at the site of orbital connective tissues (8, 9), leading to the concept of an imbalance in effector and regulatory T cells in GO autoimmunity. Fang et al. comprehensively summarized the current knowledge on T cell immunity in GO: Th1 (cytotoxic leaning), Th2 (antibody leaning) and an emerging role of Th17 (fibrotic leaning) cell subsets, in the context of key pathogenic processes such as adipogenesis and fibrosis. Hypothesizing that auto-immune responses can be locally triggered and aggravated, Lu et al. characterized the T cell repertoire infiltrating paranasal sinus mucosae in GO patients and reported increased proinflammatory effector T cells but reduced regulatory T cells.

Despite advances in understanding, the current treatment for GO is not satisfactory and long-term deformities and disabilities often persist. The problem is exacerbated by the fact that GO presentation can be heterogenous, as highlighted by an overview of asymmetric GO by Panagiotou and Perros, focusing on its clinical relevance and possible mechanisms. The Amsterdam declaration recommends that GO patients should be managed in joint endocrinology-ophthalmology clinics (10). An interesting study from Farag et al. described a 'real world' snapshot of the multidisciplinary management of a multi-ethnic GO population prior to the introduction of established GO standards.

Several existing therapies are currently used in the management of GO. In light of the inflammatory and immunological processes in operation, the first-line treatment is the administration of intravenous glucocorticoids (11). Naselli et al. reported that GO patients with high levels of low-density lipoproteins cholesterol were likely to respond poorly to glucocorticoids. In mild GO, selenium was shown to be beneficial, via its anti-oxidant and immunomodulatory effects, as reviewed by Lanzolla et al. The effects of anti-B cell therapies are somewhat controversial but indicate an effective role of rituximab for early active moderate-to-severe GO (12, 13). Campi et al. performed a post-hoc analysis on the efficacy of

rituximab in GO patients, which can be helpful in the decision-making process. In addition, Zhang et al. described the salvaging benefits of rituximab in 2 patients refractory to glucocorticoids who subsequently underwent orbital decompression.

In order to develop novel effective target therapies for GO, it is essential to clarify the key pathological mechanisms, both immune and non-immune related, occurring within the orbit. The concept of circulating bone marrow derived CD34⁺ fibrocytes in the pathogenesis of GO is original and impactful. These TSHR-expressing progenitor cells migrate from the peripheral blood into orbital connective tissues and transit into CD34⁺ OFs that upon stimulation, undergo adipocytic and myofibroblastic differentiation (14). TSHR signaling in fibrocytes and OFs are partially dependent on the insulin-like growth factor 1 receptor (IGF-1R), another putative autoantigen that has received increasing attention in the past few years (15). Years of in vitro research were recently translated into a phase 2 and then a phase 3 randomized multicenter trials which consistently demonstrated that GO patients treated with teprotumumab, a fully human monoclonal IGF-1R inhibitor, were significantly more likely to experience a meaningful improvement in proptosis compared with patients treated with placebo (16, 17). Smith, one of the main researchers involved, reviewed the encouraging findings of teprotumumab as the first targeted therapy for GO. The cross-talk between TSHR and IGF-1R mediated by PKA/PI3K-FOXO signaling highlights a feasible therapeutic strategy to attenuate signaling initiated at either receptor, thereby relieving GO processes (4, 18).

Traditional immunosuppressive agents such as mycophenolate, azathioprine, and cyclosporin mainly inhibit the activation and proliferation of T cells (4). To date, no therapy targeting a particular T cell subset has been reported. Fortunately, blocking T cell related cytokines (e.g. IL-6) such as tocilizumab shows promising results in GO (19). Fallahi et al. from Antonelli's group shared their thoughts on cytokine-based therapy in GD and GO.

Previous studies reported changes in T cell subsets during the transition from hyperthyroidism to euthyroidism and from active to inactive GO (20, 21). Thus, whilst GD and GO shared similar antibody-mediated immune attack, as highlighted by the improvement of these conditions in a patient with thyroid cancer treated with a TSHR-blocking monoclonal antibody (22), cell-mediated immunity is believed to play a central role in GO pathogenesis. A phase 1 multicenter trial revealed that ATX-GD-59, a mixture of two TSHR-derived peptides binding with HLA-DR on dendritic cells, improved free thyroid hormone levels in 70% (7/10 responders) of previously untreated mild to moderate Graves' hyperthyroidism (23).

In vitro studies have facilitated identification of other novel treatments: Lanzolla et al. reviewed the use of antioxidant agents in mild GO; Wei et al. proposed simvastatin and ROCK inhibitors for orbital fibrosis whilst Lee et al. demonstrated potential for proprotein convertase as therapeutic target and biomarker.

The field of GO research has benefitted from the development of a robust TSHR-induced *in vivo* model of GO (24, 25).

Comparison of the gut microbiota composition between mice housed in 2 centers revealed significant differences which may account for heterogeneity in the induced immune responses. When diseased and control mice were compared, disease-associated taxonomies were identified e.g. *Firmicutes* positively correlated with orbital adipogenesis (26). The same researchers modified the gut microbiota using antibiotic, probiotic and human fecal material transfer (hFMT). Experimental GO was exacerbated by hFMT and ameliorated by antibiotic treatment confirming a role for the gut microbiome in GD/GO (27). Several groups have investigated whether the same applies in human GD/GO as reviewed by Wang et al., who also provide a comprehensive summary of epigenetic and other factors implicated in the GO disease process.

In conclusion, this Research Topic provides encouraging updates made in understanding the complex interactions between genetic background and environmental factors such as the gut microbiome. It reviews the efficacies and short-comings of emerging therapies (e.g. teprotumumab and potential hearing problems) which have evolved from years of translational

research to date. It illustrates an unlimited potential of nonsurgical treatments, many of which target fundamental disease processes such as ligand-receptor binding and T cell subset imbalance underpinning GO development, and will ultimately benefit patient care in the near future.

AUTHOR CONTRIBUTIONS

SF and HZ wrote the paper. ML, IM, and KC revised the paper. ML was the senior author of the paper. All authors contributed to the article and approved the submitted version.

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Use of Rituximab After Orbital Decompression Surgery in Two Grave's Ophthalmopathy Patients Progressing to Optic Neuropathy

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Background: While orbital decompression can alleviate optic nerve compression and prevent further vision loss in dysthyroid optic neuropathy (DON), it cannot relieve inflammatory symptoms. Very high doses of intravenous glucocorticoids (GCs) are the first-line therapy for DON; however, the effective rate is only 40% and might be much lower in patients who fail high-dose GC pulse therapy and progressed to DON. The results of two case series studies indicated that rituximab treatment had a much better curative effect compared to very high doses of intravenous GCs, but some patients required urgent orbital decompression after rituximab injection because rituximab might lead to the release of cytokines, aggravated intraorbital edema, and further vision loss.

Methods: We retrospectively studied the therapeutic process of two Grave's ophthalmopathy (GO) patients complicated with DON who failed high-dose GC pulse therapy and underwent orbital decompression. Both patients received single-dose (500 mg) rituximab treatment.

Results: During more than 2 years of follow-up, rituximab treatment exhibited significant improvement in inflammatory symptoms, as manifested by a substantial decrease in Clinical Activity Score (CAS); meanwhile, the vision of both patients improved significantly and their diplopia was relieved.

Conclusions: The results of this study were consistent with those of two previous case series studies indicating the significant and lasting effect of rituximab treatment on DON, especially for patients with GC resistance or recurrence after GC therapy. Orbital decompression before rituximab treatment might reduce the incidence of rapid vision loss and urgent orbital decompression surgery caused by aggravated orbital edema after rituximab injection; however, the necessity for preventive decompression surgery requires further study.

Keywords: Grave's disease, Grave's ophthalmopathy, Dysthyroid optic neuropathy, orbital decompression, rituximab

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INTRODUCTION

Grave's disease (GD) is an autoimmune disease involving the thyroid, skin, and orbit, with an incidence in the adult population of 1–2% (1). Grave's ophthalmopathy (GO) is the most frequent extrathyroidal manifestation of GD, occurring in up to 50% of GD patients throughout the course of the disease (2). GO is generally self-limiting, with the signs and symptoms improving naturally or following thyrotoxicosis control, smoking cessation, and local treatment (3). However, for some patients, the signs and symptoms persist or aggravate gradually, which require specific treatment besides smoking cessation and anti-thyroid drug administration (4).

The pathogenesis of GO is incompletely understood, although immunological cross-reactivity between the thyroid and orbital antigens may play a key role, and disorders of inflammatory cytokines, thyrotropin receptor autoantibodies, and immunoglobulins targeting the insulin-like growth factor 1 receptor may be correlated with GO (2, 5). Hence, the current internal medicine treatments for GO mostly target immunological disorders, especially for moderate-to-severe and sight-threatening GO.

High-dose intravenous glucocorticoid (GC) pulse treatment is the first-line treatment for moderate-to-severe and active GO, with a 4.5 g cumulative dose of methylprednisolone divided into 12 weekly intravenous injections recommended. A higher dose is acceptable for severe forms; however, the cumulative doses should not exceed 8.0 g (4, 6). The response rate of high-dose intravenous GCs pulse treatment is 70–80% (4, 6, 7); however, in clinical practice, many patients responding well to intravenous GCs may re-develop active disease after therapy completion. For patients with a recurrence or poor response to intravenous GCs, shared decision-making with patients to select an appropriate second-line treatment is recommended (6). A second course of intravenous GCs, orbital radiotherapy, combination use of cyclosporine and oral GCs, and rituximab are the most common second-line treatments (6).

As an alternative second-line treatment for GO, rituximab has been proven to be effective and safe in patients with GO who fail high-dose intravenous GC pulse treatment and shows potential to become a first-line treatment (6). For GO patients progressing to DON, non-control studies have also shown that rituximab was effective in relieving DON; however, a small number of patients in the studies required urgent orbital decompression surgery after receiving rituximab injection as rituximab may have aggravated intraorbital edema (8–10). This finding is consistent with the recommendation in the 2016 European Group of Graves' Orbitopathy Guidelines for the Management of GO that rituximab should not be used in patients with impending DON (6). Thus, the use of rituximab in patients with GO complicated with DON requires further clinical studies.

This study retrospectively analyzed the therapeutic processes of two GO patients who failed high-dose intravenous GC pulse treatment and progressed to DON. Although a very high dose of

Abbreviations: CAS, Clinical Activity Score; DON, dysthyroid optic neuropathy; GC, glucocorticoid; GD, Grave's disease; GO, Grave's ophthalmopathy.

intravenous GCs is the first-line treatment for DON, patient 1 was not sensitive to GCs and the cumulative doses of GCs exceeded 8 g. Two patients underwent orbital decompression surgery first to avoid further vision loss. However, there was no significant improvement, especially the clinical activity score (CAS) and diplopia. Moreover, the recovery of vision was not ideal. Thus, rituximab treatment was initiated. During 2 years of follow-up after rituximab treatment, both patients achieved stable and significant remission.

CASE DESCRIPTIONS

Both patients were initially diagnosed with moderate-to-severe and active GO and received high-dose intravenous GC pulse treatment. The patients also received basic treatments including selenium supplementation and artificial tears. Orbital decompression surgery and sequential rituximab were initiated when the patients suffered from DON and resisted to GCs or the total methylprednisolone dose exceeded 8g. All patients received a single dose of 500 mg rituximab as a slow intravenous infusion, with 5 mg of dexamethasone to prevent allergic reactions.

Case 1: A 54-year-old man (non-smoker) was diagnosed with GD and GO in December 2017 at a local hospital, where he received methimazole treatment only. Three months later, he was transferred to Tongji Hospital, Wuhan, China, for further treatment due to rapidly declining vision (OD 0.3, OS 0.08), eyeball movement disorder, and inability to close his eyelid. The patient was then diagnosed with DON, with a CAS of 7/7. Two days after transfer to this hospital, we initiated high-dose intravenous GC pulse treatment (0.5 g methylprednisolone seven times) and performed bilateral orbital decompression and eyelid margin suture surgery as ophthalmologic examination showed persisting DON with no significant improvement in vision. The orbital decompression surgery was balanced decompression of the inner and outer orbital walls combined with lipectomy, part of the bone in both medial and lateral orbital walls and about 3ml of adipose tissue were removed in the surgery. After the above treatment, the patient's CAS decreased to 5/7, and ophthalmologic examination showed normal intraocular tension; however, the eyeball movement disorder remained and his vision did not improve (OD 0.1, OS 0.1). The patient then left the hospital and received oral GC treatment at home (methylprednisolone 40 mg/d for 1 week and 36 mg/d for another week). Two weeks later, the patient returned to the hospital. Ophthalmologic examination showed a CAS of 6/7; therefore, we administered an eighth intravenous GC treatment (0.5 g methylprednisolone). His vision improved 7 days after the intravenous GC treatment (OD 0.5, OS 0.6), but the CAS was still 6/7, and the eyeball movement disorder and inability to close the eyelids did not improve. Finally, we decided to administer rituximab treatment (a single dose of 500 mg) and oral GC treatment simultaneously.

Case 2: A 58-year-old man (smoker, quit smoking 10 years ago) was diagnosed with hyperthyroidism and moderate-to-severe GO in March 2017 at the local hospital, where he received methimazole and intravenous GC treatment (11

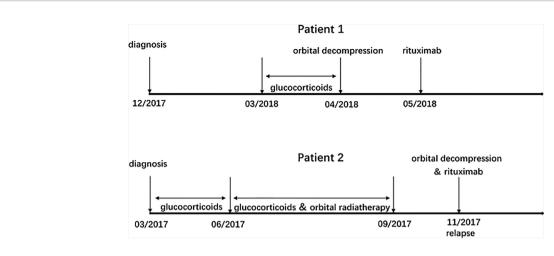


FIGURE 1 | Treatment process: corticosteroids, orbital decompression, and rituximab for patient 1 and patient 2.

weekly intravenous injections of 0.5 g methylprednisolone), without significant improvement. The patient was then transferred to Tongji Hospital, Wuhan, China. Ophthalmologic examination showed eyeball movement disorder; both eyes had 1.0 vision, and his CAS was 5/7. We administered intravenous GC treatment (12 weekly intravenous injections of 0.25 g methylprednisolone) combined with right orbital radiotherapy (2 Gy daily for 10 days). After the above treatment, his vision was normal (OD 1.2, OS 1.2), the intraocular tension was also normal, and the CAS decreased to 2/7. However, the disease relapsed approximately 2 months after completion of intravenous GC therapy, which manifested as severe exophthalmos, diplopia, vision decrease (OD 0.1, OS 1.0), and inability to close his eyelid. The CAS increased to 7/7. The patient was then diagnosed with DON and right orbital decompression surgery was performed. The surgical type and methods were similar to those described for patient 1. After orbital decompression surgery, his vision improved slightly (OD 0.4, OS 1.0); however, the CAS and diplopia did not improve. Therefore, we decided to administer rituximab treatment (a single dose of 500 mg) and oral GC treatment.

The treatment process of patient 1 and patient 2 was summarized in **Figure 1**. **Table 1** showed the baseline data of patients before rituximab treatment, **Table 2** showed in detail the patients undergoing orbital decompression surgery and rituximab treatment, and **Table 3** showed the changes in patients' vision before rituximab treatment.

RESULTS

As shown in **Table 4** and **Figure 2**, rituximab treatment was administered after orbital decompression and showed significant improvement in inflammatory symptoms, as manifested by a substantial decrease in CAS. Meanwhile, the vision of both patients improved significantly, and their diplopia was relieved after rituximab treatment.

In patient 1, the CAS decreased from 5/7 to 2/7 (OD) and from 6/7 to 3/7 (OS) 5 months after rituximab treatment, with a final CAS of both eyes of 2/7 at the last follow-up to 24 months after rituximab treatment. His vision had improved from 0.5 to 1.0 OD and from 0.6 to 0.8 (OS) at the last follow-up. The

TABLE 1 | Description of the two patients treated with Rituximab.

| Patient number | Patient 1 | Patient 2 |
|--|--|----------------------|
| Age (year) | 54 | 58 |
| Gender | Male | Male |
| Smoking | No | Yes |
| Quit smoking | N/A | 10 years ago |
| Endocrinological treatment of GD | Methimazole | Methimazole |
| Chronology of GO in relation to GD | With GD | With GD |
| Corticosteroids therapy (time relative to GO diagnosis) | 3 months | 0 month |
| Corticosteroids therapy (duration time) | 8 weeks | 23 weeks |
| Total dose of intravenous corticosteroids dose (methylprednisolone, MPS) | 4.0g | 8.5g |
| Other immunosuppressive treatments before RTX | Oral MPS for 2 weeks, total dose 532mg | None |
| Orbital radiotherapy | None | 2Gy/time × 10 times |
| Additional treatments | Selenium supplements | Selenium supplements |

GD, Grave's disease; GO, Grave's ophthalmopathy; RTX, rituximab; MPS, methylprednisolone; N/A, not available

TABLE 2 | Orbital decompression and Rituximab treatment details of the two patients.

| Patient number | Patient 1 | Patient 2 | | |
|--|---|--|--|--|
| Reason for orbital decompression | DON, no improvement after intravenous MPS | Relapse and DON developed after intravenous MPS and orbital radiotherapy | | |
| Total dose of MPS before orbital decompression | 3.5g | 8.5g | | |
| Duration of MPS use before orbital decompression | 7 weeks | 23 weeks | | |
| Orbital decompression (time relative to DON diagnosis) | At 7 weeks | At 1 week | | |
| Immunosuppressive treatments after orbital decompression | Intravenous MPS 0.5g and oral MPS | None (prior to RTX) | | |
| RTX therapy (time related to orbital decompression) | At 52 days | At 4 days | | |
| RTX therapy (time related to diagnosisi) | 5 months | 8 months | | |
| Reason for RTX therapy | Failure of glucocorticoids treatment | MPS dose exceeded 8g | | |
| RTX dose | 500mg only once | 500mg only once | | |
| Immunosuppressive treatments after RTX | Oral MPS | Oral MPS | | |

DON, dysthyroid optic neuropathy; MPS, methylprednisolone; RTX, rituximab.

TABLE 3 | Summary of changes in vision before RTX treatment.

| | Vision | | |
|---|--------|------|--|
| | Right | Left | |
| Patient 1 | | | |
| Before high-dose GCs pulse treatment | 0.3 | 0.08 | |
| After 7 times high-dose GCs pulse treatment | 0.1 | 0.1 | |
| After bilateral orbital decompression and the eighth high-dose GCs pulse treatment (before RTX) | 0.5 | 0.6 | |
| Patient 2 | | | |
| Before high-dose GCs pulse treatment combined with right orbital radiotherapy | 1.0 | 1.0 | |
| After high-dose GCs pulse treatment combined with right orbital radiotherapy | 1.2 | 1.2 | |
| Disease relapse | 0.1 | 1.0 | |
| After right orbital decompression (before RTX) | 0.4 | 1.0 | |

RTX, rituximab; GCs, glucocorticoids.

TABLE 4 | Results of the therapy with rituximab.

| | CAS score (/7) | | Vision | | IPO (mmHg) | | Proptosis (mm) | | Diplopia |
|--------------------------------------|----------------|------|--------|------|------------|------|----------------|------|------------|
| | Right | Left | Right | Left | Right | Left | Right | Left | |
| Patient 1 | | | | | | | | | |
| Before RTX | 5 | 6 | 0.5 | 0.6 | 19 | 15 | 21 | 20 | Yes |
| 5 months after RTX | 2 | 3 | 1.0 | 0.8 | 18 | 14 | 20 | 21 | Alleviated |
| Last examination 24 months after RTX | 2 | 2 | 1.0 | 0.8 | 18 | 15 | 19 | 20 | Alleviated |
| Patient 2 | | | | | | | | | |
| Before RTX | 7 | 3 | 0.4 | 1.0 | 21 | 19 | 19 | 19 | Yes |
| 6 months after RTX | 2 | 0 | 0.8 | 1.0 | 19 | 17 | 18 | 18 | Alleviated |
| Last examination 26 months after RTX | 0 | 0 | 1.0 | 1.2 | 19 | 17 | 18 | 19 | No |

CAS, Clinical Activity Score; IPO, intraocular pressure; RTX, rituximab.

severity of diplopia decreased but persisted until the last followup. Slight improvements in intraocular pressure and proptosis were also observed.

In patient 2, similar to patient 1, inflammatory signs and vision improved significantly after rituximab treatment, while intraocular pressure and proptosis improved slightly. Specifically, the CAS decreased from 7/7 to 2/7 (OD) 6 months after rituximab treatment, with a final CAS of 0/7 (OD) at the last follow-up. The patient's vision had improved from 0.4 to 1.0 OD and from 1.0 to 1.2 (OS) at 26 months after rituximab treatment. In addition, diplopia was alleviated at 6 months and completely relieved at 26 months after rituximab treatment.

DISCUSSION

Rituximab is a mouse anti-human monoclonal antibody that directly targets CD20, a B-lymphocyte-specific antigen (11). It was initially developed and used in patients with CD20-expressing lymphoid malignancies, including chronic lymphocytic leukemia and B-cell non-Hodgkin lymphoma. Rituximab induces cell apoptosis by binding to the CD20 membrane antigen in both normal and malignant B cells (12). In addition, it significantly depletes lymphocytes not only in the blood but also in the target tissue, and the effect remained significant 2 months after rituximab infusion (13). Based on the pharmacological mechanism of

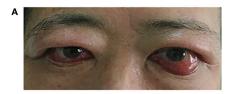








FIGURE 2 | Changes in the appearance of the two patients' eyes. (A) Before rituximab treatment (patient 1); (B) 24 months after rituximab treatment; (C) Before rituximab treatment (patient 2); (D) 26 months after rituximab treatment (patient 2).

rituximab, clinical trials on autoimmune diseases have indicated the potential use of rituximab as a candidate treatment for autoimmune diseases (14).

In 2006, two case reports indicated that corticosteroidresistant GO might be significantly relieved after treatment with rituximab (15, 16). Following the two successful treatment cases, several non-controlled studies on the effects of rituximab in GO have been published, with most showing that rituximab can be used in active moderate-to-severe GO, especially when intravenous methylprednisolone therapy fails (5). In one study, Salvi et al. reported good therapeutic effects after rituximab treatment and improved activity and severity of GO in 98 and 91% of 43 GO patients who failed to respond to corticosteroids, respectively (17). In recent years, the results of two randomized clinical trials of rituximab as a first-line treatment in moderateto-severe and active GO were published. In one study, Salvi et al. reported that GO was more significantly inactivated after 500 or 2,000 mg (administered twice over 2 weeks) rituximab injection (100 vs. 69% after 7.5 g intravenous methylprednisolone) at 24 weeks (18). However, another study by Stan et al. did not observe differences in either short- or long-term (24- or 52-week) outcomes between rituximab and placebo; meanwhile, two patients developed DON after rituximab. One possible explanation for the occurrence of DON is that the release of cytokines induced by rituximab treatment resulted in aggravated intraorbital edema, which transformed subclinical DON to DON (5, 19). The contradictory results of the two clinical trials suggest the need for more evidence before rituximab can be implemented as a first-line treatment for GO. However, the 2016 European Group of Graves' Orbitopathy Guidelines for the Management of Graves' orbitopathy recommended that, although it is still difficult to say which second-line option is more effective because of the limited evidence regarding differences in efficiency difference, treatment has relatively more evidence suggesting that rituximab is a good option for active moderate-to-severe GO when highdose intravenous GC pulse treatment fails (6).

In GO patients progressing to DON, very high doses of intravenous GCs (e.g. 500-1,000 mg of methylprednisolone for 3 consecutive days) is the first-line therapy; however, the effective rate is only 40%, and many patients require urgent decompression surgery when the response is poor (6). In the formulation of the patients' treatment plans in the present report, we considered that very high doses of intravenous GCs might be ineffective in patient 1 as he showed resistance GCs, which manifested as non-remission of the disease after seven intravenous GC treatments. Meanwhile, very high doses of intravenous GCs might have led to serious side effects in patient 2 as the cumulative doses of GCs exceeded 8 g. To relieve compression of the optic nerve and avoid further loss of vision, we performed orbital decompression surgery first instead of administering very high doses of intravenous GCs. However, as it was predictable before the surgery that the patients' condition, especially the inflammatory symptoms and eye movement disorder, did not significantly improve as orbital decompression could only relieve orbital pressure and optic nerve compression; thus, further drug treatment was urgently required.

No evidence-based medical recommendations exist regarding the further treatment of GO patients complicated with DON who fail high-dose GC pulse therapy and require orbital decompression. We finally decided to administer rituximab based on the results of several non-control studies that indicated a better curative effect for rituximab treatment than for first-line therapy in GO patients with DON (8-10). Among these studies, Chong et al. successfully treated four patients who failed to respond to GCs and developed DON. In these patients, the CAS decreased significantly 2 months after rituximab treatment and remained so in all patients. The vision also improved bilaterally in all four patients. One of the four patients received decompression surgery and fractionated orbital irradiation 2 months before rituximab infusion, and another patient received urgent decompression surgery 12 days after the first rituximab infusion due to continued DON (8). Two other studies by Salvi et al. and Mitchell et al. also reported the

successful treatment of five patients with DON, with significantly improved disease activity and vision in all five patients, similar to Chong's study. One of the five patients received urgent decompression surgery after the first rituximab infusion (9, 10). The longest interval between rituximab infusion and orbital decompression surgery was 2 months. Gess et al. administered rituximab to a patient with GO with DON who had failed very high-dose intravenous GCs. The patient improved initially but subsequently worsened 2 months later and underwent orbital decompression surgery (13).

The common characteristics of both patients in this study were significant enlarged extraocular muscle and orbital fat, which led to optic nerve compression and progression of DON. Previous studies revealed that the enlargement of extraocular muscle and orbital fat was associated with hyaluronic acid (HA) accumulation in the muscles and connective tissues and adipogenesis (20, 21). B cells play an important role in HA accumulation and adipogenesis. Briefly, B cells can present self-antigen and activate T cells through the complement receptor 2 (CR2, CD35), then activate helper T cells, recognize TSHR, ligate TSHR and TRAb together, and enhance hyaluronic acid (HA) production and adipogenesis (21, 22). Therefore, rituximab inhibits the activation of T cells by B-cell depletion and blockade of antigen presentation, which may be the main reason why rituximab is more effective compared with GC.

In conclusion, the treatment processes of these two patients indicated that rituximab might be safe and persistently effective for GO patients complicated with DON who fail high-dose GC pulse therapy and require orbital decompression. This finding is consistent with those of two previous non-control studies suggesting that rituximab treatment may have a better curative effect than very high doses of intravenous GCs. However, rituximab treatment may lead to the release of cytokines, aggravated intraorbital edema, and rapid vision loss. Orbital decompression before rituximab injection might mitigate this

side effect of rituximab; thus, the patients in this study might have obtained additional benefits from orbital decompression, although the surgeries were not performed preventively. The necessity for prophylactic orbital decompression surgery before rituximab injection requires additional evidence.

DATA AVAILABILITY STATEMENT

The raw data supporting the conclusions of this article will be made available by the authors, without undue reservation.

ETHICS STATEMENT

The studies involving human participants were reviewed and approved by Ethics Commission of Tongji Hospital, Tongji Medical College, Huazhong University of Science and Technology. The patients/participants provided their written informed consent to participate in this study. Written informed consent was obtained from the individual(s) for the publication of any potentially identifiable images or data included in this article.

AUTHOR CONTRIBUTIONS

BZ: Data curation, Project administration, Formal analysis, Writing—original draft, Writing—review and editing. YL: Data curation, Writing—review and editing. WX: Data curation, Writing—review and editing. BP: Data curation, Writing—review and editing. GY: Conceptualization, Data curation, Project administration, Formal analysis, Investigation, Methodology, Project administration, Validation, Writing—review and editing.

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Conflict of Interest: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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Epidemiology, Natural History, Risk Factors, and Prevention of Graves' Orbitopathy

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GO is the most frequent extrathyroidal manifestation of Graves' disease, although it may rarely occur in euthyroid/hypothyroid patients with chronic autoimmune thyroiditis. It is a relatively infrequent disorder, and men tend to have more severe ocular involvement at an older age. The prevalence of GO is lower than in the past among patients with recent onset Graves' hyperthyroidism, and moderate-to-severe forms requiring aggressive treatments are no more than 5–6% of all cases of GO. After an initial inflammatory (active) phase and a phase of stabilization (plateau phase), GO tends to improve and eventually inactivates (inactive or burnt-out phase). Minimal-to-mild GO often remits spontaneously, but complete restitutio ad integrum almost never occurs when GO is more than mild. Several risk factors contribute to its development on a yet undefined genetic background. Cigarette smoking is the most important of them. Early diagnosis, control and removal of modifiable risk factors, early treatment of mild forms of GO may effectively limit the risk of progression to more severe forms, which have a profound and dramatic impact on the quality of life of affected individuals, and remain a therapeutic challenge, often requiring long-lasting and multiple medical and surgical therapies.

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INTRODUCTION

Graves' orbitopathy (GO), also named Graves' ophthalmopathy, Thyroid-eye disease, or Thyroid-associated ophthalmopathy, is a relatively rare, but disabling and disfiguring disease of the orbit, in most cases associated with hyperthyroidism due to Graves' disease (1). GO is a disorder of autoimmune origin, the pathogenic mechanisms of which remain to be fully elucidated (2). The close link of GO with thyroid autoimmune disorders, mainly Graves' disease, underpin the hypothesis that GO is triggered by immune reactions against one or more antigens shared by thyroid and orbit. The thyrotropin (TSH) receptor is the most likely culprit, supporting an important role of TSH receptor antibodies (TRAbs) in the pathogenesis and course not only of thyroid disease, but also of orbital disease (3, 4). The insulin-like growth factor-1 (IGF-1) receptor seems also to be involved, which may explain the recent promising results obtained by treating patients with active, moderate-to-severe GO with an IGF-1 receptor antagonist monoclonal antibody, teprotumumab (5, 6). This would also be in keeping with a putative, but yet to be proven, protective role of antibodies to the IGF-1 receptor found in serum of patients with GO (7).

The histopathologic changes underpinning orbital remodeling (expansion of the fibroadipose tissue, swelling of the extraocular muscles), as well the increased production of hydrophilic glycosaminoglycans by orbital fibroblasts provide a mechanical basis for understanding most of the clinical features of GO (soft tissue changes/inflammation, exophthalmos, diplopia, compressive dysthyroid optic neuropathy) (8).

EPIDEMIOLOGY

Incidence

Information on the incidence of GO is scant. In a study of residents in the Olmsted County, Minnesota, through a 15-year interval (1976-1990), 120 incident cases of GO were found, with an ageadjusted incidence of all-degree GO of 16/100,000 population/year for women and 2.9/100,000 population/year for men (9) (Table 1). In a more recent prospective multicenter study from Sweden, covering a population of more than 3,500,000 individuals, newly diagnosed cases of Graves' hyperthyroidism from 2003 to 2005 were 2,200, with an incidence of 21/100,000 population/year; of these, 20.1% had eye involvement of all degrees (4.9% moderateto-severe), with an overall incidence of GO of 4.2/100,000 population/year (10). Because a 3.9:1 female to male ratio for Graves' hyperthyroidism was reported in that study, an approximate incidence of 3.3/100,000 population/year in women and 0.9/100,000 population/year in men can be derived for GO of all degrees (Table 1). Moderate-to-severe cases of GO would then have an incidence as low as 0.05/100,000 population/year, but classification criteria were not indicated in details (10). In a prospective registry-based study of patients with incident moderate-to-severe GO seen in Denmark during the period 1992-2009, 143 new cases of moderate-to-severe GO, classified following standardized criteria, were registered, with an incidence rate of 1.61/100,000 population/year (2.67/100,000 population/ year for women, 0.54/100,000 population/year for men) (11). Figures were similar before and after iodine fortification of salt (11). The Danish study did not consider patients with mild GO (11). However, because about two thirds of Graves' patients have no or mild GO (13, 14), it was calculated that the incidence of GO of all degrees derived from the Danish study (11) might approximately be 4.83/100,000 population/year (8.01/100,000 population/year in women, 1.62/100,000 population/year in men) (12) (Table 1). The considerable differences among the various studies underscore the difficulties in establishing the true

incidence of GO. Calculated figures derived from the two more recent studies (10, 11) appear to be lower than those reported in the older and pivotal study from USA (9). These differences might be apparent, due to difficulties in identifying patients with mild GO in registry-based studies (10, 11), or to persistent low referral rate of mild cases of GO, leading to an underestimation of all incident cases. On the other hand, this trend might be real and reflect a true decrease in the incidence of GO in recent years related to several factors, including changes in referral, improved interaction among general practitioners, endocrinologists and ophthalmologists, better control of risk factors, particularly smoking.

Prevalence

A recent study from the European Group on Graves' Orbitopathy (EUGOGO) used reported data on the incidence of GO (10, 11, 15) to estimate the prevalence of GO in the general population in Europe (12). By these calculations, it would appear that the prevalence of GO of all degrees in Europe might be between 90 and 155/100,000 population (12). Although this calculation is based on only three studies, these are large, epidemiological investigations and likely provide sound estimates of the overall prevalence of GO. Previous approximate calculations yielded an estimate of GO prevalence in Europe ranging from 100 to 305/100,000 population (16). The estimated prevalence derived from the Olmsted County, USA study was about 250/100,000 population (9). Other studies [reviewed in Ref. (17)] have led to an estimated prevalence of 100-300/100,000 population in Asia. Thus, although limitations inherent to the way of calculating prevalence from incidence data must be taken into account, it is reasonable to state that prevalence of GO does not substantially differ in different ethnic groups and is comprised between 90 and 300/100,000 population. GO, although relatively infrequent, does not fulfill the main criterion for being defined as a rare disease, i.e., a prevalence <50/100,000 population. However, several variants of the disease, namely, euthyroid GO, GO associated with thyroid dermopathy or acropachy are far below this threshold and can be considered as rare diseases, provided that distinct pathophysiological mechanisms can be identified (12).

Many patients with Graves' disease have no ocular involvement when first diagnosed with hyperthyroidism. A review of clinical records of the first 100 consecutive patients seen at a tertiary referral center in UK in 1960 and 1990 demonstrated that 57 and 35%, respectively, had clinically relevant GO with a parallel decrease in the proportion of severe forms in the two decades (**Figure 1**) (18) In a large and

TABLE 1 | Estimated incidence of Graves' orbitopathy (GO).

| Author (year) | Years of observation | Estimated Incidence in Women | Estimated Incidence in Men | Ref. |
|-------------------------|----------------------|---|---|------|
| Bradley (1994) | 1976-1990 | 16/100,000/year | 2.9/100,000/year | (9) |
| Abraham-Nordling (2011) | 2003-2005 | 3.3/100,000/year | 0.9/100,000/year | (10) |
| Laurberg (2012) | 1992-2009 | 2.67/100,000/year* 8.01/100,000/year** | 0.54/100,000/year* 1.62/100.000/year** | (11) |

^{*}Moderate-to-severe forms of GO.

^{**}All degrees of GO estimated by assuming that moderate-to-severe forms of GO represent 1/3 of cases (12).

more recent series of 346 patients with newly diagnosed and recent onset Graves' hyperthyroidism seen at a single center in Italy, almost three quarters had no ocular involvement, and only 6% had moderate-to-severe GO (including 0.3% with sightthreatening GO) (Figure 2) (13). A recent meta-analysis and systematic review of 57 studies including 26,804 patients reported an overall prevalence of 40% (19). It should be noted that included papers likely comprised patients with early onset GO and patients with GO of longer duration. Thus, it can be concluded that the overall prevalence of GO of all grades among patients with Graves' hyperthyroidism is comprised between 25 and 40%. It is possible that early diagnosis and treatment of hyperthyroidism, as well as use of preventive measures, including removal of modifiable risk factors (see a subsequent section of this manuscript), may reduce the prevalence of clinically significant GO to the lower end of the above spectrum, or even lower. In addition, in those patients who have GO at onset, early referral and management may be associated with a change in the clinical manifestations of GO. In the PREGO (Presentation of Graves' Orbitopathy) study, patients referred to EUGOGO centers in the year 2012 were compared to patients referred to

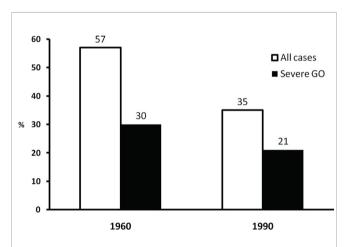


FIGURE 1 | Prevalence and severity of Graves' orbitopathy (GO) in the first 100 consecutive patients seen in a combined thyroid-eye clinic in UK in 1960 and 1990. Derived from Perros and Kendall-Taylor (18).

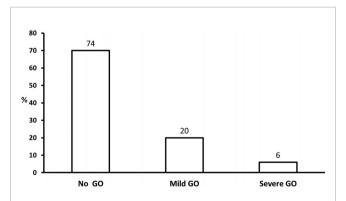


FIGURE 2 | Prevalence and severity of Graves' orbitopathy (GO) in 346 patients with recent onset and untreated Graves' hyperthyroidism. Severe GO: moderate-to-severe and sight-threatening GO. Derived from Tanda et al. (13).

EUGOGO centers in the year 2000: the 2012 cohort had a shorter referral time and showed a much higher prevalence of mild forms (60.5 vs. 41.2%) and inactive forms (63.2 vs. 39.9%) of GO when compared with the 2000 cohort (14). Nowadays, moderate-to-severe forms of GO, which remain a major therapeutic challenge, represent 5–6% of cases (11, 13). Although it is difficult to draw definitive conclusions, it seems that the proportion of Graves' patients with GO of all grades and, particularly, with severe forms of the disease is possibly declining over time (20).

Age, Gender, and Ethnicity

In the Olmsted County study, GO of all degrees showed a bimodal peak, 40-44 years and 60-64 years in women, 45-49 years and 65-69 years in men (9). In an observational Japanese study of 10,931 consecutive patients, the mean age of GO occurrence was 39 years in women and 43 years in men (21). In a study of 101 consecutive patients referred to a combined thyroid-eye clinic, the mean age was lower in patients without GO (40 years) than in those with GO (46 years) (22). In an Italian study mean age did not differ in Graves' patients without GO and in those with mild GO (46 and 44 years, respectively), but was significantly higher in patients with moderate-to-severe GO (54 years) (13). Likewise, in a Danish study of patients with moderate-to-severe GO, the median age was 50 and 56 years before and after salt iodization, respectively, and the risk of developing moderate-to-severe GO was lower in patients aged <40 years (11). Thus, age is a relevant factor affecting severity of GO, and the disease tends to be more severe in older patients (22). Although a questionnaire-based survey among European thyroidologists reported the presence of GO of all degrees in approximately one third of juvenile Graves' disease (23), clinically relevant GO in childhood is in general rarer than in adults and usually mild (24).

GO is more frequent in women than in men, although the female-to-male (F/M) ratio varies in different studies. In a study of 202 consecutive Graves' patients, the F/M ratio was 3.4 in patients without GO, 2.1 in patients with GO, and 0.7 in euthyroid GO (25). Other studies reported F/M ratios of 3.9 (21) and 4.2 (10). Gender affects also severity of GO, the F/M ratio progressively decreasing with increasing severity of GO (22). Likewise, in a cohort study of 2045 Graves' patients, although the proportion of patients with clinically relevant GO (NOSPECS class ≥2) was similar in women and men (51.5 and 52.7%, respectively), patients with more severe GO (NOSPECS class 4-6) were more frequently men (30.4 vs. 21.3%, p <0.001), and their median age was also higher than in women with similar severity of GO (52 years vs. 40 years, p <0.05) (27). Although a registry-based Danish study failed to show any significantly different risk of developing moderate-to-severe GO in men and women (11), it seems reasonable to conclude that GO tends to be relatively more frequent and severe in men, in whom it occurs at a more advanced age.

Relevance of ethnic factors in the occurrence of GO is controversial. In a study of 155 patients with newly diagnosed Graves' disease (116 Caucasians and 39 of Asian origin), the prevalence of GO was significantly higher in Caucasians than in

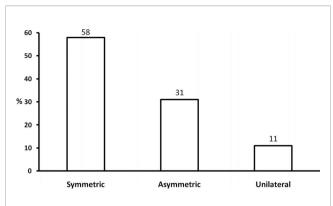


FIGURE 3 | Presentation of Graves' orbitopathy in a series of 269 patients referred to EUGOGO tertiary centers. Derived from Perros et al. (26).

Asians (42 vs. 7.7%, p = 0.0002), with a risk of developing GO 6.4-fold higher in Caucasians, possibly in relation to the high prevalence of smokers (>60%) among Caucasians (28). Conversely, a recent meta-analysis and systematic review showed a slightly lower prevalence of GO among Caucasians (37%) compared with Asians (45%) (19). A study from Australia reported that Caucasians have an increased risk (2.08, 95% CI, 1.56–2.76) of developing GO compared with non-Caucasian ethnicities (29). Conversely, other studies failed to link ethnic origin (including Caucasians, African Americans, Asian Americans, Asians, Latinos) to an increased for the occurrence of GO (30, 31). In summary, the role of ethnic factors, if any, remains, for the time being, unclear.

Phenotype

The majority of patients affected with GO have bilateral disease, but asymmetric or even unilateral GO may also develop (1). Asymmetric forms have been described in 4-14% of cases, unilateral forms in 9-34% of cases (9, 32-34). In a recent multicenter study of 269 patients referred to EUGOGO centers, although the majority (157 patients, 58%) had symmetric GO, many patients had either asymmetric or unilateral GO (26) (Figure 3). Interestingly, patients in the asymmetric group were older, were more frequently men, and tended to have more severe and active GO than the other groups (26). The reasons for asymmetric presentation of GO are unknown, but they might include differences in the anatomy of bony orbit or its vascularization. One study showed that patients with euthyroid/hypothyroid GO tend to have milder and more asymmetric forms of the disease (35). On the other hand, unilateral or asymmetric GO may sometimes progress to bilateral disease (34, 36). Asymmetric and unilateral forms of GO require an accurate diagnostic assessment to exclude other orbital diseases mimicking GO, reviewed in Ref. (37). Graves' disease may be an isolated disease or associated with other autoimmune disorders in the context of Autoimmune Polyglandular Syndromes: prevalence and severity of GO in these two different settings do not differ (38).

Relation With Thyroid Dysfunction

Most patients affected with GO are hyperthyroid (1), but a substantial proportion, ranging from 0.2 to 11% in different

studies (35, 39-41), are euthyroid or subclinical/overt hypothyroid. TSH-receptor antibodies (TRAbs) are the ultimate responsible for hyperthyroidism due to Graves' disease (42), but they also strongly correlate with the clinical activity and severity of GO (3, 4, 43). But, in a series of 700 unselected patients with chronic autoimmune thyroiditis, overt GO was present in 44 (6%): among them TRAb bioassay for stimulating activities tested positive in 30/44 (68%) patients with GO vs. 36/656 (5.5%, p <0.001) patients without GO (44). Thus, it is conceivable that TRAbs can explain the rare occurrence of GO in patients without hyperthyroidism. A note of caution should be added by mentioning that cases of so-called euthyroid/hypothyroid GO might indeed be a manifestation of another emerging disease, IgG4 ophthalmic disease, which has clinical features similar to GO (45). This underscores again the need for a careful differential diagnosis in doubtful cases.

A close relationship exists between the onset of Graves' hyperthyroidism and the onset of GO, because in about 85% of cases GO develops within 18 months before or after the onset of hyperthyroidism (25, 46). Therefore, a normal thyroid status does not exclude the diagnosis of GO, because hyperthyroidism may occur long after the development of GO. On the other hand, when GO follows hyperthyroidism, a relevant question is whether treatment for hyperthyroidism can affect the development and course of GO (47). Hyperthyroidism per se negatively influences the course of GO, and restoration of euthyroidism is often associated with a stabilization/ improvement of GO (48). But hypothyroidism can have a negative impact on the occurrence/progression of GO as well (45, 46). Thus, restoration and maintenance of euthyroidism is fundamental. While antithyroid drug treatment and thyroidectomy are apparently neutral to the course of GO, radioactive iodine (RAI) treatment is associated with a small, but definite risk of progression or de novo occurrence of GO, especially in smokers (49) (see a subsequent section).

Natural History

Following the initial description by Rundle and Wilson (50), it is widely accepted that overt GO clinically goes through an initial phase of inflammatory changes corresponding to ongoing activation of the pathogenic cascade of events occurring in the orbit (active phase); the disease then stabilizes when inflammation starts to subside (plateau or static phase), and then progressively improves in association with burning out of inflammation (inactive phase), without going back to normal (51). It is unknown how long it takes for GO to become inactive, but it is generally believed this occurs within 18-24 months. It should be mentioned that, although rarely, late reactivation of GO may occur (52). In a computer tomography-based study, the increase in the extraocular muscle volume appeared to be an early phenomenon, while the increase in the orbital fat volume occurred later (53). A precise definition of the natural course of GO of all degrees is, however, hampered by the fact that patients with moderate-to-severe and active GO cannot be left untreated and are, therefore, given disease-modifying treatments, namely intravenous glucocorticoids, orbital radiotherapy (54) or, more recently, biological agents (2).

In a series of 59 patients with untreated GO referred to a combined thyroid-eye clinic in UK, 13 patients (22%) improved substantially, 25 patients (42%) manifested only slight amelioration, 13 patients (22%) were stable, and 8 patients (14%) had a progression of GO over a median period of 12 months (55). In series of 196 Graves' patients, 81 (41%) developed GO, 53 of whom (65%) received no treatment, except for local measures: 25/53 (47%) improved substantially, 26/53 (49%) had unchanged GO at follow-up, 2/53 had a progression of GO (56). In a large series of 346 recent onset and untreated Graves' patients, 237 then completed a course of antithyroid drugs; 194 had no GO at baseline, and progression to moderate-to-severe GO occurred in <3% of cases; among the 43 patients with mild and inactive GO at baseline, progression to moderate-to-severe GO occurred in only 1 patient, while the majority of patients had a complete remission of ocular manifestations (13) (Figure 4). In another series of 65 patients, the large majority of whom had mild, minimally active GO and were followed without treatment for a median of 40 months, 51% improved spontaneously, 34% remained stable, 15% had some progression of GO (57). In summary, it would appear that mild/ minimal GO rarely progresses to severe forms, particularly if hyperthyroidism is adequately managed and controlled, while stabilization and remission is frequent in mild forms of GO (20). This favorable course of mild GO is favored by control of modifiable risk factors (see a subsequent section). No conclusion can be drawn on the natural history of moderateto-severe GO, because the latter is promptly treated by diseasemodifying therapies. However, in a retrospective study of 226 patients with initially moderate-to-severe and active GO reexamined after a median of about 4 years after GO onset and various non-surgical and surgical treatments for GO, further amelioration was observed in 60% of responders to treatments at the last visit (58). This suggests that time (i.e., the natural history) is a key factor affecting also the long-term outcome of GO after treatment (58). In addition, in the placebo group of the second teprotumumab study, at the end of 6 months of observation a reduction in exophthalmos ≥2 mm was observed in 10% patients, an amelioration in diplopia in 29%, and a CAS 0-1 (inactive GO)

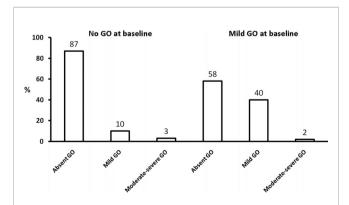


FIGURE 4 | Natural history of Graves' orbitopathy (GO) at the end of antithyroid drug treatment according the absence (n = 194) or presence (n = 43) of mild GO at baseline. Derived from Tanda et al. (13).

in 21% (6). This suggests that the Rundle's curve description applies also to patients with moderate-to-severe GO. Obviously, this by no means implies that in this category of patients therapies should be postponed, because early immunosuppressive treatments improve and inactivate GO, thereby affecting substantially the natural history of GO, and reduce the time interval required to proceed to rehabilitative surgery, if needed.

RISK FACTORS AND PREVENTION

GO results from a complex interaction of endogenous (unmodifiable) and exogenous/environmental (modifiable) risk factors. The former include age, gender, and genetic factors. As described in a previous section of this manuscript, GO tends to be more severe in men, in whom it occurs at an older age than in women. A search for genetic factors did not provide unequivocal information, with particular regard to differentiation between Graves' patients with or without GO (59–61). Likewise, results are conflicting as to the association between GO and Major Histocompatibility Complex (MHC), cytotoxic T-lymphocyte-associated antigen-4 (CTLA-4), protein tyrosine phosphatase or non-receptor type 22 (PTPN22), interleukins, intercellular adhesion molecule 1 gene polymorphisms, adipogenesis-related genes, TSH receptor or thyroglobulin (17, 62, 63).

Several modifiable risk factors for the occurrence/progression of GO have been identified. This is a very important issue, because controlling them may positively affect the course of GO (64) (**Table 2**).

Smoking

Smoking habit is probably the most important modifiable risk factor for GO (65, 66) (Table 2). The negative impact of smoking on GO is based on the following evidence: i) Graves' patients with GO are more frequently smokers (and heavy smokers) than those without GO, or patients with other thyroid disorders, including chronic autoimmune thyroiditis (67); ii) Among Graves' patients, smokers have a higher risk of developing GO than non-smokers (odds ratio, 7.7, 95% CI, 4.3-13.7, vs. 1.9, 95% CI, 1.1–3.2) (68); iii) Smokers are at high risk of developing severe forms of GO, with a dose-dependent relative risk of diplopia or exophthalmos (69) (Figure 5); iv) Response to treatments for GO is decreased and occurs later in smokers than in non-smokers (70, 71); v) De novo development or progression of GO after RAI treatment occurs more frequently in smokers (29, 49, 72); vi) Quit smoking decreases the risk of developing exophthalmos and diplopia, suggesting that current smoking is more important than lifetime tobacco consumption (69). Accordingly, refrain from smoking is a fundamental, general, preventive action strongly recommended by guidelines (54). It is possible, but yet unproven, that also passive smoking be relevant for the occurrence of GO (23). Whether e-cigarette may also concur to development/ progression of GO remains for the time being unsettled. Mechanisms whereby cigarette smoking exerts a negative effect on GO are not fully understood, but they might involve oxygen free radical generation, hypoxia in the orbit, enhanced production of cytokines, stimulation of adipogenesis (64).

TABLE 2 | Risk factors for the occurrence/progression Graves' orbitopathy (GO) and preventive actions.

| Risk Factor | Evidence (Ref.) | Preventive Action |
|---|--|--|
| Smoking Habit | Patients with GO are more frequently smokers than those without GO (67) | Urge patients to refrain from smoking |
| | Smokers have an increased risk of developing GO than non- smokers (68) | |
| | Smokers have a dose-dependent increased relative risk of developing severe GO (69) | |
| | Smokers have a lower and slower response to immunesuppressive treatments (70, 71), | |
| | Refrain from smoking reduces the risk of developing exophthalmos and diplopia (69) | |
| | Radioiodine associated progression og GO is more frequent in smokers (70, 72), | |
| Thyroid Dysfunction | Hyperthyroidism negatively influences GO and amelioration is associated with restoration of euthyroidism (48) GO may develop during uncontrolled hypothyroidism (73) | Restore and stably maintain euthyroidism by antithyroid drugs (hyperthyroidism) or levothyroxine replacement (hypothyroidism) |
| Radioiodine treatment for hyperthyroidism | • Radioiodine treatment may cause occurrence or progression of GO (49, 74), particularly in smokers (72) | Give a short course of oral prednisone (steroid prophylaxis) in at risk patients |
| Oxidative stress | GO is associated with an increased oxidative stress (75) | Provide a 6-month selenium supplementation in patients with mild GO of short duration |
| TSH receptor antibodies (TRAbs) | TRAb levels are higher in patients with GO than in those without GO (76) TRAb levels correlate with GO activity and severity (4, 43, 76), | In patients with relevant GO, control hyperthyroidism with antithyroid drugs, because this is usually associated with a decline in TRAb levels |
| Hypercholesterolemia | Serum total and LDL cholesterol levels correlate the presence and activity of GO (77) | Correct dyslipidemia in patients with newly diagnosed Graves' hyperthyroidism |
| | Serum total and LDL cholesterol levels are higher in patients with GO than in those without GO (78) | |
| | The use of statins was associated with a decreased risk of developing GO (31) | |

Thyroid Dysfunction

We mentioned above that both hyperthyroidism and hypothyroidism represent risk factors for the occurrence of GO (46, 48). Accordingly, a rapid restoration of euthyroidism and its stable maintenance, avoiding thyroid status fluctuations, are strongly recommended by guidelines as an important preventive measure (54) (**Table 2**). Mechanisms whereby thyroid dysfunction may favor the occurrence or progression of GO are probably related to the activation of the TSH receptor by either TRAbs (hyperthyroidism) or TSH (hypothyroidism) (64).

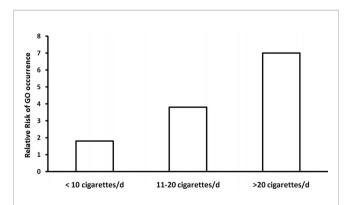


FIGURE 5 | Relative risk of developing exophthalmos and diplopia in relation to daily number of cigarettes. Derived from Pfeilschifter and Ziegler (67).

Radioiodine Treatment

RAI is a well-established and effective method of treatment for Graves' hyperthyroidism (79). RAI carries a small but definite risk of causing progression or de novo occurrence of GO, probably due to antigen release and related exacerbation of autoimmune reactions following RAI administration (1) (Table 2). In a randomized clinical trial of 168 patients, 114, aged 35 to 55 years, were randomized to either antithyroid drug treatment, RAI treatment, or subtotal thyroidectomy: GO developed or worsened in 4/38 patients (10%) treated with antithyroid drugs, 6/37 patients (16%) treated surgically, and 13/39 patients (33%) treated with RAI (74). In a randomized clinical trial of 443 patients, GO progressed in 23/150 patients (15%) treated with RAI, but in only 3/148 patients (3%) treated with antithyroid drugs (49). RAI-associated progression of GO was transient in some patients, but was permanent and required immunosuppressive treatments in 5% of patients (49). This untoward effect could be prevented in a third group of patients who received a concomitant short-term course of oral prednisone (steroid prophylaxis) (49), confirming a previous randomized clinical trial from the same group (80). This undue effect of RAI is more likely to occur in smokers (72), while it is unlikely in patients whose Graves' disease duration is >5 years (81) or whose GO is stably inactive (73). Relatively lower doses of oral prednisone (0.1–0.2 mg/bodyweight as starting dose, gradually tapered and withdrawn after 6 weeks) are equally effective as previously suggested highly doses (0.3–0.5 mg/bodyweight as starting dose, tapered and withdrawn after 3 months) (82) and safe (83). Based on the above studies, steroid

prophylaxis is an important preventive measure in most Graves' patients undergoing RAI treatment for hyperthyroidism, and as such it is recommended in most patients by recent guidelines (54, 79). In particular, it should be given in patients who have mild and active GO and/or risk factors for its development/progression, especially smoking, while It can be avoided in patients with long-standing and inactive GO.

Oxidative Stress

Graves' orbitopathy is associated with an increased oxidative stress (75). Because of its antioxidant and immunoregulatory actions, selenium has been proposed as an adjuvant therapy in patients with mild GO. In a randomized clinical trial of 159 patients with mild and active GO, selenium administration for 6 months was associated with an improved quality of life and improved ocular involvement compared with placebo; in addition, it was more effective than placebo in preventing progression of GO from mild to moderate-to-severe (84). It is unclear whether selenium should be used only in patients who are selenium-deficient or also in patients with an adequate selenium intake. It is probably useless in patients with longstanding, inactive mild GO. Likewise, there is no evidence that a course of selenium may be beneficial also in patients with moderate-to-severe GO. With these limitations, current guidelines recommend a 6-month selenium supplementation in patients with recent onset, mild GO, because it improves ocular involvement and prevents progression to more severe forms of GO (54).

TSH-Receptor Antibody Levels

TRAbs are the only specific biomarker for Graves' disease and GO (3). TRAb levels correlate with the Clinical Activity Score (4, 43). TRAbs with stimulating activities in a recently developed bioassay tested positive in 150/155 patients with Graves' disease (97%) and 148/155 of those with GO (95%), their levels were 3-fold higher in patients with GO than in those without GO, and strongly correlated with GO activity and severity (76). TRAbs were shown to be an independent risk factor for GO and predictors of severity and outcome of the disease (85). In a study of 100 consecutive patients, the combination of high TRAb levels and absent thyroid peroxidase antibodies identified Graves' patients at high risk of developing GO (86). This finding was subsequently confirmed by a 3-year prospective study from the same group (87). In terms of prevention, for the time being, there is no treatment that blocks TRAb synthesis. However, while RAI treatment for hyperthyroidism is followed by a rise in TRAb levels (probably related to the cytolytic effect of RAI), which may last for several years, antithyroid drug treatment (either directly or through restoration of euthyroidism) and thyroidectomy are generally and gradually associated with a progressive reduction of serum TRAb concentration, which may be beneficial for GO (88).

Hypercholesterolemia

In a cross-sectional study of 250 patients with Graves' hyperthyroidism of recent onset, with (n = 133) or without GO (n = 117), a correlation between serum total and LDL

cholesterol and the presence and activity of GO was reported (77). A second study, from the same group, of 86 consecutive patients with recent onset Graves' disease referred for RAI treatment, confirmed that serum total and LDL cholesterol were higher in patients than in those without GO, although there was no relationship between cholesterol and activity or severity of GO (78). The above reports might explain why the analysis of a large database from USA, including 8,404 patients with newly diagnosed Graves' disease, showed that the use of statins for at least two months in the year of observation was associated with a 40% decreased hazard ratio (HR) of developing GO (HR: 0.60, CI, 0.47-0.75) (31). Because the use of non-statin cholesterol lowering drugs was not associated with a decreased risk of GO occurrence (31), it is unsettled whether the purported protective/preventive effect of statins on GO is related to their anti-inflammatory action or to the cholesterol lowering effect. Clearly, further studies are needed to clarify this issue, but it is reasonable to state that correction of dyslipidemia in newly diagnosed Graves' patients may represent a useful preventive action against the occurrence of GO.

Prediction of GO Occurrence in Newly Diagnosed Graves' Patients

Because of the major role of exogenous factors in the development of GO, a prospective observational study was carried out in 10 EUGOGO centers to construct a predictive score (called PREDIGO, prediction of GO) for the development/ progression of GO in 348 newly diagnosed and untreated Graves' patients without obvious GO (89). Four independent variables composed the PREDIGO score, i.e., the Clinical Activity Score, TRAbs (measured as TSH-binding inhibiting immunoglobulins, TBII), duration of hyperthyroidism, and smoking. Of the 348 patients, all submitted to antithyroid drug treatment, 53 (15%) developed GO, which was mild in 46 (13%) and moderate-tosevere in 7 (2%) (78). The PREDIGO score proved to be much better in identifying patients who will not develop GO (negative predictive value: 0.91, 95% CI, 0.87-0.94) than those who will (positive predictive value: 0.28, 95% CI, 0.20-0.37) (89). Although this represents an evident limitation and underpins the need for the improvement and refinement of this score with additional biomarkers, the PREDIGO score may represent a simple and useful tool to consider when discussing the general treatment plan of newly diagnosed Graves' patients with absent or very mild GO. Moreover, as mentioned earlier, early referral of patients with mild and active GO and/or effective control of modifiable risk factors for development/progression of the disease may substantially contribute to exacerbation of GO (54).

CONCLUDING REMARKS

GO is the most frequent extrathyroidal manifestation of Graves' disease, although it may rarely develop also in euthyroid/hypothyroid patients with chronic autoimmune thyroiditis. It is a relatively infrequent disorder, and men tend to have more severe ocular involvement at an older age. After an initial

inflammatory (active) phase and a phase of stabilization (plateau phase), GO tends to improve and eventually inactivates (inactive or burnt-out phase). Mild GO tends spontaneously to remit spontaneously, but complete *restitutio ad integrum* almost never occurs when GO is more than minimal-to-mild. Several risk factors contribute to its development on a yet undefined genetic background. Cigarette smoking is the most important risk factor. Early diagnosis, control and removal of modifiable risk factors, early treatment of mild forms, stable control of thyroid dysfunction may effectively limit the risk of progression to more severe forms of GO, which have a profound and dramatic impact on the quality of life of affected individuals, and remain a therapeutic challenge, often requiring long-lasting and multiple medical and surgical therapies.

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Antioxidant Therapy in Graves' Orbitopathy

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The balance of the cell redox state is a key point for the maintenance of cellular homeostasis. Increased reactive oxygen species (ROS) generation leads to oxidative damage of tissues, which is involved in the development of several diseases, including autoimmune diseases. Graves' Orbitopathy (GO) is a disfiguring autoimmune-related condition associated with Graves' Disease (GD). Patients with active, moderate-to-severe GO, are generally treated with high doses intravenous glucocorticoids (ivGCs) and/or orbital radiotherapy. On the contrary, up to recently, local ointments were the treatment most frequently offered to patients with mild GO, because the risks related to ivGCs does not justify the relatively poor benefits expected in mild GO. However, a medical treatment for these patients is heavily wanted, considering that GO can progress into more severe forms and also patients with mild GO complain with an impairment in their quality of life. Thus, based on the role of oxidative stress in the pathogenesis of GO, a therapy with antioxidant agents has been proposed and a number of studies have been performed, both *in vitro* and *in vivo*, which is reviewed here.

Keywords: Graves' orbitopathy (GO), oxidative stress, Quercetin, Enalapril, Vitamin C, N-acetyl-I-cysteine, melatonin, β -carotene

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INTRODUCTION

Graves' Orbitopathy (GO) is an autoimmune, disfiguring disease observed in ~25%–30% of patients with Graves' Disease (GD). More rarely, GO can be found in patients with hypothyroid autoimmune thyroiditis or in subjects with subclinical evidence of thyroid autoimmunity, but with no thyroid dysfunction, and the pathogenesis of these conditions is still a matter of debate (1, 2). Patients affected with moderate-to-severe and active GO are generally treated with high dose intravenous glucocorticoids (ivGCs), orbital radiotherapy, surgical procedures (3) as well as new medications, including Rituximab, an anti-CD20 monoclonal antibody (4), Teprotumumab, an anti-insulin-like growth factor-1 receptor (IGF1R) monoclonal antibody (5, 6), Tocilizumab, an interleukin 6 (IL-6) receptor antagonist (7) and mycophenolate (8). Fortunately, the majority of GO patients have a mild disease and are typically treated with local measures (3), because the risks carried by the abovementioned therapies is not justified in view of the relatively poor benefits in mild GO, unless there is an important impairment of the quality of life of patients (9–11). Nevertheless, patients affected with mild GO still complain with a significant impairment in their quality of life, and although GO can improve spontaneously in up to 50% of them, it can worsen in ~15% of patients, thereby leading to the

search for treatments that are suitable for these patients (2, 12). Oxidative stress plays an important role in the pathogenesis of GO, being involved in the production of pro-inflammatory cytokines and hyaluronic acid (HA), as well as in promoting proliferation of fibroblasts and their differentiation into adipocytes (13). Keeping in mind that an ideal treatment for mild GO should be effective, well tolerated and widely available, several antioxidant agents have been investigated as new possible approaches for the management of these patients. The role and efficacy of selenium have been well demonstrated, as extensively discussed in another review in the same issue of this journal. In addition to selenium, various antioxidant agents, namely pentoxifylline, nicotinamide, allopurinol, quercetin, enalapril, vitamin C, N-acetyl-L-cysteine and melatonin, have been proposed to play a potential therapeutic role in GO (14–18).

OXIDATIVE STRESS IN GRAVES' ORBITOPATHY

Oxidative stress is defined as the disruption of the balance between the production and the elimination of reactive oxygen species (ROS), which causes a remarkable damage of several cell components, namely proteins, lipids, membranes, and nucleic acids (19, 20). The process ultimately results in mitochondrial dysfunction and loss of enzymatic activity. The ultimate pathogenetic mechanisms of GO are still unclear, but the most popular hypothesis implies an interplay between cellular and humoral immunity against the thyrotropic hormone (TSH) receptor (TSH-R) and possibly other autoantigens expressed by thyrocytes and orbital fibroblasts (OFs) (2). Following recognition of these antigens by autoreactive T lymphocytes, they infiltrate fibroadipose orbital tissue, thereby triggering the production of pro-inflammatory cytokines, growth factors and chemokines. The inflammatory response and the consequent local cell damage cause a dysregulated production of ROS, with saturation of the antioxidant systems. The interplay between cellular and humoral immunity, as well as the uncontrolled release of inflammatory molecules and ROS, leads to the increased proliferation of OFs, the differentiation of connective orbital cells into pre-adipocytes and then adipocytes, and the synthesis of great amounts of glycosaminoglycans (GAGs), especially hyaluronic acid (HA). The resulting orbital remodeling is characterized by extraocular muscle enlargement and orbital fat expansion, which are ultimately responsible for the clinical manifestations of the disease (12, 21, 22). Among the various players involved in the pathogenesis of GO, oxidative stress is believed to have a major role. Tissue hypoxia, as well as ROS, are involved in the typical changes of fibroadipose orbital tissue and of the perimysium of extraocular muscles (23) (Figure 1). ROS, and the consequent antioxidant defense mobilization, are present at sites of inflammation in general and, in the case of GO, together with edema and tissue hypoxia, contribute the damage of orbital tissue. A role of oxidative stress in GO is supported by a number of basic studies suggesting that ROS promote the proliferation of OFs and the release of GAGs, the synthesis and secretion of cytokines, and

the expression of other factors involved in the pathogenesis of GO, namely heat shock protein 72 (HSP-72), HLA-DR and ICAM-1 (14).

In this context, a number of studies, performed both *in vitro* and *in vivo*, investigated the effects of antioxidant agents in GO, including selenium, pentoxifylline, quercetin, enalapril, vitamin C, N-acetyl-L-cysteine and melatonin (14–18, 24, 25). Some of these studies have provided convincing or at least promising results, and in the case of selenium, have led to the clinical use of these antioxidant agents. In addition to the above mentioned compounds, statins (3-hydroxy-3-methylglutaryl-coenzyme reductase inhibitors), because of their anti-oxidative actions, have also been considered for GO treatment (26).

Oxidative Stress and Proliferation of Orbital Fibroblasts

Burch et al. performed a study on primary culture of OFs obtained from GO patients and control subjects, demonstrating that superoxide radicals induced an increased proliferation of OFs from GO patients compared with control fibroblasts (27). Moreover, preincubation of OFs with anti-thyroid drugs (ATD) and antioxidant agents as allopurinol and nicotinamide inhibited the proliferation induced by ROS (27).

The hypothesis that oxidative stress contributes GO pathogenesis was also suggested by studies performed by Tsai et al. and Hondur et al. (28, 29). They both showed that H_2O_2 , a potent ROS-inducer, and superoxide dismutase (SOD), a natural antioxidant agent, were higher in GO then control OFs, whereas the ROS antagonist glutathione peroxidase (GPX) was reduced, with a reduced ratio between glutathione and oxidized glutathione, suggesting the presence of a remarkable state of oxidative stress (28–31).

Glycosaminoglycan Production and Heat Shock Protein 72 Expression in Orbital fibroblasts

As mentioned above, the release of GAGs plays an important role in the pathogenesis of GO. The phenomenon is triggered by a number of cytokines, among which interleukin-1 (IL-1) is mostly involved (1, 2). Lu et al. reported that IL-1 β promotes an increase in ROS release in OFs from patients with GO (32). SOD and catalase, another ROS antagonist, to some extent reduced the secretion of GAGs induced by IL-1 β , suggesting that ROS may be involved in the IL-1 β -dependent release of GAGs. In addition, pentoxifylline, which has antioxidant properties, was found to be able to inhibit GAG accumulation in GO OFs (33). HSP-72 is involved in antigen recognition and T-lymphocyte activation in orbital tissue, and its expression in GO fibroblasts is promoted by H_2O_2 (34),. The effects of H_2O_2 were reduced by anti-oxidative agents or ATD (35), suggesting that ROS play a role in the aberrant expression of HSP-72 in OFs of GO.

Clinical Evidence of a Role of Oxidative Stress in Graves' Orbitopathy

In addition to basic studies, the importance of oxidative stress in GO is supported by clinical investigations. Cigarette smoking is

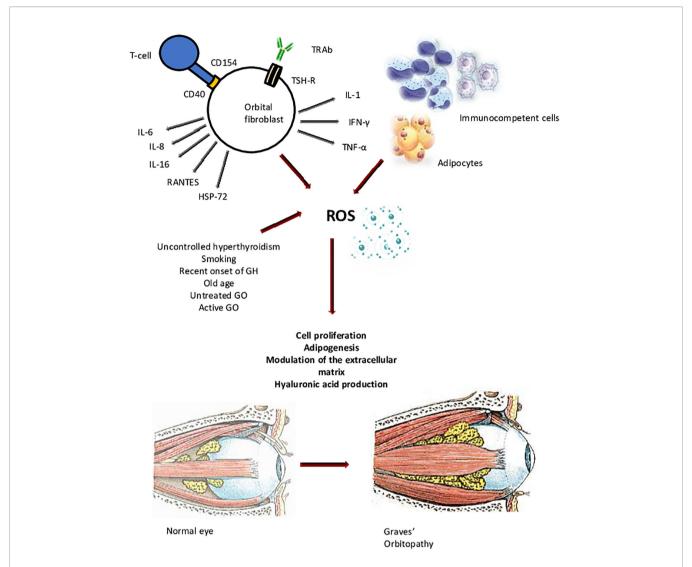


FIGURE 1 | Orbital fibroblasts (OFs) are activated by both cellular and humoral immune responses against autoantigens, among which the thyrotropic hormone (TSH) receptor (TSH-R) is the main involved. The inflammatory process triggers the production of pro-inflammatory cytokines, growth factors, chemokines, and ROS, which contribute the increased proliferation of OFs, the differentiation of connective orbital cells into pre-adipocytes and then adipocytes, and the synthesis of great amounts of glycosaminoglycans (GAGs), especially hyaluronic acid. The resulting orbital remodeling is characterized by extraocular muscle enlargement and orbital fat expansion, which are ultimately responsible for the clinical manifestations of the disease. IL-6, interleukin-6; IL-8, interleukin-8; IL-16, interleukin-16; RANTES, regulated upon activation, normal T-cell expressed and presumably secreted; HSP-72, heat shock protein-72; IL-1, interleukin-1; IFN-γ, interferon γ; TNF-α, tumor necrosis factor α; GH, Graves' hyperthyroidism; GO, Graves' orbitopathy.

the most important environmental risk factors for GO. Thus, cigarette smoke enhances the *in vitro* generation of ROS and reduces the antioxidant machinery (36). Patients with a recent onset Graves' hyperthyroidism (GH) have higher plasma levels of SOD and catalase than control subjects, regardless of the presence of GO (37). No differences in serum GPX and thioredoxin reductase (TRs), another ROS antagonist, were observed. However, the normalization of oxidative markers following restoration of euthyroidism was observed only in patients without GO, suggesting that orbital inflammation contributes the increased serum markers of oxidative stress (38–40). Moreover, higher urinary levels of 8-hydroxy-2'-deoxyguanosine (8-OHdG), a marker of oxidative DNA

damage, were found in patients with active GO (29, 31). Interestingly, urinary 8-OHdG was higher in smokers and in patients with GO relapse, and oral glucocorticoid (GC) administration led to a significant decrease of 8-OHdG levels, which was also associated with a reduction of GO activity and severity. These data were confirmed by a subsequent study performed in a larger cohort of GO patients, where there was a significant positive correlation between the GO Clinical Activity Score (CAS) and urinary levels of 8-OHdG. The relationship between oxidative stress and GC has also been studied (30) Akarsu et al. reported a significant reduction in the serum levels of malondealdyde (MAD), a marker of oxidative stress, following GC administration in GO patients (41). Similar findings were

obtained by Abalovich et al. (38) and Bednarek et al. (39), in line with an involvement of oxidative stress in the pathogenesis of GO.

ANTIOXIDANT AGENTS IN GRAVES' ORBITOPATHY

According to the guidelines of the European Thyroid Association and of the European Group on Graves' Orbitopathy (EUGOGO), a number of therapies can be offered to patients with moderateto-severe and active GO. Even though mild GO often improve spontaneously (in ~50% of cases), it worsens in up to 15% of patients (3, 12) and, because of the disfiguring features of the disease and the impairment in the quality of life, also patients with mild GO ask for some sort of treatment, which has been searched for over the last decades. Keeping in mind that an ideal treatment for mild GO should be effective, well tolerated and widely available, antioxidant agents have been proposed as a reasonable approach. Selenium has been largely investigated and currently used in the clinical practice but will not be mentioned in this review article as it is treated largely elsewhere in this special issue. In addition to selenium, a protective role in GO has been suggested by studies on other antioxidant agents, as reported below and summarized in Table 1.

Nicotinamide and Allopurinol

Allopurinol is a drug widely used in the management of hyperuricemia and gout, with powerful antioxidant activities, being an inhibitor of the enzyme xanthine oxidase (45). Vitamin nicotinamide is a largely studied nicotinamide adenine dinucleotide (NAD⁺) precursor, which is a coenzyme involved in a number of metabolic pathways and in the balance of cell redox state (46). The effects of nicotinamide are currently being investigated by some clinical trials (46). The potential role of allopurinol and nicotinamide in pathophysiologic conditions involving oxidative stress led to the investigation of their effects in GO. As mentioned above, Burch et al. reported that allopurinol and nicotinamide inhibit the proliferation of OFs induced by superoxide (27). This evidence was confirmed by a subsequent in vitro study performed in cultured OFs from patients with GO, incubated with interferon γ (IFN- γ) and tumor necrosis factor α (TNF- α) in the presence of nicotinamide (47). The authors reported an inhibitory effect of nicotinamide on cytokine production and OF proliferation. Moreover, nicotinamide was also able to reduce HSP-72 and HLA-DR as well as the expression of ICAM-1 expression, which is involved in the recruitment of T-lymphocytes to the orbit and enhances immune reactions and cytotoxicity (47-50). Bouzas et al. performed the first clinical, pilot study on antioxidants, which included 22 consecutive patients with mild or moderately severe GO treated with placebo or allopurinol (200 mg daily) plus nicotinamide (300 mg daily) for 3 months (42). All patients recruited had an active GO of recent onset (<6 months) and were euthyroid for at least 2 months. The response to treatment was evaluated by a complete ophthalmologic examination 1 and 3 months after the beginning of treatment. A significant

improvement of GO was observed in 9 (82%) and 3 (27%) patients who respectively received antioxidant agents or placebo. The inflammation of soft tissue was the feature that responded more to the treatment, whereas proptosis was little affected. Clearly, further studies in larger series of patients are needed to confirm these promising results.

Pentoxifylline

Pentoxifylline is an analog of the methylaxantine theobromine with anti-inflammatory and antioxidant properties (51). It is commonly used in the management of patients with vascular diseases (52). It has been demonstrated that pentoxifylline inhibits the synthesis of GAG and the proliferation of human skin fibroblasts (53). Similar results were reported in primary cultures of fibroblasts derived from patients with fibrotic diseases, including hypertrophic scar, scleroderma and keloid (54). In 1993 Chang et al. reported that treatment with pentoxifylline inhibits GAG release and fibroblast proliferation in primary cultures of OFs derived from patients with GO and control subjects, with no significant difference between them (33). Based on this evidence, some clinical studies investigated the potential beneficial effect of pentoxifylline in patients with GO. Balazs et al. performed a pilot study, neither randomized nor controlled, in 10 patients with GO selected for having contraindications to GCs (43). They were euthyroid for at least 4 weeks and they had a mild or moderate-to-severe GO. The patients were treated with a daily infusion of pentoxifylline (200 mg/die) for 10 days, following which treatment was continued orally (1800 to 1200 mg/day) and stopped after 3 months. At the end of treatment, a significant improvement of GO was observed in 8 of 10 patients. The beneficial effect was more evident on soft tissue inflammation whereas diplopia and proptosis were less affected by treatment. The randomized clinical trial performed by Marcocci et al. which opened for the clinical use of selenium in patients with GO, also investigated the effects of pentoxifylline. The authors showed that pentoxifylline was not associated with a significant improvement of the eye disease in terms of overall clinical outcome as well as the quality of life of patients (44).

Quercetin

Quercetin is a member of the flavonoid family contained in vegetables and fruits, which assures an adequate intake with diet (16). Quercetin has anti-inflammatory, antioxidant, anti-viral activity, and is also able to promote apoptosis in tumor cells (55, 56). It has been reported that quercetin blocks the transforming factor- β (TGF β)/Smad signaling pathway, leading to the inhibition of the proliferation of scar-derived fibroblasts (57, 58). In addition, an antifibrotic activity in fibroblasts and adipocytes from GO patient has been reported, and quercetin seems also to be able to reduce the adipogenesis induced by cigarette smoke-extract in GO fibroblasts. Based on these evidences, a number of in vitro studies evaluated the possibility that quercetin may reduce the proliferation of OFs in GO. A study performed in primary cultures of OFs from GO patients and control subjects demonstrated that quercetin exerts an inhibitory effect on fibroblasts proliferation and HA release, with no difference between GO and control fibroblasts (16).

TABLE 1 | Antioxidant agents in Graves' orbitopathy (GO).

| Compound | Main finding | Methods | Results | Reference |
|--|--|---|--|----------------------------------|
| Nicotinamide and Allopurinol | Clinically significant improvement of GO | A pilot study including 22 consecutive patients with mild or moderately severe, active GO of recent onset (<6 months). They were treated with placebo or allopurinol (200 mg daily) plus nicotinamide (300 mg daily) for 3 months. | After 3 months, a significant improvement of GO was observed in 9 (82%) and 3 (27%) patients who respectively received antioxidant agents or placebo. The inflammation of soft tissue was the feature that responded more, whereas proptosis was little affected. | (42) Bouzas EA. et al. |
| Pentoxifylline | Inhibition of GAG release and fibroblast proliferation | Primary cultures of OFs from GO patients and control subjects were obtained. Cell proliferation and GAG production were measured after the addition of pentoxifylline (0.1–1,000 mg/L). | The exposure of OFs cultures to pentoxifylline resulted in a dose-dependent inhibition of fibroblast proliferation and GAG synthesis. | (33) Chang C.C et al. |
| | Clinically significant improvement of GO | A pilot study, neither randomized nor controlled, which involved 10 patients with active, mild or moderate-to-severe GO. Patients were treated with a daily infusion of pentoxifylline (200 mg/die) for 10 days, following which treatment was continued orally (1,800 to 1,200 mg/day) and stopped after 3 months. | At the end of treatment, a significant improvement of GO was observed in 8 of 10 patients. The beneficial effect was more evident on soft tissue inflammation, whereas diplopia and proptosis were less affected. | (43) Balazs C. et al. |
| | No improvement of GO | A prospective, multicenter, placebo-controlled clinical trial in which 159 patients with mild GO were randomized to receive sodium selenite (100 µg twice daily orally), pentoxifylline (600 mg twice daily orally) or placebo for 6 months. | Unlike selenium, no beneficial effects of pentoxifylline were observed | (44) Marcocci NEJM |
| Quercetin | Reduction of cell proliferation and HA release in GO fibroblasts | Primary cultures of OFs from GO patients and control subjects were obtained. Cell proliferation, cell necrosis, apoptosis and HA release were measured after incubation with medium without compounds, or containing: i) quercetin, ii) quercitrin, or iii) rutin. (at concentrations ranging between 1 and 150 µMCell). | Beginning at a 30 µM concentration, quercetin reduced cell proliferation and HA release, acting by the induction of necrosis and cell cycle blockade. No difference between GO and control fibroblasts was observed. | (16) Lisi S. et al |
| N-acetyl- cysteine, vitamin C and melatonine | Reduction of proliferation and release of cytokines in GO fibroblasts | Primary cultures of OFs from GO patients and control subjects. Oxidative stress was induced by H ₂ O ₂ . Cells were pretreated with N-acetylcysteine (100 and 200 μM) or vitamin C (250 and 500 μM). Cell proliferation, cell necrosis, apoptosis and HA release were measured | Treatment with H_2O_2 at low concentrations of H_2O_2 (3.125–25 μ M) increased the survival of GO fibroblasts. 6.25 μ M H_2O_2 led to a significant elevation of TGF β , IL-1 β and superoxide anion in GO fibroblasts, in GO, but not in control OFs. Pretreatment with N-acetylcysteine or vitamin C reversed the enhanced proliferation and the production of TGF- β , IL-1 β and superoxide anion of GO fibroblasts in response to 6.25 μ M H_2O_2 . | (24) Tsai CC. et al. |
| | Reduction of cell proliferation and HA release in GO fibroblasts | GSSG, as a measure of oxidative stress, cell proliferation, HA, TNF α , IFN γ , and IL-1 β were measured in primary cultures of GO and control OFs treated with H $_2$ O $_2$ and incubated with N-acetyl-cysteine, vitamin C and melatonine | Oxidative stress was reduced by all of the three antioxidant agents. Vitamin C reduced proliferation in GO, but not in control fibroblasts. N-acetyl-l-cysteine reduced proliferation and IFNy in GO, and HA and IL-1β in both GO and control fibroblasts. Melatonin reduced IL-1β and HA in GO and control fibroblasts, and IFNy only in GO fibroblasts. | (17) Rotondo Dottore G. et al |
| Enalapril | Reduction of cell proliferation and HA release in GO fibroblasts | Primary cultures of GO and control fibroblasts were treated with enalapril (2 or 5 mM, for 3 or 5 days) or with a control compound (lisinopril). Cell proliferation assays, lactate dehydrogenase release assays (as a measure of cell necrosis), apoptosis assays, and measurement of HA in the cell media were performed | Enalapril reduced OFs proliferation and HA release in the cell media in both GO and control fibroblasts. Because enalapril did not affect cell necrosis and apoptosis, the effects on proliferation probably reflected an inhibition of cell growth and/or a delay in cell cycle. | (15) Botta R. et al. |
| Retinol, β- carotene, vitamin E | Reduction of cell proliferation in GO fibroblasts | Primary cultures of GO and control fibroblasts. Oxidative stress was induced by treatment with H_2O_2 . Cells were pre-incubated for 2 days at 37°C with complete medium without compounds, or with medium containing either one of the compounds investigated at various concentrations. GSSG, cell proliferation, HA, TNF α , IFN γ , and IL1 β were measured | Retinol, β -carotene and vitamin E significantly reduced the release of GSSG and IL-1 β induced by H_2O_2 in GO, but not in control fibroblasts. β -carotene reduced OFs proliferation in GO, but not in control fibroblasts, whereas retinol and vitamin E had no effect. Retinol reduced IFN γ in GO and control fibroblasts. | (16) Rotondo Dottore G. et al |

OFs, orbital fibroblasts; GAG, glycosaminoglycans; HA, hyaluronic acid; H_2O_2 , hydrogen peroxide; TGF β , transforming growth factor β ; IL-1 β , interleukin-1 β , IFN γ , interferon γ , GSSG, Glutathione disulfide; TNF α , tumor necrosis factor α

The reduction of cell proliferation was likely due to a pronecrotic, rather than to a pro-apoptotic, effect of quercetin. These findings suggest that quercetin may have a therapeutic role in GO, but *in vivo* studies are needed to confirm this hypothesis, also considering that quercetin seems to act through necrosis, which may worsen autoimmunity against orbital antigens due to antigen exposure to the immune system.

Enalapril

Enalapril is a rather common anti-hypertensive medication, and it has been reported to inhibit fibroblasts proliferation and HA production in cheloid scars (59, 60). In view of these effects, a study was carried out using primary cultures of OFs from patients with GO and control subjects, which were treated with enalapril or with a control compound (lisinopril) (15). Cell proliferation, apoptosis, cell necrosis and HA release were measured. A significant reduction of OFs proliferation and HA release upon enalapril, but not lisinopril treatment, was observed in both GO and control OFs. Because enalapril did not cause an increase in necrosis or apoptosis, its inhibitory effects on proliferation were interpreted as the consequence of a direct action on cell growth and/or of a delay in cell cycle. No clinical studies are available on the use of this promising medication in GO.

Vitamin C, N-Acetyl-I-Cysteine, and Melatonin

Additional antioxidant agents, namely Vitamin C, N-acetyl-lcysteine and melatonin, were tested in another in vitro study performed in primary cultures of OFs from six GO patients and six control subjects (17). After treatment with H₂O₂ to induce oxidative stress, primary cultures were incubated with the various compounds. Glutathione disulfide (GSSG), as gage of oxidative stress, cell proliferation, HA, TNF-α, IFN-γ, and interleukin 1- β (IL1- β) were measured (17). H_2O_2 induced oxidative stress, as demonstrated by an increase in GSSG, and promoted cell proliferation and cytokine release, but did not affect HA. Vitamin C reduced proliferation in GO, but not in control fibroblasts. N-acetyl-l-cysteine reduced fibroblasts proliferation and release of IFN-y, a cytokine involved in the pathogenesis of GO, in OFs from GO patients, and it was also able to inhibit HA and IL1-β secretion in both GO and control OFs. Melatonin reduced IL1-β and HA release in GO and control OFs, whereas its inhibitory effect on IFN-γ release was significant only in GO OFs (17). Overall, the results of this study support a beneficial role of Vitamin C and N-acetyl-L-cysteine, suggesting a possible clinical use of these compounds. Furthermore, this study showed, for the first time, a beneficial effect of melatonin in terms of cytokines and HA synthesis. However, based on the lack of an inhibitory effect on fibroblasts proliferation, melatonin does not seem to be a strong candidate for a clinical use.

β-carotene, Retinol, and Vitamin E

Among other bio-available substances with antioxidant effects, β -carotene seems to be rather promising for a use in GO patients. In a study performed in our laboratory (18), again using primary

cultures of OFs from GO patients and control subjects, cells were pre-incubated with medium containing β-carotene, retinol or Vitamin E, or with medium without compounds, following induction of oxidative stress with H₂O₂. As in previous studies (17), oxidative stress promoted fibroblast proliferation, but it did not affect HA release. All of the three substances were able to reduce oxidative stress in GO, but not in control OFs (18). Cell proliferation, HA synthesis, as well as of pro-inflammatory cytokines were measured. Cell proliferation was reduced only by β -carotene. No effects in terms of HA synthesis were observed, whereas all of the cytokines measured, namely IL-1β, IFN-γ and TNF-α, which were increased by H₂O₂, were reduced by β -carotene as well as by retinol and Vitamin E (18). We can speculate that Vitamin E and retinol might not have an effect on GO in vivo, because they do not affect cell proliferation. On the contrary, \(\beta \)-carotene displayed antiproliferative effects in addition to an antioxidant and anti-inflammatory action, making it the most suitable candidate, among the three compounds studied, for a clinical use. Clearly, clinical studies are needed.

CONCLUSIONS

The balance of the cell redox state is a key point in cellular homeostasis. The disruption of the balance between oxidative and reductive activities in the cells leads to oxidative stress, by an increase in ROS production and a reduction of their scavenging. ROS interfere with intracellular reactions, thereby damaging various cellular components. Oxidative stress is involved in a number of diseases and seems to play an important role in GO. The management of patients with mild GO represents a challenge for endocrinologists and ophthalmologists. In addition to selenium, studies in vitro on the effect of other antioxidant agents provided, at least in part, interesting results which, unfortunately, have not been yet completely confirmed by clinical studies. Nicotinamide and allopurinol seem to be effective in reducing the inflammation of soft tissue but the data sustaining this conclusion were obtained in a rather small sample of GO patients affected with both moderate-to-severe and mild GO (42). The in vitro results on the use of pentoxifylline are promising, but clinical studies provided conflicting results (43, 44). Among the other compounds, because of the in vitro findings reported above, one can speculate that quercetin and β-carotene might be the most promising, but clinical trials are needed to confirm this hypothesis.

AUTHOR CONTRIBUTIONS

GL wrote the manuscript. CM and MM contributed to the conception of the manuscript and revised the paper critically. All authors contributed to the article and approved the submitted version.

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Asymmetric Graves' Orbitopathy

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Graves' Orbitopathy (GO) is an autoimmune orbital disorder usually presenting as a sequala of autoimmune thyroid disease. The presence of GO is associated with increased psychological burden and, in severe cases may cause blindness. While most patients with GO present with bilateral disease, asymmetric or unilateral GO may affect a significant proportion of patients diagnosed with GO. Older age, male sex, active and severe disease correlate with asymmetric disease. However, the exact mechanisms causing asymmetry remain elusive. Herein, we review the literature on asymmetric GO and highlight its differences compared with bilateral GO.

Keywords: asymmetry, Graves' disease, exophthalmos, unilateral, hyperthyroidism

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INTRODUCTION

Graves' orbitopathy (GO) is the most common extrathyroidal feature of Graves' disease (1) with an estimated prevalence of 10/10,000 persons in European populations (2), while more recent data have shown variable prevalence between 25% and over 50% in Graves' hyperthyroidism cases (3) worldwide. GO is closely associated with thyroid autoimmunity, and although it is classically linked with hyperthyroidism, GO features have been described in hypothyroid or euthyroid individuals (4).

Autoimmune processes resulting in proliferation of orbital fibroblasts, increased adipogenesis, and extracellular matrix expansion are involved in its pathophysiology (5) and Thyroid Stimulating Hormone (TSH) Receptor Antibodies (TRAb) seem to be a key determinant. In fact, TRAb titer together with smoking, duration of thyroid dysfunction and clinical activity score (CAS) at baseline are regarded as main risk factors for developing GO (6).

The disease most often presents bilaterally and symmetrically with lid retraction, exophthalmos. and diplopia (3) being the most common features. In severe cases GO can be sight-threatening, thus requiring a prompt referral to specialist services (7) upon diagnosis. GO may also cause a psychological and financial burden (8), primarily but not exclusively due to cosmetic concerns, as well as increase the risk for suicide (9), and impairment of quality of life of patients (10).

However, some patients might exhibit asymmetric or unilateral symptoms, for yet unknown reasons. It has been suggested that unilateral GO may progress to bilateral disease (11) and we have also found that asymmetry was associated with more severe and active GO (12). Asymmetry and/or unilateral GO can impose diagnostic challenges (13) and it is therefore important that awareness for asymmetric GO be raised. Clinicians involved in the care of patients with GO need to be able to identify unilateral or asymmetric GO, as timely recognition of asymmetry may expedite referral to specialist services and facilitate further management. However, whether unilateral and/or asymmetric disease represent a distinct variant of GO remains unclear and a better understanding of their underlying mechanisms may provide useful insight on the pathogenesis of

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GO, in general. Herein, we review the available literature on asymmetric GO and highlight differences compared with bilateral disease.

ASSESSMENT OF ASYMMETRIC AND/OR UNILATERAL GO

Asymmetry of the human face is well documented (14, 15), while asymmetry in the orbital anatomy of normal human skulls has been shown to be the norm. In a study of 127 human skulls (254 orbits) of individuals aged between newborn to 76 years, asymmetry in orbital anatomy (greater horizontal diameter, greater vertical diameter, orbital perimeter and orbital base area) was present in all cases except four, with variability between 2.47 and 4.47% between the right and left orbit (16). In the same study asymmetry was more prevalent in females and measurements from the right orbit were greater than the left (16). This is of importance, as facial asymmetry in itself can be a cause of significant distress (17).

On the other hand, orbital volume calculation using CT scanning has shown no differences between orbits or gender in a Taiwanese normal population (18). A large study of 653 normal Caucasians subjects aged 21–80 years found asymmetry to be rare (2%), minor (difference in exophthalmos readings <2mm), and unrelated to gender or age. Another study however showed a linear negative correlation between proptosis and age between the ages of 31 and 80 (19), which was also confirmed independently (20). In the latter study, recruiting a large sample (n = 1,063) of normal Iranian subjects including children, a significantly greater proptosis in the right eye compared with the left via exophthalmometry was shown, though the difference was never greater than 2 mm (21).

In the past, asymmetry has been variably defined as difference in proptosis between eyes by $\geq 2 \text{ mm } (13)$, > 2 mm (11, 22), or any one of the following criteria: retrobulbar pain or > or =1 grade in soft tissue involvement, and/or of > or =2 mm in exophthalmos, and/or > or =8 degrees in elevation (23), or repeatable asymmetry with regard to more than one symptom and more than one external or anterior segment finding for a duration of two or more visits at least 1 month apart at any time during the initial or follow-up period (23). Radiological criteria for asymmetry, such a right-to-left ratio of more than 1.4 in extraocular muscle diameter, as obtained by CT measurements based on normative data, have also been proposed (24). Unilateral GO is likewise variably defined as one or more features in one eye without any such manifestations in the other eye (20), proptosis >2 mm in one eye with normal examination of the other eye (25), or proptosis difference between eyes >4 mm, and/or if clinical signs and symptoms of GO found unilaterally (26). In our recently published study, asymmetry was defined as bilateral disease with one or more of the following features: difference between the two eyes in exophthalmos by ≥2 mm; difference in palpebral aperture by

 \geq 2 mm; difference in eyelid swelling by \geq 1 grades; difference in eyelid erythema by \geq 1 grades; difference in conjunctival redness by \geq 1 grades; presence of dysthyroid optic neuropathy in one eye only. Unilateral disease was defined as one or more clinical features of GO in one eye without any evidence of GO in the contralateral eye (12). Moreover, disease severity is assessed similarly to bilateral disease, using clinically evaluated standardized tools, such as CAS (27) and Vision, Inflammation, Strabismus, Appearance (VISA) (28). Clearly, there is a need for a consensus in the definitions of asymmetric and unilateral disease.

EPIDEMIOLOGY OF ASYMMETRIC AND/OR UNILATERAL GO

Different studies have estimated prevalence of unilateral disease between 4.5 and 14% (13, 25, 29-31), while asymmetry was evident in 9-34% of patients with GO (13, 26, 32). More recently, in our multi-center prospective cohort recruiting 269 newly diagnosed patients from 13 different centers across Europe, we have found a prevalence of 30.9% for asymmetric and 10.7% for unilateral GO (12). However, in a recently published retrospective hospital-based report from India, unilateral disease was much higher at 36% and significantly more common among silent presenters compared to the clinically active group (33). It is important to note that published studies have used different definitions of asymmetry and/or unilateral GO and this might explain the big variation in reported epidemiology among published data. Furthermore, less is known in regards to the epidemiology of unilateral or asymmetric GO in children.

Many risks factors have been associated with asymmetry. Shorter duration of symptoms (29) and thyroid status might be associated with asymmetry. In specific, small studies have suggested that euthyroid and primarily hypothyroid patients develop more asymmetrical GO (34, 35), which tends to present more mildly. In a larger study, Ponto et al. showed that a sevenfold higher risk for unilateral GO in hypothyroid or euthyroid subjects, compared with hyperthyroid GO (26). Moreover, previous studies have shown no difference in regards to race (13) and sex (29), but more recent data, including our recent multicenter study (12) and a cohort of 354 Chinese patients (36), have shown that male sex is associated with asymmetry. In another study, male subjects exhibited asymmetric disease (proptosis and overall asymmetry) three-fold more frequently compared with women, while women with hyperthyroidism demonstrated more symmetry compared with euthyroid and hypothyroid individuals of either sexes (23). With regards to correlation with older age, we have confirmed previous findings supporting higher prevalence and more severe asymmetry in older and especially men (12, 37), in keeping with previously described associations between age and male gender with severity of GO in patients with bilateral disease (38).

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SPECIFIC CONSIDERATIONS IN ASYMMETRIC GO

It is important that unilateral GO be differentially diagnosed from orbital diseases affecting one eye, such as orbital tumors, such as lymphoma (39, 40). Other differentials affecting the orbit including other orbital tumors or pseudotumors, carotid cavernous fistulae, and dermoid and/or epidermoid cysts need to be excluded, too. Diagnosing unilateral or asymmetric GO without any other signs of GO and especially in the absence of hyperthyroidism or high TRAb titer suggestive of Graves' disease can be challenging (41) and requires increased clinical awareness and usually imaging of the orbits.

With regards to pathophysiology, the autoimmune processes in asymmetric and unilateral GO, causing expansion of orbital contents seem to be similarly to bilateral disease. However, mechanical, vascular, and inflammatory factors as well as anatomical variances may contribute to development of asymmetric disease.

More specifically, Soroudi et al. have speculated that asymmetric distribution of antigen or inflammatory processes may be the cause of asymmetrical expansion of orbital contents (13), though this was not explored in any studies so far. They have also postulated, that anatomical differences causing differential blood flow or lymphatic drainage may be present (13). Elasticity of orbital septae or other local factors, associated with unilateral triggers such as infections or difference in potential for adipogenesis have also been suggested (12). Others have examined the effect of sleeping position but found no significant correlation with asymmetric GO (32). However, despite previous postulations the exact mechanisms remain elusive. Therefore, more studies are needed to study asymmetric GO and shed light on the mechanisms leading to asymmetry. Such studies may provide further answers in regards to GO development and management.

TREATMENT OF UNILATERAL AND/OR ASYMMETRIC GO

Given that the pathophysiological mechanisms causing unilateral or asymmetric GO appear to be similar to the well-described bilateral GO, treatment of asymmetric GO is generally the same with bilateral disease and is usually dependent on disease severity. However, since asymmetric disease may progress to bilateral GO thereby increasing patients' anxiety and deteriorating their mental health status and quality of life as alluded to above, it is important that patients are managed promptly. Therefore, the goal in managing asymmetric or unilateral GO should be: a) to ensure symptoms are alleviated and sight is not threatened, similarly to bilateral GO; b) to take measures that may prevent progression to bilateral symptomatology; c) surgical rehabilitation when indicated.

Smoking cessation advice and restoring the euthyroid state are important pillars in the management of asymmetric GO. There is no contraindication in regards to any treatment modality including antithyroid drugs, radioactive iodine, and/ or total thyroidectomy for mild asymmetric and/or unilateral GO, while in moderate to severe cases radioiodine should be avoided, similarly to bilateral symmetric disease (42).

In mild cases of asymmetric GO patients may still benefit from selenium supplementation (43). However, there are no prospective data evaluating whether early selenium supplementation in asymmetric GO cases may halt progression to bilateral symptoms. For more severe cases, methylprednisolone infusion (44) and/or administration of immune-modifying therapies, such as rituximab (45) and/or mycophenolate mofetil (46) may need to be considered. Targeted therapies such as teprotumumab, an anti-Insulin Growth Factor 1 (IGF-1) receptor monoclonal antibody, were shown to be effective in patients with active GO (47), but its role in asymmetric or unilateral GO is unexplored. Decompression surgery may be used in appropriate cases.

DISCUSSION AND CONCLUSIONS

Asymmetric and unilateral GO are recognized features of GO with variable prevalence among different studies; however, they do not seem to represent a distinct variant of classical GO, rather than the extreme of the spectrum, and thus focused studies are scarce. Therefore, prospective multicenter studies recruiting patients with different socioeconomic, demographic, ethnic, and anthropometric background are needed to better elucidate the epidemiology as well as other parameters of asymmetry and/ or unilateral disease. Furthermore, previous studies have shown associations of asymmetric/unilateral GO with disease activity and severity and it is crucial that large cohorts include patients of different clinical status. There is a need for better documentation of the suggestions that asymmetric/unilateral GO might run a milder course or be a prelude to bilateral disease. The mechanisms leading to asymmetric disease discussed are speculative. Animal and mechanistic studies are needed to provide more in depth understanding of the mechanisms leading to disease progression and bilaterality and could identify possible novel therapeutic targets at a molecular level. Wide agreement and consensus among experts on the definition of asymmetry and unilaterality is paramount and will facilitate diagnosis, management, and research into this fascinating clinical entity.

In conclusion, although the available literature is limited, asymmetric and/or unilateral GO tend to be present in older age and male patients and is associated with more active and severe GO. Current evidence suggests that patient presenting with asymmetric or unilateral GO may progress to bilateral disease, which clinicians treating patients with GO need to be aware of. The present mini-review summarizes important information for both clinicians and researchers and also provides the impetus for further research. More specifically, in everyday clinical practice, unilateral disease needs to be differentiated from other pathologies affecting one eye. In the future, mechanistic studies

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are needed to explore the underlying pathological processes in asymmetric/unilateral GO. Most importantly, longitudinal studies evaluating individuals who are at higher risk for developing bilateral symptoms as well as the effect of various treatments in impeding progression of unilateral and/or asymmetric disease to bilateral GO are warranted.

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AUTHOR CONTRIBUTIONS

GP: literature search, data acquisition, and writing of the manuscript. PP: conceptualization, literature search, and writing of the manuscript. All authors contributed to the article and approved the submitted version.

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Conflict of Interest: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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Teprotumumab as a Novel Therapy for Thyroid-Associated Ophthalmopathy

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Thyroid-associated ophthalmopathy (TAO) has remained a vexing and poorly managed autoimmune component of Graves' disease where the tissues surrounding the eye and in the upper face become inflamed and undergo remodeling. This leads to substantial facial disfigurement while in its most severe forms, TAO can threaten eye sight. In this brief paper, I review some of the background investigation that has led to development of teprotumumab as the first and only US FDA approved medical therapy for TAO. This novel treatment was predicated on recognition that the insulin-like growth factor I receptor plays an important role in the pathogenesis of TAO. It is possible that a similar involvement of that receptor in other autoimmune disease may lead to additional indications for this and alternative insulin-like growth factor I receptor-inhibiting strategies.

Keywords: autoimmune, insulin-like growth factor I receptor, monoclonal antibody, Graves' disease, thyroid-associated ophthalmopathy, connective tissue

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INTRODUCTION

Thyroid-associated ophthalmopathy (TAO), also known as thyroid eye disease or Graves' orbitopathy, remains a vexing disease process most frequently occurring in individuals with Graves' disease (GD) (1). Less commonly, TAO can occur in patients with Hashimoto's thyroiditis. Autoimmunity similar to that found in the thyroid is presumed to underlie TAO, although responses to the specific autoantigen(s) in the two tissues may not be identical. The medical treatment of TAO has been woefully inadequate, in large part as a consequence of our poor understanding of its pathogenesis. Thus, until very recently, no drug had been approved by the U.S. Food and Drug Administration (FDA) for TAO. TAO can present with a variety of physical signs and symptoms, many of which are shared with other more common diseases of the tissues surrounding the eye. In this brief review, I attempt to provide the historical background underpinning efforts to identify an effective and safe medical therapy for TAO. That direction of study has yielded substantial evidence for meaningful involvement of the insulin-like growth factor-I receptor (IGF-IR) in the development of TAO (2). At the heart of this evidence is the overexpression of IGF-IR by orbital fibroblasts, T and B cells (3-5), the generation of autoantibodies targeting the receptor in patients with GD (3, 6), and the apparent physical and functional collaboration between IGF-IR and the thyrotropin receptor (TSHR) (7). Based nearly entirely on a series of studies conducted in vitro, a β arrestin-biased agonist that acts as an IGF-IR inhibitor, teprotumumab, now marketed as Tepezza, was repurposed from its initially intended clinical use as

an anti-neoplastic agent, to its consideration as a medical therapy for TAO (8). On the strength of two successful clinical trials involving patients with active, moderate to severe disease, teprotumumab has recently been approved by the FDA for use in TAO (9). This approval has therefore ushered in to clinical practice a new era for medically managing this serious and historically underserved manifestation of thyroid autoimmunity.

CURRENT UNDERSTANDING OF TAO PATHOGENESIS

A number of key insights have been generated in the recent past by several laboratory groups across North America and abroad. At the heart of TAO is the growing and sometimes reluctant recognition that orbital fibroblasts from diseased orbits (GD-OF) comprise a heterogeneous cell population and are involved in pathogenesis as a consequence of the unique presence of CD34⁺ cells where CD34 indicates a cell phenotype with specific characteristics (10–14). These cells have been proposed as the dominant mediators of TAO development by virtue of their extraordinary responsiveness to inflammatory mediators such as cytokines and growth factors (15–19) (**Figure 1**). They express key promoters of inflammation including both prostaglandin endoperoxide H synthase 1 and 2, the latter of which is highly inducible (15, 16, 20, 21) and 15-lipoxygenase (22). We contend that the identification of CD34⁺CXCR4⁺Col I⁺ fibroblasts in the

TAO orbit as a discrete subset of GD-OF and their putative derivation from circulating fibrocytes represents a plausible explanation for the markedly heterogeneous behavior found among these disease-derived fibroblasts (14). The aggregate of these markers identifies fibrocytes and distinguishes them from fibroblasts and other cell types (23). These cells can undergo adipogenic differentiation (24). The Thy-1 subset of GD-OF is particularly susceptible to pro-adipogenic factors while those of the Thy-1⁺ phenotype can undergo differentiation into myofibroblasts through the actions of TGF-β and the activation of the Smad pathway (25, 26). CD34+ OF exhibit unique phenotypic attributes which can be attributed, at least in part, to their promiscuous expression of the autoimmune regulator protein (27). They synthesize several proteins, the expressions of which were previously thought to be restricted to the thyroid, including TSHR, thyroglobulin, thyroperoxidase, and sodium iodide symporter (28, 29). But the expression of tissue-specific proteins that behave as autoantigens is not limited to those relevant to the thyroid. Fibrocytes have also been found to express those protein antigens implicated in type 1 diabetes mellitus (29). Fibrocytes cross-talk with T cells, and especially with those polarized to the Th17 paradigm, the cytokines from which appear to play important roles in driving orbital inflammation in TAO (30). IL-17A promotes the expression of regulated on activation, normal T cell expressed and secreted (RANTES, CCL5) by orbital fibroblasts, effects mediated through the CD40/CD154 (CD40 ligand) bridge (31).

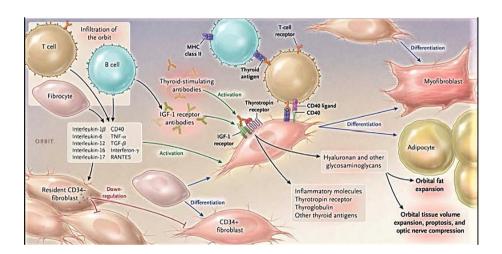


FIGURE 1 | Cartoon proposed model of thyroid-associated ophthalmopathy (TAO) pathogenesis. At the center of the disease are orbital fibroblasts which exhibit particularly robust responses to inflammatory mediators. Among them are CD34+ cells which we propose derive from fibrocytes, monocyte-derived progenitor cells trafficked from bone marrow. Fibrocytes circulate in Graves' disease at higher frequency than that found in healthy individuals. When cultivated from the peripheral circulation, fibrocytes express several thyroid-specific proteins, including thyrotropin receptor (TSHR), thyroglobulin, thyroperoxidase, and sodium-iodide symporter. They also express Class II major histocompatibility complex (MHC) when unstimulated by interferon γ and can present antigens. When exposed to the appropriate culture conditions, they undergo differentiation into myofibroblasts (through Smad pathway activation by TGF- β) and adipocytes (through the activation of PPAR- γ). Many of the genes expressed by fibrocytes are detected at considerably lower levels in CD34+ orbital fibroblasts. We have found that these lower levels of expression result from the actions of Slit2. When activated, CD34+ fibrocytes and CD34+ fibroblasts generate several pro-inflammatory or anti-inflammatory cytokines, including interleukins 1 β , 6, 8, 10, 12, 16, tumor necrosis factor α , and regulated on activation, normal T expressed and secreted (RANTES), CXCL-12 and CD40-CD154. Both CD34+ and CD34+ orbital fibroblasts cell-surface display insulin-like growth factor-I receptor (IGF-IR). Orbital fibroblasts express three mammalian hyaluronan synthase isoenzymes and UDP glucose dehydrogenase and synthesize hyaluronan, the glycosaminoglycan associated with expanding orbital tissue in TAO. The vast majority of hyaluronan synthesis occurs in CD34+ orbital fibroblasts. From N. Engl. J. Med, Smith T.J. and Hegedus L., Graves' Disease, 375; 1552-1565. Copyright $^{\odot}$ (2016) Massachusetts Medical Society. Reprinted with permission.

GD-OF and fibrocytes have been found to express TSHR (14, 32). This receptor was subsequently found to be functional and could mediate effects of both TSH and thyroid-stimulating autoantibodies (7, 14). Activation of several genes, including those encoding cytokines implicated in the pathogenesis of TAO, has been reported (19, 33–35). Fibrocytes are trafficked to sites of tissue injury through several chemokine networks, most notably the CXCL-12/CXCR4 pathway (36) which is under the control of TSHR signaling. A potentially important molecular conduit, the activation of which can influence gene expression in GD-OF, is the CD40–CD154 bridge (21, 37). Ligation of CD40 displayed on the surface of these cells results in substantial upregulation of hyaluronan synthesis and the generation of prostaglandin E₂ through the induction of prostaglandin endoperoxide H synthase 2.

TAO HAS LONG BEEN A DISEASE THE TREATMENT FOR WHICH EXEMPLIFIES INADEQUACY

Despite its initial published descriptions, dating to the 19th century (38), precious little progress had been made in identifying safe and effective medical therapy for TAO (39, 40). Of the available medications, glucocorticoids have been the most widely used. Standardization of their use, including indications, exclusions, treatment duration, and dosages has yet to be established with placebo-controlled, prospective, and adequately powered clinical trials. Some clinicians taking care of patients with TAO extoll them as the "gold standard", but the basis for this often strongly expressed opinion appears to rest on dubious grounds rather than on firm scientific evidence. Both oral and intravenous routes of steroid administration have been advocated. More recent reports suggest that IV pulse may be more effective and associated with fewer side effects than those delivered by the oral route (41). But even staunch advocates of steroid use in TAO typically concede that these agents fail to consistently provide benefit to patients and that only 50% respond meaningfully (42). Further, many experts acknowledge that responses to steroids are essentially limited to amelioration of inflammatory signs and symptoms. Among the most informative reports of steroid use in active, moderate to severe TAO, was that of Bartalena and colleagues (43). They examined three different cumulative dosages (2.25, 4.98, and 7.47 g) administered as 12 weekly infusions. Their study demonstrated that short-term response, as measured by improvement in the clinical activity score (CAS), was greatest in those receiving the highest dosage. Patients receiving 7.47 g also showed a 0.6 mm proptosis reduction at 12 weeks compared to baseline, a result widely considered to be lacking clinical importance. Evidence suggests that glucocorticoids in combination with external beam radiation may be somewhat superior to steroids alone (44-47). Other agents have been proposed for combination with steroids, including rituximab, mycophenolate and azathioprine (48-50). In my view, whether used as a single agent or in combination with radiotherapy, systemic steroids should play a progressively

less frequent role in the treatment of TAO. In distinction, locally administered steroids, such as those delivered in eye drops or intraorbital injections, may have some utility, especially in mild but symptomatic disease. Steroid therapy in TAO poses substantial risk. Given their limited efficacy, these agents are deemed unsafe since they are associated with several potentially serious side-effects, including hypertension, bone loss, psychiatric illness, and peptic ulcer disease. Moreover, highdosage intravenous steroids can deleteriously affect liver function and in rare cases, result in fatal hepatic failure (48, 51, 52). Having articulated these objections to steroid use, several biological agents are currently under development or already have been repurposed for potential use in severe TAO. Among them tocilizumab, an IL-6 receptor antagonist, has been subjected to a placebo controlled trial of 32 patients with steroid-resistant TAO at 10 performance sites in Spain (53). Drug or placebo was administered at weeks 0, 4, 8, and 12. Of those receiving active drug, 93.3% met the primary response of a ≥2 point reduction in CAS at week 16 while 58.8% of those treated with placebo met the response (p = 0.04). Notably, a 1.5 mm reduction in proptosis was detected in those receiving the active drug versus 0.0 mm in the placebo group at week 16. Observations made later in the study revealed that the small change in proptosis seen with tocilizumab was not durable. A clinical trial of belimumab is currently underway for TAO (54). In addition, preliminary studies examined the potential for immune retolerizing of TSHR in GD and TAO (55).

IGF-IR AND ITS SIGNALING PATHWAY AS THERAPEUTICALLY EXPLOITABLE TARGETS

The IGF-I pathway is complex, comprising several molecules that either enhance signaling initiated by IGF-I or play inhibitory/modulatory roles. IGF-I influences regulation of both normal and abnormal development, metabolism/energy expenditure and immune surveillance (56, 57). It is generated either by the liver or in peripheral tissues. In the latter, it can act locally. IGF binding proteins (IGF-IBP), of which six have been identified, can enhance the half-life of IGF-I within the circulation (58) and can modulate the interactions between IGF-I and IGF-IR (59). The actions of IGF-I and IGF-II are mediated through their interactions with the two cell surface receptors, IGF-IR and IGF-IIR/mannose-6-phosphate receptor (60). IGF-IR is a member of the insulin receptor (IR) family of cell surface-displayed tyrosine kinases. We had suggested some time ago that this pathway might be exploitable for the therapeutic targeting of autoimmune diseases, including GD and TAO (61). The first clue that the IGF-IR pathway might be involved in TAO emanated from the laboratory of Kendall-Taylor nearly three decades ago (62). They reported that IgGs found in sera of patients with TAO could compete for binding of radiolabeled IGF-I to the surface of orbital fibroblasts collected from patients with the disease. While the identity of the binding site was surmised to be IGF-IR by that group, results in a later

study conducted by those in another laboratory definitively demonstrated that IGF-IR harbored the binding site and therefore the epitope for autoantibodies circulating in TAO (3). These later studies indicated that IgGs from patients with GD (GD-IgG) could initiate signaling and that the PI3 kinase/FRAP/mTor/p70^{S6k} and Erk p42/44 pathways become activated in GD-OF (3, 6) and thyroid epithelial cells (63). Those studies failed to examine whether GD-IgGs could initiate autophosphorylation of tyrosine residues intrinsic to IGF-IR. Their findings did however raise the possibility that GD-IgG was initiating signaling directly through interactions with IGF-IR. Subsequently, several laboratory groups reexamined this question that GD-IgGs activate IGF-IR autophosphorylation. One group reported that GD-IgGs from a subset of patients could activate receptor kinase activity above levels seen with control (healthy) IgGs (64). In contrast, other investigators have failed to detect GD-IgG-stimulation of IGF-IR kinase activity (65, 66). The concept of anti-IGF-IR antibodies playing a role in TAO has been vigorously argued against by some investigators [summarized in ref (2)]. Their viewpoints have relied heavily on selective reviews of the literature rather than on comprehensive consideration and reconciliation of opposing opinions. The discrepancy between the findings generated by several laboratory groups has yet to be explained satisfactorily. It should be noted that the methods used for detecting IGF-IR activity have varied widely, experimental conditions have not been standardized, and cell treatment protocols have generally relied on single time points, potentially inappropriate target cells, and the uniform absence of assessing endogenous IGF-I concentrations in the culture media. GD-IgG was found to induce the synthesis of hyaluronan in orbital fibroblasts from patients with TAO but not in orbital fibroblasts from healthy individuals (67). Those effects could be mimicked by recombinant IGF-I but not by recombinant human TSH. Subsequent to the detection of anti-IGF-IR autoantibodies in GD and TAO, Tsui et al. reported that IGF-IR and TSHR colocalize in thyroid epithelial cells, orbital fibroblasts and in situ in orbital fat (7). Further, in pull-down studies, these investigators found that monoclonal antibodies directed against either receptor protein could precipitate both. They also found that the IGF-IR inhibitory monoclonal antibody, 1H7, could attenuate the activation of Erk 42/44 MAPK initiated by rhTSH, GD-IgG and IGF-I. Those results suggested that signaling emanating from either receptor depended on the activation of IGF-IR. Another report demonstrated that the cellular distribution of IGF-IR was altered in fibroblasts derived from TSHR-null mice (68). A similar pattern of response to disease-derived IgGs was subsequently observed in synovial fibroblasts from patients with rheumatoid arthritis (69). Based on the aggregate of findings implicating IGF-IR in the pathogenesis of TAO, the receptor became a plausible therapeutic target. Further it was suggested that multiple autoimmune diseases, besides GD and rheumatoid arthritis, might share IGF-IR-dependent disease mechanisms and that therapeutic development could be successfully focused on that receptor (61).

TWO CLINICAL TRIALS OF TEPROTUMUMAB REVEAL AN EFFECTIVE AND SAFE THERAPY FOR MODERATE TO SEVERE, ACTIVE TAO

Phase 2 Trial

Based on the preclinical studies outlined above, which demonstrated the plausibility of IGF-IR inhibition as a potential treatment strategy for GD and TAO, an initial trial was organized by River Vision Development Company beginning in 2010. The phase 2 study of teprotumumab (RV001, R1507), a fully human monoclonal β -arrestin biased agonist IGF-IR inhibitor, involved the repurposing of that drug from its initially intended use in cancer where it had proven ineffective in treating several disease types (70, 71). This trial was a double-masked, randomized, placebo-controlled and multicentered study with 15 performance sites in North America and Europe (72). Enrolment commenced on 2 July, 2013 and was completed 23 September, 2015. Exclusionary factors included uncontrolled hyper- and hypothyroidism, a history of receiving >1 g of steroids for the treatment of TAO, prior rehabilitative ocular surgeries for TAO and prior treatment with either rituximab or tocilizumab. A total of 88 patients, age range from 18 to 75 years, with moderate to severe, active [clinical activity scores (CAS) ≥4 on a 7-point scale] TAO were randomized to receive either teprotumumab or placebo in a 1:1 ratio. Participants were uniformly within 9 months of the onset of TAO, included both cigarette smokers and nonsmokers, and were clinically euthyroid. Once randomized, patients were included in the intention-to-treat (ITT) cohort. Participants in both treatment groups received dosing every 3 weeks for a total of eight infusions over a 24-week intervention phase. The primary response was defined as improvement in the study (worse) eye of both, improvement in CAS of ≥2 points, AND reduction in proptosis of ≥ 2 mm. This improvement must have occurred in the absence of a similar degree of worsening in the fellow eye. Secondary endpoints, measured as independent variables, included improvement from baseline of CAS of ≥2 points, reduction in proptosis of ≥2 mm, improvement in subjective diplopia, and improved results from a fully validated quality of life questionnaire (GO-QoL). The occurrence and severity of adverse events were also assessed. Any subjects failing to complete the 24-week treatment phase or the assessment at week 24 for any reason were considered treatment failures.

The responses used to judge drug efficacy included reduction in CAS, proptosis, improvement in diplopia, and GO-QoL. These parameters were comparable at the baseline assessment in the two treatment arms. Despite attempts to stratify participants with regard to cigarette smoking, an imbalance in group assignment was detected, with a great number who used tobacco randomized to the placebo group. Analysis of the results demonstrated that of those in the ITT group receiving teprotumumab, 29 of 42 (69%) exhibited a response at 24 weeks. In contrast, nine of 45 (20%) patients receiving placebo

responded (p < 0.001). Similar numbers of participants in the two treatment arms completed the intervention phase (87% in the placebo and 88% in the teprotumumab cohorts). The time to response was considerably shorter in the teprotumumab cohort than in those receiving placebo. The onset of response was very rapid. Nearly one-half of those receiving teprotumumab achieved the primary response at week 6, i.e. those who had received the half-dose and a single full-dose. The proportion of patients achieving a response was greater in the teprotumumab group and continued to increase throughout the duration of the treatment phase with striking differences compared to placebo which continued to be significant at every time point (all p < 0.001). When the responses were graded, considerably more patients exhibited a high response at week 24 (≥3 point improvement in CAS and ≥3 mm proptosis reduction in the study eye). With regard to the secondary outcomes, those patients receiving teprotumumab responded with a change in proptosis

from baseline and CAS from baseline when compared to those receiving placebo. Both responses were significantly different when compared to controls at the 6-week time point (p < 0.001 for both), and the differences continued to increase throughout the treatment phase which ended at week 24 (both p < 0.001). On the basis of this trial, the FDA designated teprotumumab a "breakthrough" therapy for TAO.

Phase 3 Trial

A pivotal phase 3 study was initiated shortly after the results of the phase 2 trial had been reported (73). The study was organized and funded by Horizon Pharmaceutical (now Horizon Therapeutics) and included many of the same investigators and institutions participating in the preceding phase 2 trial. It was conducted between October 24, 2017 and August 31, 2018, when 107 patients were screened and 83 underwent randomization (**Figure 2A**). The trial followed an experimental

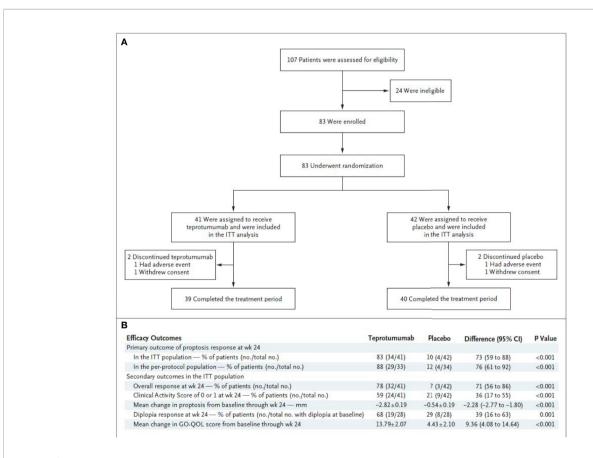


FIGURE 2 | (A) Trial profile. (B) Efficacy endpoints. * CMH weighting was used to estimate the common risk difference and the 95% Confidence Interval (95% CI) of the common risk difference for the primary and secondary endpoints of overall responder, percent with CAS 0 or 1, and diplopia responder; least squares mean difference was calculated for secondary endpoints of change in proptosis from baseline and change in Graves' Orbitopathy quality of life (GO-QOL) questionnaire from baseline using the Mixed-Model Repeated-Measures (MMRM) analysis of covariance (ANCOVA) model described below. * 28 patients in each treatment group had diplopia at baseline * GO-QOL score was calculated and transformed to a 0 to 100 scale. Transformed score = [(sum of each score – number of completed items)/(2 * number of completed items)] * 100 \$ Change from baseline in proptosis or GO-QOL as a continuous variable is based on MMRM ANCOVA model with an unstructured covariance matrix including the following terms: baseline score, tobacco use status (non-user, user), treatment group, visit, and visit-by-treatment and visit-by-baseline-score interactions; data presented as least squares mean ± standard error. All analyses were controlled for multiplicity. From N. Engl. J. Med, Douglas R.S., Kahaly G.J., Patel A., Sile E.H.Z., Thompson R et al. Teprotumumab for the treatment of active thyroid eye disease. 382; 341-352. Copyright © (2020) Massachusetts Medical Society. Reprinted with permission.

protocol that was very similar to that of the earlier study. It allowed a widened patient age limit (18 years to 80 years) and the inclusion of a study extension, which enabled treatment crossover for patients failing to respond during the 24-week treatment period and those who had responded initially and then relapsed. Individuals with a history of inflammatory bowel disease were excluded since at least one participating patient in the earlier trial with that diagnosis experienced worsening bowel symptoms. The remainder of the inclusion and exclusion criteria were unchanged from the phase 2 trial. The primary endpoint was proptosis responder rate (percentage of patients with ≥2 mm

reduction in the study eye without ≥ 2 mm increase in the fellow eye) at week 24 for teprotumumab *versus* placebo. Secondary outcomes included overall responder rate (the primary outcome of the phase 2 study) which was defined as the percentage of patients with both ≥ 2 -point reduction in CAS AND ≥ 2 mm reduction in proptosis in the study eye in the absence of a corresponding worsening in the fellow eye, percentage of patients with a CAS 0 or 1, and diplopia improvement at week 24 and change in GO-QOL overall score through week 24. Unlike the phase 2 trial, this study included an extension. Specifically, those patients who did not achieve the primary

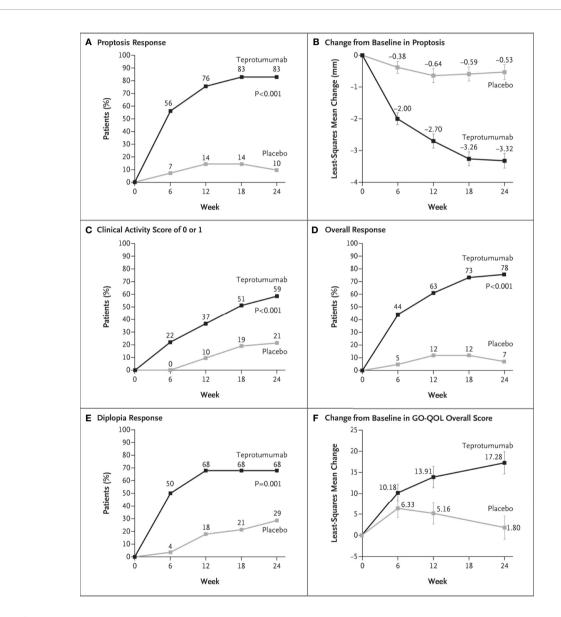


FIGURE 3 | (A) Proptosis responder analysis (percent of patients with ≥2 mm reduction in proptosis from baseline in the study eye). (B) Change from baseline in proptosis (least squares mean ± standard error). (C) Percent of patients with clinical activity score (CAS) of 0 or 1 in the study eye. (D) Overall responder rate (percent of patients with ≥2-point reduction in CAS and ≥2 mm reduction in proptosis from baseline in the study eye). (E) Diplopia response (percent of patients with improvement of at least one grade from baseline). (F) Change from baseline in transformed GO-QOL score (least squares mean ± standard error). From N. Engl. J. Med, Douglas R.S, Kahaly G.J., Patel A., Sile E.H.Z., Thompson R et al. Teprotumumab for the treatment of active thyroid eye disease. 382; 341-352. Copyright (2020) Massachusetts Medical Society. Reprinted with permission.

outcome *i.e.* the proptosis non-responders, were eligible for participation in an open-label extension study (OPTIC-X (NCT03461211)]. This trial involved eight additional infusions of teprotumumab. Those patients not included in the extension were followed for 48 weeks. Those patients who responded during the initial treatment phase, regardless of treatment arm but then relapsed during follow-up could also enter the extension study.

Efficacy outcomes are shown in **Figure 2B**. The results from the trial revealed that 9.5% of patients receiving placebo achieved the primary response compared with 82.9% of those treated with

teprotumumab at week 24 (delta 73.45%; 95% CI 58.89 to 88.01%; p < 0.001) (**Figure 3A**). A majority of patients responding did so at week 6. Mean proptosis response from baseline with teprotumumab at week 24 was -3.32 mm *versus* -0.53 in placebo or a mean difference from the placebo group of 2.79 mm (**Figure 3B**).

Secondary outcomes were achieved by significantly more patients receiving teprotumumab than those in the placebo cohort. CAS scores of 0 or 1 (**Figure 3C**) and over-all responders were also more numerous in the teprotumumab group at all study visits (**Figure 3D**). At baseline, both treatment groups included 28 patients with diplopia;

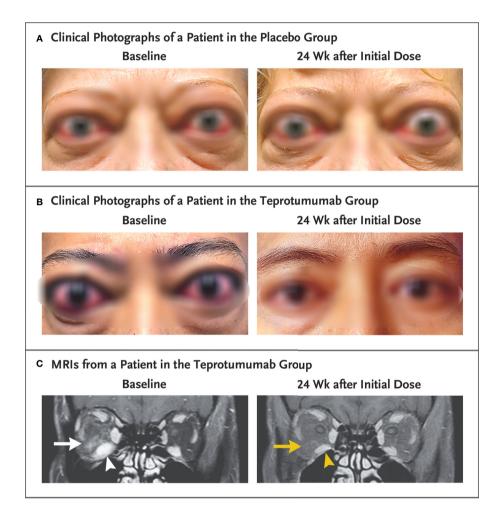


FIGURE 4 | Facial photographic images and MRIs at Baseline and 24 Weeks following treatment with either placebo or teprotumumab. (A) Clinical photographs of a patient receiving placebo. At baseline, the patient exhibits substantial proptosis (left eye, 29 mm and right eye, 27 mm) as well as multiple inflammatory signs (left eye Clinical Activity Score of 7 and right eye 5). At week 24, considerable proptosis and inflammatory signs remain. (B) Images of a teprotumumab-treated patient. Baseline proptosis (both eyes 24 mm), edema, upper and lower eyelid retraction, and multiple inflammatory signs (CAS 5 bilaterally). At week 24, considerable bilateral reductions in proptosis (–5 mm) and CAS (–4 points). (C) Coronal, contrast-enhanced, fat-saturated, T1-weighted MRIs in a single patient receiving teprotumumab at baseline and at week 24. Note marked enhancement of the inferior rectus muscle (white arrowhead) and orbital fat (white arrow) as well as inferior reduced by 49% (yellow arrowhead). Proptosis reduction decreased from 23 mm at baseline to 18 mm at week 24. From N. Engl. J. Med, Douglas R.S, Kahaly G.J., Patel A., Sile E.H.Z., Thompson R. et al. Teprotumumab for the treatment of active thyroid eye disease. 382; 341–352. Copyright © (2020) Massachusetts Medical Society. Reprinted with permission.

improvement of ≥1 diplopia grade was experienced in 50.0% of patients receiving teprotumumab compared to 3.6% of those administered placebo at week 6. Diplopia improved with teprotumumab, regardless of disease severity at baseline when compared to placebo (**Figure 3E**) as did overall quality of life using the GO-QOL score (**Figure 3F**).

Additional assessments in the phase 3 trial were "before and after" facial photographs (**Figures 4A** and **4B**) and limited orbital imaging performed prior to and following treatment of six patients enrolled (off study protocol) at a single study site. Those imaging studies revealed decreased extraocular muscle volume with reduction of the inferior rectus muscle exhibiting the greatest change in 4/6 patients (**Figure 4C**). Orbital fat volume was reduced in two of these patients. Thus it would appear that teprotumumab affects both extraocular muscle and orbital fat compartments, in at least some patients with TAO.

CONCLUSIONS AND CONSIDERATION OF WHAT MIGHT LAY AHEAD FOR TEPROTUMUMAB IN TAO AND BEYOND

In aggregate, the two clinical trials demonstrate the substantial potential for IGF-IR inhibition in the treatment of active, moderate to severe TAO. Because both phase 2 and phase 3 trials excluded TAO of a duration longer than 9 months and did not allow enrollment of individuals with sight-threatening disease, neither

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study could inform the potential impact of teprotumumab in patients with either long-standing disease or compressive optic neuropathy. Thus, additional studies examining these patients will broaden our insights into the temporal dimensions of benefit and durability the drug can provide. Since its approval by the US FDA in January 2020 (9), case reports have revealed potential effectiveness in clinical stable TAO (74) and in worsening compressive optic neuropathy (75, 76). In addition, the impact of teprotumumab on thyroid function in patients retaining intact thyroid glands should be assessed prospectively. Since other autoimmune diseases, including rheumatoid arthritis, are also associated with abnormalities in the IGF-I/IGF-IR pathway (61, 69), the potential therapeutic effects of IGF-IR inhibition may yield advances in treatments of those conditions as well.

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The author confirms being the sole contributor of this work and has approved it for publication.

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Conflict of Interest: TS has been issued patents covering his inventions concerning the use of IGF-IR inhibitors as therapy in Graves' disease. These patents are held by UCLA School of Medicine and Los Angeles Biomedical Research Institute.

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Evidence That Baseline Levels of Low-Density Lipoproteins Cholesterol Affect the Clinical Response of Graves' Ophthalmopathy to Parenteral Corticosteroids

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Background: High dose intravenous glucocorticoid (ivGC) therapy is the first line treatment in moderate to severe Graves' ophthalmopathy (GO) and is associated with a clinical response rate ranging from 50% to 80%. Recently, a positive correlation between total cholesterol and low-density lipoproteins cholesterol (LDLc) with GO presentation and activity has been described.

Objective: We aimed at evaluating whether, in patients with moderate to severe active GO treated with ivGC therapy, cholesterol, and LDLc could represent valuable predictive factors of medium-term GO outcome.

Methods: This single center retrospective study was conducted in a consecutive series of 87 patients undergone ivGC therapy because affected by moderate to severe active GO. Clinical outcome of GO was evaluated at week 6 (W6) and 12 (W12) in respect to baseline conditions (week 0) by the seven points CAS according to EUGOGO recommendations. Univariate analysis and binary logistic regression were performed for the outcome variable W12CAS.

Results: In patients with active GO, an early positive clinical response to ivGC therapy (as evaluated by CAS at 6W) was a strong determinant (OR=13) of the clinical outcome at week 12. Moreover, high levels of LDLc at baseline were positively associated with a reduction in the likelihood of being classified as improved at 12W. Patients with LDLc >193.6 mg/dl were very likely to respond negatively to ivGC therapy independently from the response at 6W. Based on these results, we propose a predictive decision-making model to be tested in future prospective studies.

Discussion: We found that, in patients with active GO, both an early clinical response to ivGC therapy and baseline LDLc levels are significant determinants of GO outcome

(W12CAS). These data support the need of a cholesterol-lowering treatment before addressing these patients to ivGC therapy.

Keywords: cholesterol, low-density lipoprotein cholesterol, Graves' ophthalmopathy, parenteral corticosteroids, Graves' disease

INTRODUCTION

Graves' disease (GD) is an autoimmune disorder associated with the production of activating autoantibodies to the thyroidstimulating hormone receptor (TSH-R) in the thyroid gland, leading to hyperthyroidism. A major extra-thyroidal complication of GD is Graves' ophthalmopathy (GO), an autoimmune and inflammatory condition characterized by orbital disfigurement, double vision, and decreased visual performance up to blindness (sight-threatening GO). All these factors are associated with proven decrease of quality of life and negative social impact (1, 2). Innate and adaptive immunity as well as a large number of inflammatory factors are implicated in GO pathogenesis (3). High dose ivGC therapy is the first choice in moderate to severe or severe active GO; other immunosuppressive drugs and/or retrobulbar irradiation are widely used in the treatment of GO. Surgical orbital decompression is, however, the best choice for treating patients with moderate to severe or severe inactive GO. Treatment with 131-iodine, severe hyperthyroidism and elevated anti TSH-R antibodies (TRAbs) are considered risks factors for GO presentation or exacerbation (4). Smoking is considered to be the most important risk factor for presentation, exacerbation of GO and resistance to corticosteroids (5). Low levels of miR-224-5p also appear to reduce the clinical efficacy of medium dose of ivGC therapy in patients with moderate to severe active GO (6). Recently a retrospective analysis of a large number of patients showed an independent effect of 3-hydrossi-3methylglutaryl CoA reductase inhibitors (statins) on the risk to develop GO (7). In patients with GD, statin use for approximately 60 days during a one-year observation period significantly reduced the risk of developing GO, an effect not obtained when patients were treated with cyclooxygenase-2 (COX-2) inhibitors alone (8). A significant correlation of GO presentation and activity with total cholesterol and LDLc has been described (9) revealing a role of cholesterol in the processes involved in GO pathogenesis. Those data suggest the possibility that cholesterol might not only increase the risk of GO but also its activity, corroborating previous data showing that chronically elevated cholesterol increases systemic inflammation and modulates innate and adaptive immunity (10). However, the possible impact of cholesterol levels on the clinical efficacy of ivGC therapy in patients with active GO is not known (8, 11, 12).

Abbreviations: GD, Graves' Disease; GO, Graves' ophthalmopathy; ATD, antithyroid drugs; LDLc, low density lipoproteins cholesterol; PREDIGO, predictive score for the development or progression of Graves' orbitopathy;, ivGC, intravenous glucocorticoid; W6, week 6; W12, week 12; EuGoGo, The European Group on Graves' Orbitopathy; FT3, triiodothyronine; FT4, thyroxine or tetraiodothyronine; TSH, thyroid stimulating hormone; TRAbs, thyrotropin (TSH)-receptor autoantibodies; RAI, radioiodine; ROC, receiver operator characteristics. CTEC, combined thyroid eyes clinic.

We aimed at evaluating whether serum cholesterol levels may serve as predictive factor of medium-term clinical activity GO outcome in patients with moderate to severe active GO treated with ivGC therapy.

PATIENTS AND METHODS

Patients

We retrospectively selected 87 consecutive patients with Graves' disease and GO referred to our out-patients thyroid clinic from January 2013 to July 2019 and treated with ivGC therapy because of moderate to severe active GO. Patients with a history of treatment with statins, corticosteroids, radioiodine (RAI) for Graves' hyperthyroidism, retrobulbar radiotherapy or with incomplete metabolic evaluation were excluded from the study. Six patients previously treated with thyroidectomy 11.1 (14.6) months earlier and stably euthyroid were included.

Graves' Ophthalmopathy Clinical Evaluation at Baseline

All patients were evaluated at the combined thyroid eye clinic (CTEC) of Endocrinology Unit at Garibaldi-Nesima Hospital, Catania, Italy. The same expert endocrinologist carried out the clinical examination of GO according to European Group of Graves' Ophthalmopathy (EuGoGo) criteria (4). The lid fissure width was evaluated in millimeters by a router, proptosis of each patient was evaluated by the same Hertel exophthalmometer. Diplopia was classified according to the Gorman score by four levels of severity: absent, intermittent, inconstant, or constant. A complete ophthalmological evaluation was carried out by the same expert ophthalmologist. The GO was defined as moderate to severe when eye disease had a sufficient impact on daily life with one or more of the following clinical signs, each part of a composite index (CI): lid retraction 2 mm or more, moderate or severe soft tissue involvement, exophthalmos 3 or more mm above 21, and inconstant or constant diplopia. GO activity was evaluated according to seven points CAS and was considered active when reaching a CAS ≥ 3 .

Graves' Ophthalmopathy Clinical Outcome Evaluation

Clinical outcome of GO was evaluated at week 6 (W6) and 12 (W12) in respect to baseline conditions (week 0) by the seven points CAS and was defined clinically improved when CAS improved by almost 2 points in at least one eye. Deterioration was defined by CAS worsening of at least 2 points or when dysthyroid optic neuropathy (DON) or corneal breakdown occurred. No changes were defined as the condition in which

changes less than those described above occurred. For each assessment, patients who improved were considered as group I, patients who did not improve or who worsened were defined as group NI. According to these groups and to the time of assessment we defined the following categories: week 6 CAS (I_{W6CAS} or NI_{W6CAS}), week 12 CAS (I_{W12CAS} or NI_{W12CAS}).

Pulse Therapy

Corticosteroid treatment for GO consisted of ivGC (Solumedrol; Pfizer, Karlsruhe, Germany) injections with a median cumulative dose of 52.3 mg/kg subdivided in 12 weekly infusions. Patients were asked to report to our endocrinology day hospital on the appointed day, an indwelling venous catheter was inserted into an antecubital vein between 8.30 and 9.30 and Solumedrol diluted in 250 ml of a 0.9 sodium chloride solution % was administered at an infusion rate of 120 ml/h in the post-absorption state and in a lying position. Blood pressure, blood glucose, lipid levels, thyroid, liver and kidney function were evaluated prior to initiating the corticosteroid infusion.

Analytical Methods

Serum hormones were measured by microparticle enzyme immunoassay (Abbot AxSYM-MEIA) with inter-assay coefficients of variation of less than 10% over the analytical ranges of 1.7-46.0 pmol/L for FT3, 5.15-77.0 pmol/L for FT4, and 0.03-10.0 mU/L for TSH. The within-run and between-run precisions for FT3, FT4, and TSH assays showed coefficients of variation <5%. Thyrotropin receptor antibodies (TRAbs) were measured by a III generation assay (SELco TRAbs Human, Dahlewitz/Berlino (Germany). Glycaemia, total cholesterol, high density lipoprotein (HDL) cholesterol and triglycerides were evaluated by standard methods. We calculated LDLc values by the formula of Martin/Hopkins that is reliable for a wide range of triglyceride values (13). The thyroid was evaluated by ultrasound and the volume of each lobe was calculated with the formula for ellipsoid volumes: Volume = length*width*depth* $(\pi/6)$. The calculated thyroid volume was the sum of the volumes of the two lobes (14).

Statistical Methods

Statistical analyses were performed with the SPSS package (IBM SPSS Statistics for Windows, Version 26.0. Armonk, NY: IBM Corp). For the descriptive analysis, continuous variables were expressed as median (25th-75th percentile); categorical variables were expressed as numbers and percentages. We calculated that, to detect a moderate size effect (OR = 2) with a p value \leq 0.05 and a statistical power of 0.8, a total of 84 patients would have needed (see supplementary data).

Univariate Analysis

Univariate analysis was performed to identify predictive variables significantly associated with the clinical outcome at W12CAS ("Improved" and "Not Improved"). Continuous variables were analyzed using the non-parametric Mann Whitney U test. The shapes of distribution of each variable were evaluated by visual inspection of the population pyramid charts; for distributions of similar shape we reported the medians, for distributions of different shapes we reported average ranks. The estimate of the effect size "r" was calculated

when appropriate (15). Categorical variables were analyzed by the Chi-square test or, if cells with less than five expected cell numbers were found, by Fisher's exact test. The strength of the associations was quantified by calculating the Odds Ratio.

Binary Logistic Regression

A binary logistic regression was performed for the outcome variable (W12CAS). Covariates were selected on the basis of the results of univariate analysis and the final model was built using forced entry and a hierarchical method. Linearity of the continuous variables with respect to the logit of the dependent variable was assessed by the Box-Tidwell procedure (16) and a Bonferroni correction was applied using all terms in the model to assess its statistical significance (17). Multi-collinearity was excluded after checking tolerance (18) and variance inflation factor (VIF) (19) statistics and the proportion of the variance of each predictor's b value attributed to each eigenvalue.

The possible presence of outliers was verified by examining standardized residuals and values ≥1.96 or ≤-1.96 standard deviations were reported; cases whose standardized residuals exceeded such thresholds were inspected closely and the decision to eliminate or to keep them in the analysis was made after assessing if they exerted an undue influence on the model using the following influence statistics: Cook's distance, DFBeta, and Leverage statistics. The adequacy of the models was tested with the maximum likelihood method. The Wald test was used to verify that coefficients differed from 0. Odds ratios were calculated as the exponential of b values to give an indicator of the change in odds resulting from a unit change in the predictor. The proportion of variation in the dependent variable explained by each model was assessed by Nagelkerke pseudo-R² value (20). To predict the probability of improving at week 12 (W12CAS), we constructed our models on the basis of the theoretical model of the binary logistic regression, expressed by the following equation [Eq. 1]:

$$P(Y) = \frac{1}{1 + e - \left(b_0 + \sum_{i=1}^n b_i x_i\right)} \tag{1}$$

Where:

P(Y): probability of Y occurring; in our study $Y = I_{W12CAS}$. e: Euler's number

 b_0 : constant regression coefficient (interception at x axis)

 b_i : regression coefficient of x_i

 x_i : predictor variable

n: number of predictor variables

Logistic regression models assigned a predicted category on the basis of P (Y) values (predicted probability of improving). Using the standard cut-off P (Y) = 0.5 the models assigned a predicted category = Improved (I_{W12CAS}) when P (Y) \geq 0.5; the remaining predicted category (NI_{W12CAS}) was assigned if P (Y) was < 0.5.

Receiver Operating Characteristic Curve Analysis and Youden's Test

These analyses were performed for the W12CAS binomial logistic regression model on the basis of the regression outputs.

The ability of the model to discriminate between outcome categories was investigated in more details by elaborating the ROC curve (21). The $P(I_{W12CAS})$ (predicted probability of improving at W12) provided by regression analysis was set as test variable, in order to study the model accuracy for different P (I_{W12CAS}) cut-offs. The cut-off with the best compromise between sensitivity and specificity was assessed using Youden's test (22).

All tests were considered statistically significant at a 2-tailed p value <0.05.

RESULTS

Patient characteristics at study entry are shown in **Table 1**. Eighty-seven patients completed the 12-week (12W) GO clinical assessment according to the seven-points CAS and 71.3% of them improved. Fifty-three patients who had a complete metabolic assessment were included in the analysis.

Univariate and Binomial Logistic Regression Analysis according to CAS Evaluation

On univariate analysis, the only two variables positively associated with GO improvement at 12W (I_{W12CAS}) were serum levels of LDLc (r=-0.25, p=0.045) and early improvement at W6 (I_{W6CAS}) (p<0.001) (**Figure 1**, **Table 2**). Notably, 90.6% patients classified as "improved" at W6 were also "improved" at W12. In contrast, 41.2% of patients 'not improved' at W6 (NI_{W6CAS}) were classified as 'improved' at W12 (OR=13.714, CI 95% 3.61–52.095, p<0.001, I_{W6CAS} vs NI_{W6CAS}).

We found a negative association between male gender and GO improvement at W12; however, this difference did not reach statistical significance (OR = 2.778, CI 95% 1.003–7.691, p=0.058).

TABLE 1 | Characteristics of patients at study entrance.

Variables Number of patients 87 ivGC treatment period 1/2013-7/2019 Sex: male/female (n, %) 22 (25.3)/65 (74.7) Age (vears) 45 (35-55) BMI (Kg/mg) 24.1 (21.6-28) Current smokers (n, %) 45 (51.7) Previous Smokers (n, %) 29 (33.3) Non Smokers (n, %) 13(14.9) 18.1 (13-24.3) Thyroid volume (ml) 52.3 (40.2-68.2) ivGC cumulative dose (mg/kg) CAS 3 (3-4) GO duration (months) 8 (3-12) 0.05 (0-0.9) TSH (mU/L) FT3 (pg/ml) 2.86 (2.53-3.64) 0.98 (0.86-1.29) FT4 (ng/dl) 5.31 (2.44-18.15) TRAbs (IU/I) Total cholesterol (mg/dl) 200 (173-226) HDL (mg/dl) 53 (47.8-61.3) 81.5 (60.8-115) Triglycerides (mg/dl) LDLc (mg/dl) 123.3 (104.1-151.9) Glycaemia (mg/dl) 94 (87.8-102.3)

Data are reported as number and percentage or median and interquartile range.

Similarly, higher total cholesterol and older age showed some association with the lack in GO improvement at W12, without reaching statistical significance (p=0.094 and p=0.08, respectively) There was no association between smoking habit and W12 outcome and no association with any of the other continuous variables considered (**Table 2**). Prior thyroidectomy was not a contributing factor as at 12W three out of six patients were classified as improved and three as not improved.

Variables identified at univariate analysis (LDLc and W6CAS) were then evaluated by binomial logistic regression to further ascertain their effects on the likelihood to affect GO improvement at W12. The continuous independent variable was found to be linearly related to the logit of the dependent variable. There were two standardized residuals with a value of -5.986 and -4.706 standard deviations, which were kept in the analysis. The logistic regression model was statistically significant, $\chi 2(2) =$ 15.985, p < 0.001. The model explained 41.1% (Nagelkerke R^2) (20) of the variance in W12CAS outcome and correctly classified 84.6% of cases. Sensitivity was 95.1%, specificity was 45.5%, positive predictive value was 86.7% and negative predictive value was 71.4%. Both predictor variables were statistically significant. I_{W6CAS} had 13 higher odds to be classified as I_{W12CAS} than NI_{W6CAS}. Increasing LDLc was associated with a reduction in the likelihood of being classified as IW12CAS, since for each unit of LDLc reduction, the odds of improving at W12CAS increased by a factor of 1.03. **Table 3** reports the regression output of the final model. Based on the results of the binary logistic regression, the following model was constructed to predict the P(IW12CAS) (probability of improving at W12CAS) for individual patients [Eq. 2]:

$$P(I_{W12CAS}) = \frac{1}{1 + e^{-(2.95 - 0.025x_1 + 2.57x_2)}}$$
 (2)

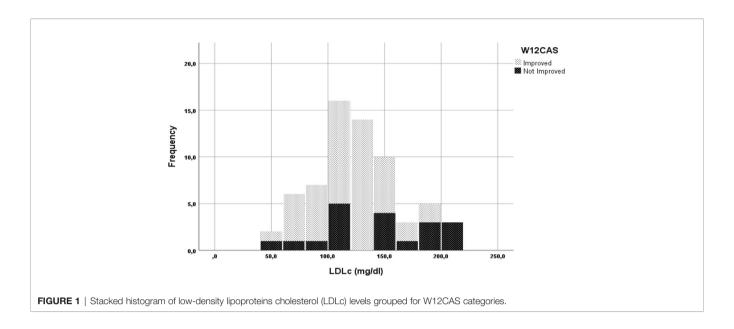
Where: $x_1 = \text{LDLc}$ (mg/dl); $x_2 = \text{W6CAS}$ outcome ($x_2 = 1$ for I_{W6CAS} ; $x_2 = 0$ for NI_{W6CAS}). We tested the reliability of our model by performing several multivariate logistic regression analyses, with the aim of verifying that LDLc is an independent predictor for the W12CAS outcome. We also constructed an overfitted model constructed with 5 covariates (see supplementary data).

Receiver Operating Characteristic Curve Analysis

The ROC curve processed using the regression $P(I_{W12CAS})$ resulted in an excellent level of discrimination (0.805; 95% CI, 0.608–1.000) according to Hosmer et al. (23). (**Figure 2**). Youden's test identified a $P(I_{W12CAS}) = 0.664$ as best cut-off, with sensitivity 90.2%, specificity 81.8%, positive predictive value 94.9%, negative predictive value 69.2% and accuracy 88.5%. **Figure 3** highlights the impact that different $P(I_{W12CAS})$ cut-offs have on the prediction performance of the model.

W12CAS Binomial Logistic Regression Implementation

Almost every patient classified as "improved" at W6CAS had a P (I_{W12CAS}) between 0.8 and 1.0, allowing a correct diagnosis in



almost all of them. Within the I_{W6CAS} group the lower P (I_{W12CAS}) (ranging from 0.59 to 0.7) was assigned to the patients with a very high LDLc and the Youden's cut-off (unlike the standard cut-off $P(I_{W12CAS}) = 0.5$) allowed to correctly predict their W12CAS outcome. Considering that the two independent variables (LDLc and W6CAS outcome) are not available at the same time, since the first one is obtained at baseline and the second one after a 6 weeks pulse therapy, we aimed at identifying a LDLc cut-off that would give a $P(I_{W12CAS}) < 0.664$ independently of W6CAS outcome; knowing this value may allow the physician to delay pulse therapy until this value is corrected. This LDLc cut-off was calculated using the binomial logistic regression model (Eq. 3) setting $P(I_{W12CAS}) < 0.664$ (as indicated by Youden's test):

$$P(I_{W12CAS}) = \frac{1}{1 + e^{-(2.95 - 0.025x_1 + 2.57x_2)}} < 0.664$$
 (3)

The resulting inequality was solved for both x_2 valid values (0 and 1) with the purpose to calculate which LDLc values (x_1) would certainly give a $P(I_{W12CAS}) < 0.664$, independently from W6CAS outcome (x_2) . Calculations resulted in $x_1>193.6$. According to this data, a patient showing a baseline LDLc >193.6 mg/dl can't reach a $P(I_{W12CAS}) > 0.664$; in fact, in the best scenario $(x_2 = 1)$, he would reach a $P(I_{W12CAS})$ value slightly lower than 0.664, while in the worst one $(x_2 = 0)$ its $P(I_{W12CAS})$ should fall below 0.132. For W6CAS Improved patients, the outcome of the W6CAS itself is the most important predictor, with LDLc making a predominant contribution when it exceeds 190 mg/dL. Based on these results, a proposal for a decision algorithm was developed (**Figures 3** and **4**).

TABLE 2 | Univariate analysis for W12CAS outcome.

| Variables | I _{W12CAS} | NI _{W12CAS} | р | |
|--------------------------|---------------------|----------------------|--------|--|
| Glycaemia (mg/dl) | 93.5 (87.8–102) | 95 (86.3–104.5) | 0.885 | |
| Age*(years) | 41.01 | 51.42 | 0.082 | |
| Females (%) | 76.9 | 23.1 | 0.058 | |
| BMI (Kg/m ²) | 24.1 (21.7–27.9) | 25.6 (21.3–28.6) | 0.929 | |
| Smokers (%) | 68.9 | 31.1 | 0.643 | |
| ivGC CD (mg/Kg) | 50.4 (40.2–67.2) | 60 (50.6-73.2) | 0.131 | |
| Thyroid (ml) | 18.6 (11.9–24.3) | 18.1 (14.5–39.9) | 0.546 | |
| CAS | 3 (3–4) | 3 (3–4) | 0.927 | |
| TSH (mU/L) | 0.05 (0-0.9) | 0.05 (0.01–1) | 0.661 | |
| FT3 (pg/ml) | 2.87 (2.56–3.43) | 2.69 (2.39–3.76) | 0.699 | |
| FT4 (ng/dl) | 0.98 (0.84–1.25) | 1.05 (0.89–1.3) | 0.975 | |
| TRAbs (IU/I) | 4.6 (2.2–16.5) | 8.4 (3.3–28) | 0.156 | |
| Cholesterol* | 31.67 | 39.89 | 0.094 | |
| HDL* | 35.06 | 29.63 | 0.298 | |
| Triglycerides (mg/dl) | 76.0 (58–112) | 89.0 (74-131) | 0.315 | |
| LDLc* | 30.49 | 40.95 | 0.045 | |
| I-W6CAS (n, %) | 48 (90.6) | 5 (9.4) | <0.001 | |

Data are expressed as number or percentage percentage, median and interquartile range or mean ranks as indicated (*); CD, cumulative dose.

TABLE 3 | Outputs of the binary logistic regression for the outcome variable W12CAS.

| | В | SE | Wald | df | р | Odds Ratio | 95% CI of Odds Ratio | |
|-------------------|----------------|----------------|----------------|--------|----------------|------------------|----------------------|--------|
| | | | | | | | Lower | Upper |
| LDLc | -0.025 | 0.013 | 4.025 | 1 | 0.045 | 0.975 | 0.951 | 0.999 |
| W6CAS Constant | 2.570 2.950 | 0.863 1.707 | 8.858 2.987 | 1 1 | 0.003 0.084 | 13.065 19.099 | 2.405 | 70.976 |

Final model built with covariates LDLc and W6CAS.

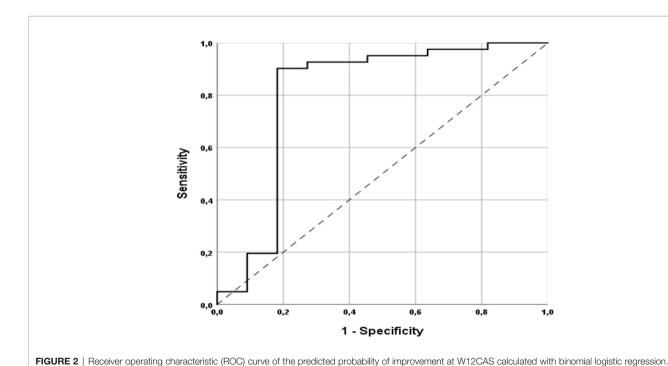
DISCUSSION

Herein we found that in patients with active GO addressed to pulse therapy with corticosteroids, baseline serum levels of LDLc and early clinical response at week 6 (W6CAS) were the strongest determinants of the clinical outcome at week 12 according to CAS evaluation (W12CAS). A previous study correlated serum total cholesterol and LDLc levels with the onset and severity of GO, but their impact on the GO response to corticosteroids has not been determined (9). In our study, we modeled the predicted probability P (I_{W12CAS}) that each patient is classified as improved at W12CAS based on baseline serum levels of LDLc plus W6CAS outcome, showing that the LDLc level provides an independent contribution to the W6CAS data by overcoming the effect of W6CAS when it exceeds 190 mg/dl with a positive predictive value of 86.7% and a negative predictive value of 71.4%. Furthermore, the ROC curves indicated a sensitivity of 90.2% and a specificity of 81.8% in predicting a positive response of GO to corticosteroids at 12W.

Active GO is often an auto resolving condition, but several factors independently modulate its severity and ultimate clinical outcome. Although ivGC therapy is the most widely used medical treatment for active GO, its clinical efficacy is variable,

and it is therefore important to have biomarkers that can predict its efficacy in the individual patient.

Few variables are known to influence the severity and outcome of GO. For example, smoking is a proven independent risk factor for the development and exacerbation of GO, and radioiodine (131-I) treatment of hyperthyroidism can also trigger GO and worsen its outcome. Furthermore, failure to achieve euthyroidism and persistence of high levels of TRAbs are negatively correlated with the clinical outcome of GO (24). Recently, a predictive risk score for the development of GO in patients with newly diagnosed Graves' hyperthyroidism has been proposed (25). This score (PREDIGO) aims to estimate the odds ratio of GO exacerbation according to CAS based on serum levels of TSH and TRAbs, duration of hyperthyroidism and smoking habits. In our study, GO was moderate to severe and active in patients with stable thyroid function and with similar TRAbs levels between responders and non-responders, therefore the risk of GO exacerbation was similar across the study population. In addition to the aforementioned risk factors, GO occurrence and severity have been positively associated with diabetes mellitus (DM) (12, 26) and serum levels of total cholesterol and LDLc (9). These data are of interest since both DM and hypercholesterolemia induce a chronic inflammatory



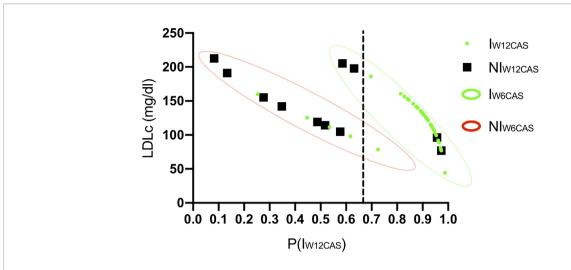
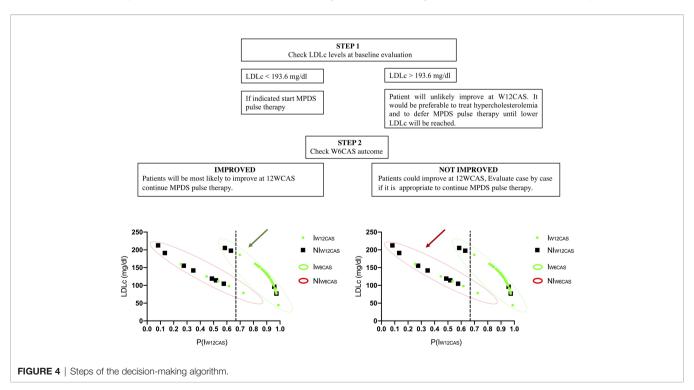


FIGURE 3 | Suggested decision-making algorithm. The graphs are constructed on the basis of binomial logistic regression prediction using the Youden's best cutoff $P(I_{W12CAS}) = 0.66$. Each dot refers to a single patient; green dots refer to patients classified as Improved at W12CAS, while black dots refer to Not Improved Patients. The predicted probability of improvement at W12CAS $[P(I_{W12CAS})]$ provided by regression analysis is positioned on the x axis; cut-off = 0.664 (dotted line). The continuous predictive variable included in the model (LDLc) is positioned on the Y axis. The categories of the dichotomous predictive variable (W6CAS outcome) are highlighted in green (I_{W6CAS}) or red (I_{W6CAS}) contour lines.

state that adds to GO immunological process and contributes to remodeling of the orbital tissues. Elevated LDLc levels have been hypothesized to increase the influx of free fatty acids into the liver causing the production of reactive oxygen radical species (ROS) and the secretion of interleukin-6 (IL-6). The increase in IL-6 activity within the orbital environment can promote greater secretion of insulin growth factor 1 (IGF-1) by orbital fibroblasts of GO patients and improve the proliferation of fibroblasts and the expansion of soft orbital tissues favoring

IGF-1R - TSHR crosstalk (8, 27–29). These studies raise the need to investigate the impact of LDLc levels on the GO response to immunosuppressive therapy.

In our series of patients, none had previously received corticosteroids or retrobulbar radiation (30) and none had received radioiodine therapy to control hyperthyroidism. At week 6 after initiation of corticosteroid therapy, more than 90% of patients classified as responders maintained the same clinical rating at week 12. Corticosteroid responders at week 6



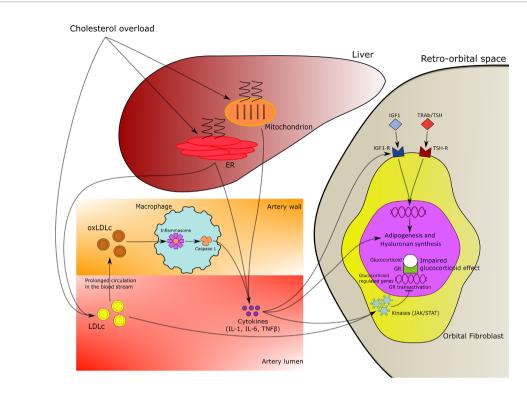


FIGURE 5 | Mechanistic hypothesis of relationship of hypercholesterolemia with chronic inflammation and resistance to corticosteroids effects in patients with Graves' ophthalmopathy (GO).

were significantly more likely than non-responders to have a positive outcome at week 12 (OR 13.7; p <0.001) confirming the previous observations (31). In our series, total cholesterol levels were higher in non-responders than in responders, but this difference did not reach statistical significance, while baseline LDLc levels were inversely related to GO clinical outcome at 12W. Other variables recognized as risk factor for GO presentation and/or exacerbation, such as cigarette smoking (24), male gender and age older than 60 yrs (32, 33) were not significantly predictive of GO response to ivGC at week 12. Although these findings may appear unexpected especially for smoking habit, the impact of smoking on the clinical response of GO to ivGC in a similar setting has not been previously investigated and deserves to be studied in prospective trials.

Our results are in agreement with the concept that hypercholesterolemia can promote chronic inflammation leading to resistance to the effects of corticosteroids characterized by inhibition of tissue macrophage functions, such as chemotaxis, phagocytosis, proliferation and antigen presentation (34). In particular, oxidized low-density lipoprotein (oxLDL) regulates the expression of dipeptidyl dipeptidase IV (DPP4) in macrophages leading to the increase of CD36 + cells which are representative of the inflammatory processes of atherosclerosis in obese and insulin resistant patients. Moreover, a direct effect of LDLc on transcriptional and translational activities of corticosteroids at the cellular levels has also been hypothesized (35, 36; **Figure 5**).

Our study has some limitations. Although all patients in our study were treated following a rigorous protocol, the study is retrospective, and the regression analysis was built on the basis of a small patient population. For these reasons, the reported algorithm is a suggestion that should be evaluated in a prospective context. However, our data indicate that patients with active GO may have a clinically limited response to ivGC therapy if LDLc levels are above 190 mg/dL.

CONCLUSIONS

Early clinical response to ivGC therapy in patients with active GO is a strong determinant of clinical outcome at week 12 when assessed by CAS. Additionally, baseline serum LDLc levels are also an independent predictor of response to ivGC therapy, and LDLc levels above 190 mg/dL greatly reduce the likelihood of a positive clinical outcome at week 12. Based on our data, we suggest measuring baseline LDLc levels in all patients with active GO and strongly considering cholesterol-lowering treatment before referring these patients to ivGC therapy.

DATA AVAILABILITY STATEMENT

The raw data supporting the conclusions of this article will be made available by the authors, without undue reservation.

ETHICS STATEMENT

The studies involving human participants were reviewed and approved by the Ethics Committee of Garibaldi Hospital—Catania. Written informed consent for participation was not required for this study in accordance with the national legislation and the institutional requirements.

AUTHOR CONTRIBUTIONS

RL conceptualized and designed the study. RL, AN, FG, MA, CR, FF, and DM collected the data. AN, RL, and FG performed the

statistical analysis. RL and AN were in charge of the figures. RL, AB, FF, and AN wrote and revised the manuscript. All authors read and approved the final manuscript. All authors contributed to the article and approved the submitted version.

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Conflict of Interest: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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Role of Proprotein Convertase Subtilisin/Kexin Type 9 in the Pathogenesis of Graves' Orbitopathy in Orbital Fibroblasts

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Background: The proprotein convertase subtilisin/kexin type 9 (PCSK9) has been implicated in the pathogenesis of inflammatory diseases. We sought to investigate the role of PCSK9 in the pathogenesis of Graves' orbitopathy (GO) and whether it may be a legitimate target for treatment.

Methods: The *PCSK9* was compared between GO (n=11) and normal subjects (n=7) in orbital tissue explants using quantitative real-time PCR, and in cultured interleukin-1 β (IL-1 β)-treated fibroblasts using western blot. Western blot was used to identify the effects of PCSK9 inhibition on IL-1 β -induced pro-inflammatory cytokines production and signaling molecules expression as well as levels of adipogenic markers and oxidative stress-related proteins. Adipogenic differentiation was identified using Oil Red O staining. The plasma PCSK9 concentrations were compared between patients with GO (n=44) and healthy subjects (n=26) by ELISA.

Results: The *PCSK*9 transcript level was higher in GO tissues. The depletion of PCSK9 blunted IL-1β-induced expression of intercellular adhesion molecule 1 (ICAM-1), IL-6, IL-8, and cyclooxygenase-2 (COX-2) in GO and non-GO fibroblasts. The levels of activated nuclear factor kappa-light-chain-enhancer of activated B cells (NF- κ B) and phosphorylated forms of Akt and p38 were diminished when PCSK9 was suppressed in GO fibroblasts. Decreases in lipid droplets and attenuated levels of peroxisome proliferator-activated receptor gamma (PPARγ), CCAAT/enhancer-binding protein β (C/EBPβ), and leptin as well as hypoxia-inducible factor 1α (HIF- 1α), manganese superoxide dismutase (MnSOD), thioredoxin (Trx), and heme oxygenase-1 (HO-1) were noted when PCSK9 was suppressed during adipocyte differentiation. The plasma PCSK9 level was significantly higher in GO patients and correlated with level of thyrotropin binding inhibitory immunoglobulin (TBII) and the clinical activity score (CAS).

Conclusions: PCSK9 plays a significant role in GO. The PCSK9 inhibition attenuated the pro-inflammatory cytokines production, oxidative stress, and fibroblast differentiation into adipocytes. PCSK9 may serve as a therapeutic target and biomarker for GO.

Keywords: adipogenesis, Graves' orbitopathy, inflammation, oxidative stress, proprotein convertase subtilisin/kexin type 9, PCSK9, thyroid eye disease

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INTRODUCTION

Graves' orbitopathy (GO) is an inflammatory autoimmune disorder, and it is the most frequent extrathyroidal manifestation of Graves' disease (1). Clinical features of GO include upper eyelid retraction, edema and erythema of the periorbital tissue and conjunctiva, proptosis, corneal ulceration, and optic neuropathy. Three cell types which predominantly contribute to the development and progression of GO are B cells, T cells, and orbital fibroblasts (2). Stimulated by interactions with T cells and autoantibodies produced by B cells, orbital fibroblasts play a key role in the establishment of inflammation by producing cytokines, chemokines, and lipid mediators. Furthermore, they proliferate, synthesize extracellular matrix, and differentiate into adipocytes, leading to tissue remodeling characteristic of GO. The mainstay treatment for moderate-to-severe GO is systemic glucocorticoids therapy (3). Due to inadequate responses and adverse effects to glucocorticoids, however, there have been several investigations for other possible biological therapies (3).

The proprotein convertase subtilisin/kexin type 9 (PCSK9), which was first reported in 2003, is the ninth member of the protein convertase family (4). PCSK9 targets low density lipoprotein receptors (LDLR) on the hepatic cell surface, toward lysosomes for degradation, resulting in elevated serum LDL cholesterol levels (5). Now, PCSK9 inhibitors have emerged as novel therapeutics to treat cardiovascular diseases (6). However, current data suggest that PCSK9 inhibitors may have pleiotropic effects, affecting targets beyond LDLR (7-9). According to other studies, PCSK9 may be a key molecule in the pathophysiology of diseases such as atherosclerosis, myocardial ischemia, Alzheimer's disease, psoriasis, and fatty liver disease (10-14). Studies on the PCSK9 in atherosclerosis, a chronic inflammatory disorder of vessel walls, showed that PCSK9 inhibition suppressed inflammatory cytokines production and decreased the activity of nuclear factor kappalight-chain-enhancer of activated B cells (NF-κB) and intracellular cell adhesion molecule 1 (ICAM-1). Additionally, silencing PCSK9 decreased oxidative stress, apoptosis, proliferative capacity, and accumulation of macrophages (10, 14, 15). Numerous studies reporting the benefit of PCSK9 suppression in vivo and in vitro suggest that PCSK9 may be an attractive target in chronic inflammatory disorders (16). However, no previous studies have reported the effect of PCSK9 inhibition in GO.

In light of what is said above, this study was designed to investigate the role of PCSK9 in the pathogenesis of GO. We used small interfering RNA (siRNA) to promote cleavage of intracellular PCSK9 mRNA in orbital fibroblasts obtained from GO and normal subjects. We tested whether PCSK9 siRNA counteracts inflammation, proliferation, and adipocyte differentiation in orbital fibroblasts, the main pathogenic mechanisms in GO. In addition, we examined whether the plasma PCSK9 levels reflect the presence and the activity of GO using the clinical activity score (CAS).

MATERIALS AND METHODS

Reagents

The antibodies were purchased from Santa Cruz Biotechnology (Santa Cruz, CA, USA), Cell Signaling Technology (Beverly, MA, USA), Novus Biologicals (Centennial, CO, USA), and Abcam (Cambridge, UK). The antibodies used in the study are listed in detail in Supplementary Table 1. PCSK9 siRNA and control siRNA were obtained from Santa Cruz Biotechnology, Inc. (Dallas, TX, USA). TransIT-siQUEST siRNA Transfection reagent was purchased from Mirus Bio, Inc. (Madison, WI, USA). The 3-(4,5-dimethyl-thiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT) assay and Oil Red O were products from Sigma-Aldrich, Inc. (Merck KGaA, Darmstadt, Germany). Dulbecco's modified Eagle's medium (DMEM), fetal bovine serum (FBS), penicillin, and gentamicin were purchased from Hyclone Laboratories, Inc. (Logan, UT, USA). Recombinant human interleukin-1β (IL-1β) and the enzymelinked immunosorbent assay (ELISA) kit for PCSK9 were obtained from R&D Systems (Minneapolis, MN, USA).

Subjects and Preparation of Tissues and Cells

Orbital tissue specimens were collected from GO subjects during orbital decompression surgery (nine females and two males; age 38-54 years). Non-GO orbital tissue was obtained during the course of upper (n=3) and lower lid (n=4) blepharoplasty from patients without history or clinical evidence of any thyroid disease (5 females and 2 males; age 36-57 years). Out of 11 GO and seven non-GO tissues, three GO and three non-GO tissues were randomly chosen for primary orbital fibroblast cultures. For gene expression analysis, nine out of 11 GO tissues were randomly selected, while all seven non-GO tissues were used. The study protocol was approved by the Institutional Review Board of Severance Hospital, and all participants provided written informed consent. This research adhered to the tenets of the Declaration of Helsinki. At the time of surgery, all GO patients were in euthyroid state and had not been administered steroid or radiation therapy for at least three months.

For plasma PCSK9 evaluation, 70 subjects were recruited: 22 with active GO (15 females and 7 males; age 42.41 ± 17.91 years), 22 with inactive GO (16 females and 6 males; age 40.64 ± 16.91 years), and 26 healthy volunteers (23 females and three males; age 36.69 ± 14.20 years). GO was considered "active" based on CAS, a grading system based on the seven classic features of inflammation in GO (17). Out of seven, GO was considered "active" if the CAS was ≥ 3 . **Table 1** shows the demographic, clinical, and serologic data of the subjects.

Orbital fibroblasts were isolated from the harvested tissue and cultured as described previously (18). After being minced, the tissue was placed directly in DMEM/F12 (in 1:1 ratio) medium containing 20% FBS, penicillin (100 U/ml), and gentamycin (20 μ g/ml). Following incubation, tissues were maintained in solution containing DMEM, antibiotics, and 10% FBS. Once the growth of the fibroblasts was confirmed, the cells were treated with trypsin/ethylenediaminetetraacetic acid and passaged in

TABLE 1 | Clinical and serological characteristics of patient population for ELISA.

| | Active GO (n=22) | Inactive GO (n=22) | Non-GO (n=26) | p-value |
|---|-------------------|--------------------|-------------------|---------|
| Sex (male/female) | 7/15 | 6/16 | 3/23 | 0.213 |
| Age (years), $M \pm SD$ | 42.41 ± 17.91 | 40.64 ± 16.91 | 36.69 ± 14.20 | 0.575 |
| Smokers, n (%) | 6 (27.27) | 4 (18.18) | 2 (7.69) | 0.202 |
| PCSK9 (ng/ml) | 256.46 ± 53.49 | 223.48 ± 36.42 | 190.83 ± 28.77 | < 0.001 |
| CAS | 4.45 ± 1.47 | 1.36 ± 0.66 | _ | < 0.001 |
| Duration GD (months), median (IQR) | 8.55 (2-16) | 7.18 (2–22) | _ | 0.396 |
| T3 (0.58–1.59 ng/dl), $M \pm SD$ | 1.17 ± 0.25 | 1.17 ± 0.26 | _ | 0.796 |
| Free T4 (0.70-1.48 ng/dl), M ± SD | 1.19 ± 0.22 | 1.23 ± 0.21 | _ | 0.751 |
| TSH (0.35–4.94 μ IU/ml), $M \pm SD$ | 1.40 ± 0.84 | 1.36 ± 1.00 | _ | 0.605 |
| TBII (0-1.75 IU/L), M ± SD | 18.09 ± 12.26 | 12.99 ± 7.42 | _ | 0.231 |

ELISA, enzyme-linked immunosorbent assay; GO, Graves' orbitopathy; SD, standard deviation; PCSK9, proprotein convertase subtilisin/kexin type 9; CAS, clinical activity score; GD, Graves' disease; IQR, interquartile ranges; T3, triiodothyronine; T4, thyroxine; TSH, thyrotropin; TBII, thyrotropin-binding inhibitory immunoglobulin.

monolayers. Strains were stored in liquid nitrogen and only those between the third and fifth passages were used for experiments.

Cell Viability Assay

Cell viability was assessed with an MTT assay, following the manufacturer's protocol (Sigma-Aldrich, Inc.). Orbital fibroblasts obtained from GO patients were seeded into 24-well culture plates (1×10^5 cells/well) and treated with PCSK9 and control siRNAs (50 nM) for 10, 24, and 48 h. Thereafter, cells were washed and incubated in MTT solution (5 mg/ml) for 3 h at 37° C. Dimethyl sulfoxide (DMSO) was applied for solubilization and the absorbance of the converted dye was measured with a microplate reader (EL 340 Bio Kinetics Reader; Bio-Tek Instruments, Winooski, VT, USA) at 540 nm, with background subtraction at 630 nm.

Quantitative Real-Time PCR

The RNA was extracted from cells using TriZol (Invitrogen, Carlsbad, CA, USA). Out of the extract, 1 μ g was reverse-transcribed into cDNA (Qiagen, Valencia, CA, USA) and amplified with SYBR green real-time PCR master mix in a StepOne Plus real-time PCR thermocycler (Applied Biosystems, Foster City, CA, USA). The sequence of primers is listed in **Supplementary Table 2**. The PCR results for each type of mRNA were normalized to the level of GAPDH, and expressed as fold-change in the Ct value relative to the control group using the $2^{-\Delta\Delta Ct}$ method (19).

Western Blot Assay

Equal amounts of protein (50 $\mu g)$ were separated by 10% SDS polyacrylamide gel electrophoresis. The resolved proteins were transferred to nitrocellulose membranes and incubated overnight with primary antibodies at 4°C. Then, the membranes were probed with horseradish peroxidase–conjugated secondary antibodies. The bands were detected on X-ray films (GE Healthcare, Piscataway, NJ), and their intensities were quantified and normalized to that of the $\beta\text{-actin}$ in the same sample.

Adipogenesis

Using a previously published protocol, adipocyte differentiation of GO fibroblasts was induced (20). Cells were cultured in serum-free DMEM supplemented with T3, insulin (Boehringer-Mannheim,

Mannheim, Germany), carbaprostaglandin (cPGI2; Calbiochem, La Jolla, CA, USA), and dexamethasone, along with proliferator-activated receptor gamma (PPAR γ) agonist, rosiglitazone (10 μ M; Cayman, Ann Arbor, MI, USA) for 7 days. To evaluate the effect of PCSK9 siRNA on adipogenesis, cells were transfected with PCSK9 siRNA for the entire 7-day differentiation period according to the manufacturer's instructions.

Oil Red O Staining

Cells were stained with Oil Red O as described by Green and Kehinde (21). A working solution was prepared by diluting 6 ml of a stock solution (0.5% Oil red O in isopropanol) with 4 ml of distilled water. The cells were fixed with 3.7% formalin at 4°C for 1 h before being washed with PBS and mixed with Oil Red O solution for 1 h at room temperature. The cell-solution mixture was visualized under on a light microscope (Olympus BX60; Olympus Corp., Melville, NY, USA).

Blood Sampling and Measurement of Plasma PCSK9 and TBII Concentrations

Blood samples were drawn into test tubes containing sodium citrate. Platelet-free plasma was obtained after centrifugation at 1,500 g for 15 min at 4°C and stored at -80°C until analysis. Plasma PCSK9 levels were determined with a commercially available ELISA kit. All samples were tested in triplicate, and all sera were run in the same assay. The average value of three repeated assays was used for statistical analyses. Thyrotropin (TSH) binding inhibitory immunoglobulin (TBII) was measured with a third-generation TBII assay using the automated Cobas electrochemiluminescence immunoassay (Elecsys; Roche Diagnostics GmbH, Penzberg, Germany).

Statistical Analysis

All experiments were performed in duplicate or triplicate on samples from each patient, and the results were expressed as mean ± standard deviation (SD). Comparisons of data between groups were performed with the independent *t*-test or ANOVA. The Bonferroni test was performed as a *post hoc* test. The Mann–Whitney *U*-test and Kruskal–Wallis test were used for nonparametric or not normally distributed data. Spearman's rank correlation coefficient was used to analyze the correlation of plasma PCSK9 concentrations with CAS and plasma TBII levels. The SPSS for Windows, version 20.0 (SPSS, Inc.,

Chicago, IL, USA) was used. A *p*-value < 0.05 denoted statistical significance.

RESULTS

GO Tissues Show Increased Expression of PCSK9, LDLR, and HIF-1 α

To investigate its potential role in GO, we measured the expression of PCSK9 in orbital tissues taken from GO and non-GO subjects. The RT-PCR results showed that PCSK9

transcript levels were greater in GO tissues (n=9) than in non-GO tissues (n=7) (**Figure 1A**). Additionally, the mRNA levels of LDLR and hypoxia-inducible factor 1α (HIF- 1α) were higher in GO tissues (n=9) than in non-GO tissues (n=7) (**Figures 1B, C**).

IL-1β Induces PCSK9 and LDLR in Orbital Fibroblasts

We challenged GO and non-GO orbital fibroblasts with IL-1 β , a key mediator in GO (18, 22), for 1, 3, 6, 16, and 24 h. The western blot results showed that IL-1 β led the GO and non-GO fibroblasts to increase PCSK9 and LDLR expression in a time-

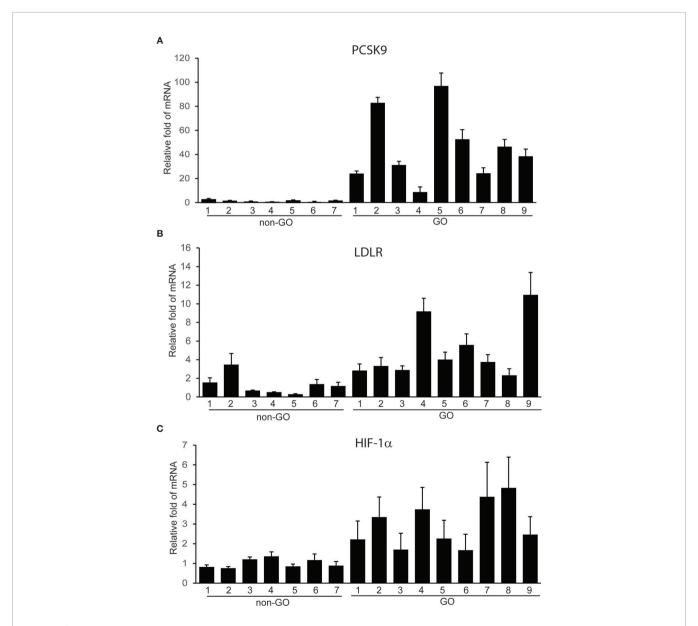


FIGURE 1 | Expression of proprotein convertase subtilisin/kexin type 9 (PCSK9), low density lipoprotein receptor (LDLR), and hypoxia-inducible factor 1a (HIF-1a) mRNAs in Graves' orbitopathy (GO) and non-GO orbital tissues. The RNA extracted from GO (n=9) and non-GO (n=7) orbital tissues was reverse-transcribed by real-time PCR and quantified. Experiments were performed in triplicate for each donor. The results showed elevated transcript levels of PCSK9 (A), LDLR (B), and HIF-1a (C) in GO tissues than in non-GO tissues. Data in the column indicate the mean ± SD fold elevation relative to the control.

dependent manner (**Figure 2**). The increase in PCSK9 and LDLR was more prominent in GO than in non-GO fibroblasts.

Silencing PCSK9 Suppresses IL-1β-Induced Expression of Pro-Inflammatory Mediators

The western blot results showed that when orbital fibroblasts were transfected with PCSK9 siRNA, the production of pro-inflammatory mediators, ICAM-1, IL-6, IL-8, and cyclooxygenase-2 (COX-2), in response to the challenge with IL-1 β was significantly suppressed in GO and non-GO fibroblasts (**Figure 3**).

Silencing PCSK9 Reduces Activation of Signaling Molecules

As shown in **Figure 4**, the IL-1 β -treatment (10 ng/ml for 60 min) increased levels of nuclear NF- κ B p65 and phosphorylated (p-) forms of Akt and p38 in GO and non-GO fibroblasts in western blot analyses. In GO fibroblasts, PCSK9 interference with siRNA reduced IL-1 β -stimulated expression of nuclear NF- κ B p65, p-Akt, and p-p38. In non-GO fibroblasts, the PCSK9 inhibition suppressed IL-1 β -stimulated expression of p-Akt, but not NF- κ B p65 and p-p38.

Silencing PCSK9 Inhibits Proliferation of GO Fibroblasts

The enhanced proliferative capacity of GO fibroblasts at baseline and in response to certain cytokines may play a role in the pathogenesis of GO (23). According to the MTT assay results, the proliferation of GO fibroblasts slowed down 48 h after being transfected with PCSK9 siRNA compared to control siRNA-treated group (**Supplementary Figure 1**).

PCSK9 Inhibition Suppresses Adipocyte Differentiation and Oxidative Stress-Related Protein Production in GO Fibroblasts

The transfection of differentiating fibroblasts with PCSK9 siRNA attenuated adipogenesis on day 7 according to the Oil Red O staining (**Figure 5A**). When quantified by measuring optical density of Oil Red O-stained cell lysates at 490 nm, the same pattern was identified (**Figure 5B**). Throughout adipogenic differentiation, the PCSK9 levels gradually increased (**Figure 5C**), but PCSK9 siRNA substantially diminished levels of adipogenic transcription factors, PPAR γ and CCAAT/enhancer-binding protein β (C/EBP β), and mature adipocyte marker, leptin. In addition, silencing PCSK9 markedly decreased the levels of oxidative stress-related protein, HIF-1 α , and antioxidant proteins, manganese superoxide dismutase (MnSOD), thioredoxin (Trx), and heme oxygenase-1 (HO-1) (**Figure 5C**).

Plasma PCSK9 Protein Levels Are Elevated in GO and Plasma PCSK9 Concentrations Show Positive Correlations With Plasma TBII Levels and CAS

The plasma PCSK9 levels were measured in GO and healthy subjects by ELISA (**Figure 6A**). The mean plasma PCSK9 level was significantly higher in GO patients (n=44, 239.97 \pm 48.20 ng/ml) than in healthy subjects (n=26, 190.83 \pm 28.77 ng/ml; p < 0.01). Additionally, the mean plasma PCSK9 level was significantly higher in patients with active GO (n=22, 256.46 \pm 53.49 ng/ml) than in patients with inactive GO (n=22, 223.48 \pm 36.42 ng/ml; p=0.01). The plasma PCSK9 level was correlated

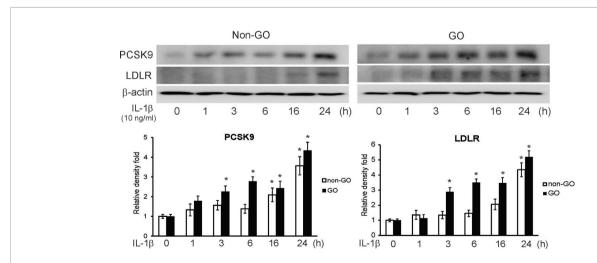


FIGURE 2 | Western blot of proprotein convertase subtilisin/kexin type 9 (PCSK9) and low density lipoprotein receptor (LDLR) after interleukin-1β (IL-1β) treatment. Confluent orbital fibroblasts obtained from Graves' orbitopathy (GO) (n=3) and non-GO subjects (n=3) were treated with 10 ng/ml of IL-1β for increasing lengths of time (0–24 h). Western blot analyses were performed to investigate the levels of PCSK9 and LDLR. The treatment with IL-1β increased the levels of PCSK9 and LDLR in both GO and non-GO tissues in a time-dependent manner. Representative gel images are shown. Data in the columns indicate the mean density ratio \pm SD, normalized to the level of β-actin in the same sample (*p < 0.05 vs. 0 h in each group).

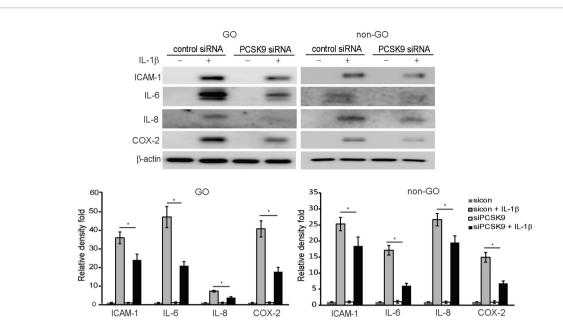


FIGURE 3 | Effect of silencing proprotein convertase subtilisin/kexin type 9 (PCSK9) on the expression of pro-inflammatory cytokines protein. Confluent fibroblasts obtained from Graves' orbitopathy (GO) patients (n=3) were treated with either control siRNA or PCSK9 siRNA (50 nM, 24 h). Then, they were challenged with 10 ng/ml of interleukin-1β (IL-1β) and compared to non-IL-1β-treated counterparts. Western blot analyses were conducted to compare the levels of pro-inflammatory cytokines, intercellular adhesion molecule 1 (ICAM-1), IL-6, IL-8, and cyclooxygenase-2 (COX-2). The same experiment was repeated with fibroblasts obtained from non-GO subjects (n=3). Representative gel images are shown. The mean density ratio \pm SD from fibroblasts were normalized to the level of β -actin in the same sample (*p < 0.05 between sicon + IL-1β and siPCSK9 + IL-1β; sicon, control siRNA; siPCSK9, PCSK9 siRNA).

with TBII (r = 0.576, p < 0.001, n = 44; **Figure 6B**) and CAS (r = 0.631, p < 0.001, n = 44; **Figure 6C**).

DISCUSSION

In this study, we examined the role of PCSK9 in GO pathogenesis. GO tissues showed increased expression of PCSK9, LDLR, and $HIF-1\alpha$. The IL-1 β challenge and adipogenic stimulation led to the increase of PCSK9 and LDLR. PCSK9 silencing with siRNA significantly decreased pro-inflammatory cytokines production, oxidative stress-related proteins, adipogenic transcription factors, and adipocyte differentiation. Importantly, the plasma level of PCSK9 was elevated in GO patients when compared to the non-GO subjects. It also showed a positive correlation with CAS, a measure of GO activity as well as TBII, a predictor of GO severity (17, 24).

Numerous studies have attempted to clarify the pro-inflammatory roles of PCSK9 in a variety of disorders including atherosclerosis, sepsis, psoriasis, steatosis, and myocardial ischemia (12–14, 25–30). For example, Tang et al. suggested an atherogenic role of PCSK9 as the suppression of PCSK9 expression in *apoE* null mice by means of small hairpin RNA decreased expression of TNF- α , IL-1 β , monocyte chemotactic protein 1, Toll-like receptor 4, and NF- κ B and reduced macrophage infiltration in the atherosclerotic plaques (26). In other *in vivo* and *in vitro* studies, the PCSK9 inhibition

diminished pro-inflammatory cytokines production and macrophage accumulation by inhibiting NF- κ B activation (12, 27, 30). In the context of GO, however, there have been no studies on the role of PCSK9. To the best of our knowledge, this study is the first of its kind to identify the pro-inflammatory properties of PCSK9 in GO. Based on our results, silencing PCSK9 ameliorated inflammation by modulating NF- κ B pathway. Furthermore, the PCSK9 level was higher in GO tissues compared to the control at baseline, and IL-1 β , a key mediator in GO inflammation (18, 22), increased PCSK9 levels more prominently in GO fibroblasts than in the control. Our study suggests that PCSK9 may serve as a therapeutic target for GO inflammation.

Mounting evidence has shown the anti-adipogenic effects of PCSK9 depletion (13, 31–33). Currently, PPARγ and C/EBPβ are believed to be responsible for terminal differentiation of fibroblasts into adipocytes (34). The PPARγ activation leads to the expression of adipogenic markers including leptin and fatty acid synthase (FAS) (35). Upstream of PPARγ, molecules such as E2F1 is thought to be involved (36). Recently, a study on rat models with alcohol-induced steatosis showed that the treatment with alirocumab, a human PCSK9 monoclonal antibody, attenuated expression of FAS and alleviated alcohol-induced lipid accumulation. Moreover, PCSK9 inhibition reduced mRNA expression of E2F1 as well as sterol regulatory element-binding protein (SREBP)-1 and SREBP-2 (13). SREBP-1 and SREBP-2 have been found to regulate cholesterol- and fatty acid metabolism-related genes (37). Ruscica et al. also showed that, in

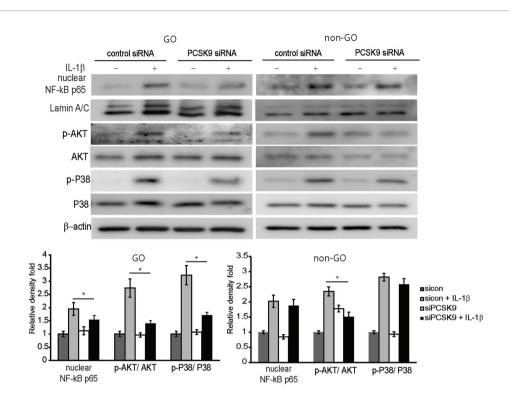


FIGURE 4 | Effect of silencing proprotein convertase subtilisin/kexin type 9 (PCSK9) on the activation of signal molecules by interleukin-1 β (IL-1 β) treatment. Confluent orbital fibroblasts obtained from Graves' orbitopathy (GO) patients (n=3) were treated with or without 10 ng/ml of IL-1 β after transfection with control siRNA or PCSK9 siRNA (50 nM, 24 h). Treatment with IL-1 β (10 ng/ml, 60 min) resulted in an increase in the levels of nuclear NF- κ B p65 and phosphorylated forms of Akt and p38. The treatment with PCSK9 siRNA in GO cells significantly blunted the increases in the transcription factors. However, in fibroblasts from non-GO subjects (n=3), the PCSK9 inhibition only suppressed the phosphorylated Akt. Representative gel images are shown. Data in the columns indicate the mean density ratio \pm SD of the bands obtained from the GO patients, normalized to the level of β -actin in the same sample (*p < 0.05 between sicon + IL-1 β and siPCSK9 + IL-1 β).

201 patients with suspected nonalcoholic steatosis, hepatic PCSK9 mRNA levels were correlated with hepatic SREBP-1 and FAS expression (31). These results are consistent with the those of our own study; the PCSK9 inhibition attenuated adipogenesis as identified by Oil Red O and blunted the expression of PPARγ, C/EBPβ, and leptin. Furthermore, the PCSK9 expression gradually increased throughout the adipogenic differentiation. In this study, we present the evidence for the adipogenic role of PCSK9 in GO. However, whether PCSK9 directly modulates PPARγ or C/EBPβ activity needs to be further investigated. Furthermore, as PPARγ agonists have recently been suggested to modulate helper T cell-related chemokine production in GO (38), further studies are needed to examine if PCSK9 intervention modulates PPARγ-mediated inflammation.

Adipogenesis is thought to be closely associated with oxidative stress (39), and both are found by our study to be significantly suppressed by the PCSK9 inhibition. Multiple studies have proven the anti-oxidative effects of PCSK9 suppression in several disorders including atherosclerosis and myocardial infarction (14, 40). Locally produced reactive oxygen species (ROS) leads to the oxidation of LDL and contributes to atherogenesis (41). Transfection of endothelial cells and vascular

smooth muscle cells with PCSK9 siRNA substantially decreased the production of ROS (40). In several in vitro studies, PCSK9 inhibition reduced ROS generation, while PCSK9 overexpression produced the opposite results (14, 42-44). In an in vivo study, PCSK9 knockout mice expressed significantly less NADPH oxidase and subsequently less ROS in aorta (40). These results are in line with those of our study. As antioxidants, Trx, MnSOD, and HO-1 are induced by oxidative stress and protect tissues from oxidative injuries (45-47). Under hypoxia, HIF-1 α is activated to increase the expression of genes involved in adipogenesis and tissue remodeling in GO (39). Our results demonstrated that PCSK9 inhibition blunted the level of oxidative stress-related proteins which was induced by adipocyte differentiation. Previously, we have found that quercetin inhibits cigarette smoke extract-induced adipogenesis in GO fibroblasts by reducing ROS (48). Additionally, we have reported several molecules with anti-oxidative properties such as resveratrol, caffeine, and curcumin suppress adipogenesis in GO fibroblasts (49–51). Given that oxidative stress contributes to the proliferation of orbital fibroblasts (52), the impeded proliferation of GO fibroblasts by PCSK9 inhibition demonstrated in this study may be attributed to the anti-oxidative property of the PCSK9 inhibitor.

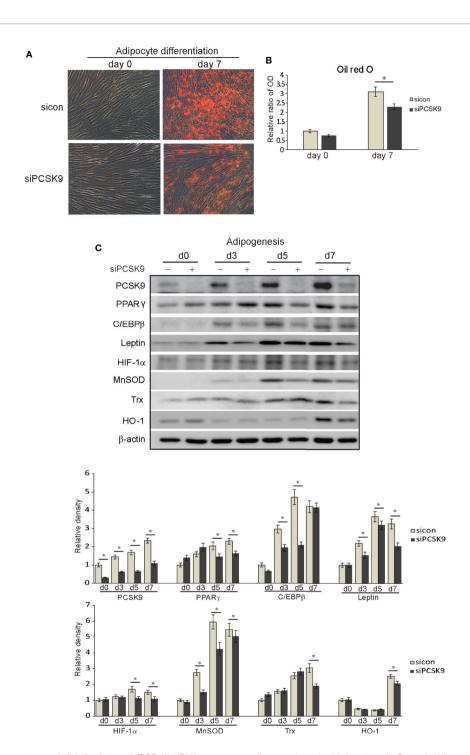


FIGURE 5 | Proprotein convertase subtilisin/kexin type 9 (PCSK9) siRNA suppresses adipogenesis and oxidative stress in Graves' orbitopathy (GO) fibroblasts. Orbital fibroblasts from GO (n=3) patients were cultured in adipogenic medium to induce differentiation into adipocytes for 7 days. (A) Oil red O staining showed treatment with PCSK9 siRNA (50 nM, 24 h) attenuated adipogenesis. (B) Quantification by measurements of optical density of cell lysates at 490nm echoed the histochemical results. The results are presented as the mean optical ratio ± SD (*p < 0.05 between sicon and siPCSK9). (C) Western blot analyses showed that throughout the 7-day period of adipogenesis, the cell lysates of fibroblasts collected at different time points showed a gradual increase in production of PCSK9, which was markedly decreased when PCSK9 siRNA (50 nM, 24 h) was treated. The levels of adipogenic transcription factors, peroxisome proliferator-activated receptor gamma (PPAR*) and CCAAT/enhancer-binding protein β (C/EBPβ), were substantially curtailed in fibroblasts transfected with PCSK9 siRNA. The levels of mature adipocyte marker, leptin, were also significantly reduced in the PCSK9 siRNA-treated group. The levels of antioxidants, manganese superoxide dismutase (MnSOD), thioredoxin (Trx), and heme oxygenase-1 (HO-1), and oxidative stress-related protein, hypoxia-inducible factor 1α (HIF-1α) were significantly decreased in the PCSK9 siRNA-transfected group. Data in the columns indicate the mean density ratio ± SD, normalized to the level of β-actin in the same sample, and representative gel images are shown (*p < 0.05 between sicon and siPCSK9 on days 0, 3, 5, and 7 of adipogenesis).

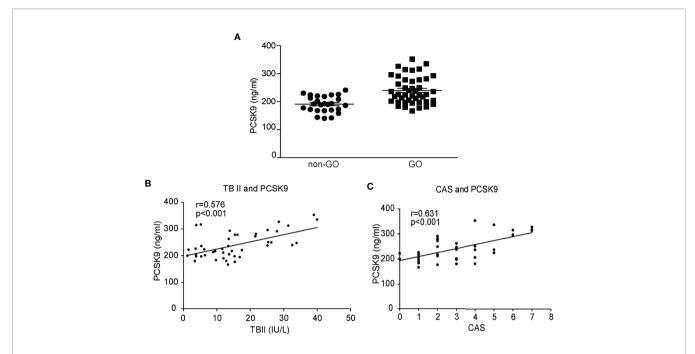


FIGURE 6 | Comparison of plasma levels of proprotein convertase subtilisin/kexin type 9 (PCSK9) between Graves' orbitopathy (GO) patients and non-GO subjects, and correlation analyses between plasma levels of PCSK9, thyrotropin-binding inhibitor immunoglobulin and clinical activity score (CAS). The plasma levels of PCSK9 were measured in GO patients and non-GO subjects using ELISA. The samples were assayed in triplicate. **(A)** The mean PCSK9 plasma level was significantly higher in the GO patients (n=44, 239.97 \pm 48.20 ng/ml) than the healthy subjects (n=26, 190.83 \pm 28.77 ng/ml; p < 0.01). A single dot represents the value obtained from a single donor. The results of Spearman's rank correlation test between plasma levels of PCSK9 (GO patients, n=44) and **(B)** plasma levels of thyrotropin binding inhibitory immunoglobulin (TBII) (GO patients, n=44) or **(C)** clinical activity score (CAS) (GO patients, n=44) are shown. The plasma PCSK9 concentrations showed significant associations with the plasma TBII levels (r = 0.576, p < 0.001) and CAS (r = 0.631, p < 0.001).

The changes in the level of phosphorylated forms of transcription factors upon the transfection of PCSK9 siRNA in GO fibroblasts indicate that a complex network of signaling pathways may exist. In this study, the PCSK9 inhibition decreased the activation of Akt and attenuated adipogenesis in GO fibroblasts. Our study with siRNA transfection is limited by the lack of mRNA data, which would have provided additional insights to its effect at transcriptional or translational level. However, like other studies that have employed similar methods (53, 54), we believe that we have shown a concrete evidence with western blot and ELISA that PCSK9, regardless of the mechanism with which its level is modified, affects inflammation and adipogenesis in GO. In line with our study, Kumar et al. have previously reported that an autoantibody against TSH receptor stimulated the phosphoinositide 3-kinase (PI3K)/Akt pathway and induced adipogenesis of orbital preadipocytes in GO (55). They asserted that this pathway triggered the terminal stages of adipogenesis. Another recent study by our group has presented that an Akt inhibitor suppressed adipogenesis in GO orbital fibroblasts (56). Another downstream effector of PI3K/Akt pathway, Forkhead box O (FOXOs), has also been demonstrated to repress excessive adipogenesis and hyaluronan overproduction in GO fibroblasts (57). As multiple studies, including our own, continue to highlight the importance of the PI3K/Akt pathway in the GO

pathogenesis, further studies are necessary to identify the interaction between the signaling molecules as well as PCSK9.

Finally, we show that plasma PCSK9 levels were significantly higher in active GO patients than inactive GO patients as well as healthy subjects. These results, along with the higher PCSK9 mRNA levels in GO tissues, strongly suggest the involvement of PCSK9 in GO pathogenesis. Moreover, the plasma PCSK9 level revealed a strong correlation with plasma TBII level as well as CAS. Its role in many other inflammatory and autoimmune diseases such as systemic lupus erythematosus, rheumatoid arthritis, and type 1 diabetes demonstrates its complex biological activity (58–60). Identifying new biomarkers and therapeutic targets such as PCSK9 can further our knowledge of these disorders and lead to the development of effective treatments.

In conclusion, we demonstrated that PCSK9 inhibition countered pro-inflammatory cytokines production, oxidative stress-related proteins, adipogenic transcription factors, and adipocyte formation in GO fibroblasts. The PCSK9 level was increased during the IL-1 β challenge and adipogenic stimulation. The plasma PCSK9 level was elevated in GO patients and positively correlated with clinical inflammation and thyrotropin receptor antibody titer, indicating that PCSK9 is a potential biomarker for diagnosis and prognosis of GO. Further studies are needed to establish the response to PCSK9 inhibitors *in vivo* and explore the use of the inhibitor as an effective therapeutic strategy for GO.

DATA AVAILABILITY STATEMENT

The raw data supporting the conclusions of this article will be made available by the authors, without undue reservation.

ETHICS STATEMENTS

The studies involving human participants were reviewed and approved by Institutional Review Board of Severance Hospital. The patients/participants provided their written informed consent to participate in this study.

AUTHOR CONTRIBUTIONS

GL drafted the manuscript; JiK acquired and interpreted the data. JL, JaK, and EL revised the work. JY designed the study and revised the work. All authors contributed to the article and approved the submitted version.

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SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at: https://www.frontiersin.org/articles/10.3389/fendo.2020. 607144/full#supplementary-material

SUPPLEMENTARY FIGURE 1 | siRNA-mediated knockdown of PCSK9 in GO and non-GO fibroblasts. GO (n=3) and non-GO (n=3) orbital fibroblasts were transfected with PCSK9 siRNA (50, 100 nM) for 24 h. As shown in western blot results, the level of PCSK9 was significantly decreased following RNA interference. β -actin was included as a loading control. NC, normal control.

SUPPLEMENTARY FIGURE 2 | Effect of PCSK9 siRNA on viability of GO fibroblasts. Orbital fibroblasts of GO patients (n=3) were seeded in 24-well culture plates, 1×10^5 cells per well. PCSK9 siRNA (50 nM) were applied to wells for 10, 24, and 48 h and MTT assay was conducted to test cell viability. Results are presented as mean \pm SD. Assays were carried out in triplicate and repeated at least three times. The proliferation of GO fibroblasts was impeded in PCSK9 siRNA-treated fibroblasts compared to the control.

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Conflict of Interest: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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Selenium in the Treatment of Graves' Hyperthyroidism and Eye Disease

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Based on the role of oxidative stress in the pathogenesis of Graves' hyperthyroidism (GH) and Graves' Orbitopathy (GO), a therapy with the antioxidant agent selenium has been proposed and a number of studies have been performed, both *in vitro* and *in vivo*. In GH, reactive oxygen species (ROS) contribute to the thyroid and peripheral tissues damage. In GO, tissue hypoxia, as well as ROS, are involved in the typical changes that occur in fibroadipose orbital tissue and the perimysium of extraocular muscles. Antioxidants have been proposed to improve the effects of antithyroid drugs in GH patients, as well as the remodeling of orbital tissues in patients with GO. Here, we reviewed the literature on the possible beneficial effects and clinical use of selenium in the management of patients with GH and GO. A randomized clinical trial on the use of selenium in patients with mild GO provided evidence for a beneficial effect; no data are available on more severe forms of GO. Although the real effectiveness of selenium in patients with GH remains questionable, its use in the management of mild GO is generally believed to be beneficial, and selenium administration has been included in the clinical practice for the patients with mild eve disease.

Keywords: Graves' hyperthyroidism, Graves' orbitopathy, oxidative stress, reactive oxygen species, selenium, selenocysteine, selenoproteins

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INTRODUCTION

Selenium has been proposed for the management of thyroid diseases, including Graves' disease (GD), Graves' Orbitopathy (GO), and chronic autoimmune thyroiditis (1–7). The efficacy of selenium in GD and GO is based on its anti-oxidant properties, being oxidative stress involved in the pathogenesis of both conditions (8–12).

GD is an autoimmune disease with a prevalence of ~1% (13), which affects mainly the thyroid. Extrathyroidal manifestations, including GO, pretibial myxedema, and acropachy, can be observed to various extents (13–17). The major pathogenetic mechanism of GD is the stimulation of the thyroid stimulating hormone receptor (TSH-R) by autoantibodies that bind to it and promote thyrocyte proliferation and activity, resulting in thyroid hyperfunction (13). The symptoms of thyrotoxicosis are often nonspecific, so patients with GD may present in various ways. Heat intolerance, tachycardia, inappropriate feelings of anxiety and apprehension, hyperactivity and weight loss are common. A diffuse goiter can be visible or palpable, with a systolic phase bruit found over it. Systolic blood pressure may be elevated, and hepatomegaly or splenomegaly may be observed (16). The complex pathogenetic interplay of GD includes, among others, oxidative stress (16–18), because of which selenium and other antioxidant agents have been proposed for the

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management of Graves' hyperthyroidism (GH). In this paper we review the role of oxidative stress in GH, as well as the most significant studies on the use of selenium in the management of patients with GH and GO. The molecular mechanisms by which the disruption of the cell redox state plays a role in the pathogenesis of GO, as well as the potential beneficial effect of other antioxidant agents in patients with GO, are largely discussed in another review article in this issue of the journal.

CELL REDOX STATE AND EVALUATION OF OXIDATIVE STRESS

The balancing of the cell redox state is a key point in cellular homeostasis. Reactive oxygen species (ROS), including hydroxyl radicals (OH⁻), hydrogen peroxide (H₂O₂), superoxide anions (O²), and lipid peroxides, are molecules highly reactive because of the presence of unpaired electrons (19, 20). The increase in ROS production, as well as the decrease in their elimination, can disrupt the balance of the cell redox state, resulting in oxidative stress and breaking of cellular homeostasis (19, 20). As oxidizing agents, ROS interfere with intracellular functions, being capable of damaging various cellular elements, including cell membranes, proteins, lipids and nucleic acids, and ultimately resulting in mitochondrial dysfunction and loss of enzymatic activity (19, 20). Under physiological conditions, antioxidant agents, namely glutathione (GSH), superoxide dismutase (SOD), glutathione peroxidase (GPX) and catalase, act as ROS antagonists, therefore contributing to the maintenance of the cell redox state (21).

The oxidant/antioxidant cell balance can by evaluated by several plasma and/or erythrocyte markers (20, 21). In healthy subjects, markers of oxidative stress, as well as of the antioxidant system, are lowly represented. The induction of an oxidative stress state promotes two different reactions in the cells, depending on how long it lasts. Initially, an increase in oxidative markers is accompanied by an increase in antioxidant markers, the latter being a homeostatic response aimed at avoiding cell damage. A long-lasting state of oxidative stress is however characterized by low levels of antioxidant markers, because of exhaustion of the antioxidant cell systems (20, 21).

OXIDATIVE STRESS IN GH

Thyrotoxicosis is a hypermetabolic state characterized by high consumption of intracellular ATP and oxygen, and by dysfunction of the mitochondrial respiratory chain, which leads to saturation of the physiological antioxidant systems, resulting in an uncontrolled production of ROS in peripheral tissues as well as in the thyroid (22–25). Studies in thyrotoxic rats suggest that the oxidative stress may be involved in the pathogenesis of thyrotoxic myopathy and cardiomyopathy (26) and *in vitro* studies have shown that exposure of thyroid cell to H₂O₂, particularly when made selenium-deficient,

induces DNA double strand breaks, apoptosis, necrosis and mutagenesis (24). On this basis, we might speculate that the oxidative stress might exert a dual action in patients with GH. The ROS-induced damage of thyroid epithelial cells could lead to an increased release of autoantigens and production of TSH-R autoantibodies (TRAb), and the peripheral tissue damage might contribute the clinical manifestations of hyperthyroidism.

Animal Studies

Studies in animal models suggest that thyrotoxicosis, induced by the administration of triiodothyronine (T3) or thyroxine (T4), promotes oxidative stress as well as the response of the physiological antioxidant machinery (22–25). It has been demonstrated that MDA, is significantly higher in thyrotoxic than in control and euthyroid rats (25, 26). Moreover, the oxidative stress induced by thyrotoxicosis promotes an increase in erythrocyte antioxidant parameters, namely SOD and GPX. Interestingly, the administration of Vitamin E, which has antioxidant properties, reduces MDA, SOD and GPX in hyperthyroid rats (26), and it seems also to be protective against a thyroxine-induced increase of lipid peroxidation in cardiac and skeletal muscles (27).

Human Studies

The most relevant clinical studies on the role of oxidative stress in thyroid diseases have been performed in GH patients, either with uncontrolled hyperthyroidism, or after restoration of euthyroidism with antithyroid drugs (ATD) or radioiodine (28-33). It has been reported that the levels of oxidative stress markers, including hydrogen, lipid peroxides, MDA and thiobarbituric acid-reacting substances (TBARS) in the serum, plasma and erythrocytes of hyperthyroid patients were higher than in euthyroid subjects (28-33). Moreover, a correlation between serum thyroid hormones and lipid peroxidation products was observed in hyperthyroid patients (28-33). These findings do not seem to be specific for GH, but rather for hyperthyroidism in general, having been observed also in patients with subclinical hyperthyroidism due to multinodular goiter (31). The evaluation of the antioxidant defense system in thyrotoxic patients resulted in conflicting findings (10, 28, 29, 33). Komosinska-Vassev et al. reported higher levels of SOD, CAT and GPX in the erythrocytes of GH patients than in age-matched controls. However, no differences in serum glutathione reductase (an antioxidant enzyme involved in GSH synthesis) and in the total antioxidant status were found, likely suggesting rapid exhaustion of the antioxidant system (10). Bednarek et al. reported an increase in plasma SOD and catalase in patients with GH of short duration (1-2 months) compared with healthy subjects, but not of GPX and glutathione reductase, which were, conversely, decreased (29). On the contrary, Aslan et al. reported a significant reduction of the total antioxidant activity in 36 hyperthyroid patients with an average duration of hyperthyroidism of 2.3 ± 1.5 months compared with controls

(33). Finally, a study involving 69 GD patients with hyperthyroidism lasting more than 6 months showed decreased levels of erythrocyte SOD and catalase activities compared with controls, with no difference in erythrocyte GPX and total antioxidant activities (28). The duration of the hyperthyroid status at the time of the evaluation can explain, at least in part, the conflicting data on antioxidant activity in patients with GD. Probably, patients with hyperthyroidism of longer duration have exhausted the antioxidant defense system, which leads to the reduction of the antioxidant capacity (34, 35). It has been also reported that the monoclonal thyroid stimulating antibody M22 as well as polyclonal serum thyroid stimulating antibodies (TSAbs) from patients with GH, promote ROS generation and lipid peroxidation in HEK cells that stably overexpressed the human TSHR (HEK-293 TSHR) (36). ATD reduce the levels of oxidative stress markers, thereby improving the activity of the intra- and extracellular antioxidant defense systems, due to restoration of euthyroidism and possibly to their antioxidant properties (28-30, 32). As discussed in another review on the same issue of this journal, oxidative stress also plays a role in GO, and the beneficial effect of antioxidant agents are supported by in vitro and human studies (1-6, 11, 37-46).

SELENIUM

Selenium is a trace mineral which acts following incorporation as selenocysteine into selenoproteins, among which thioredoxin reductases (TRs), GPX, and iodothyronine deiodinases (D1, D2 or D3) are the best known (3-5, 47). In this regard, D1-3 are homologous proteins consisting of 250-300 amino acids, with a single transmembrane domain located at the N-terminus. They are involved in the reductive deiodination of thyroid hormones. The most remarkable feature of all three deiodinases is the presence of a selenocysteine residue in the center of the amino acid sequence. D1, encoded by DIO1 gene, is predominantly expressed in the liver, kidney and thyroid. It catalyzes the outer and inner ring deiodination of iodothyronine derivatives, with a preference for reverse T3 (rT3) and iodothyronine sulfates. D2, encoded by DIO2 gene, is primarily expressed in the thyroid, brain, anterior pituitary and brown adipose tissue, where contributes local conversion of T4 in T3. Moreover, the pituitary enzyme is involved in the negative feedback regulation of TSH and TRH secretion. The DIO3 gene codes for D3, which is detected in brain, skin, liver, intestine, placenta and pregnant uterus, where it catalyzes the inactivation of T4 and T3. Since D1-3 are selenoproteins, selenium deficiency would be expected to result in their reduced activities in different tissues, although this effect is observed only for D1 in the liver and kidney and not for D2 and D3 in other tissues (48).

Selenoproteins have antioxidant and enzymatic capacity (47) and, in the thyroid, where they are highly expressed, influence the balance of the cell reduction-oxidation activities

(49). Furthermore, selenoproteins are essential for activated Tcell function, being involved in the proliferation of T-cells in response to T-cell receptor stimulation. It has been also reported that selenium supplementation promotes differentiation of CD4+ T cells into T-helper-1 (Th1) rather than T-helper-2 (Th2) effector. Among the immunomodulating functions, selenoproteins increased cytotoxic lymphocyte-mediated tumour cytotoxicity and natural killer activity (50). It has been also reported that selenium may regulate the inflammatory response by reducing the release of tumor necrosis factor α (TNFα) and cyclooxygenase (COX2), as well as the activation of nuclear factor κB (NF-κB) which is one of the transcriptional factors involved in the immune and pro-inflammatory response. Elevated selenium levels increase GPX, which inhibits $I\kappa B-\alpha$ phosphorylation and consequently the translocation of NFκB (51).

Dietary sources of selenium include meat, seafood, shellfish, offal, eggs and cereals (47, 49). The bioavailability of selenium varies depending on the type of food and content in the soil for growing crops and fodder. Furthermore, additional factors, such as selenium speciation, soil pH or the presence of ions complexed with selenium, play a role in the bioavailability of the mineral (47). Selenium dietary intake is followed by absorption in the gastrointestinal tract and transport to the liver, where it is incorporated into selenoglycoproteins containing up to 10 selenium residues per molecule. Selenoglycoproteins finally reach peripheral tissues, where their concentration is proportional to the degree of oxidative stress (47–50).

The measurement of total serum selenium concentration or circulating selenoproteins, including GPX-3 and selenoprotein P, can be used to evaluate the selenium status, which is variable depending on the geographical area, being high in North America and relatively low in most European countries, in particular in the Eastern Europe (50-52). The Office of Dietary Supplements has established that the recommended daily allowance (RDA) is 55 µg in men and women. Selenium intake in several European countries is lower than in the USA, because of its lower content in the soil. Values range from around 30 µg in UK, Germany, Sweden and Slovakia to around 70 in Netherlands and Switzerland (47, 50). To properly interpret these data, we need to have appropriate standards against which to compare them. There is no consensus on this issue. The UK reference nutrient intake of 75 µg per day in men and 60 in women has been determined as the amount needed to maximize the activity of the GPX in plasma (which occurs at a selenium concentration of about 95 µg/L). Current UK intake are about half the value. On the other hand, selenium excess has been suggested to increase the risk of type 2 diabetes and of malignancies (50, 52). However, the issue is still matter of debate and no adverse events for selenium doses not exceeding 200 µg/day were reported by a number of studies, including one in which relatively high levels of selenium were reached with supplementation (~190 μg/L after 90 days of treatment) (3, 50). It is therefore possible that selenium might lead to subclinical alterations, for example on glucose metabolism, in selected patients, with no real clinical impact (1, 4, 53, 54).

In this regard, several trials have shown a benefit of selenium supplementation on the risk of cancer incidence and death, as well as the risk of developing hyperglycemia. However, two large trials performed in USA have shown that these benefits are no longer seen in patients whose baseline serum selenium concentration was above 122 µg/L, in whom, conversely, these risks were increased. On this basis, Rayman proposed that 122 µg/L represents the concentration of baseline serum selenium that "delineates a change in risk, from lower to higher, of developing cancer and type 2 diabetes in patients treated with selenium supplementation of 200 µg per day" (50). This observation may be relevant for individuals living in countries, like the USA, where a large proportion of people has a serum selenium concentration above this cut-off. Therefore, it has been proposed to measure serum selenium concentrations before administering the supplement, in order to avoid overdosage in subjects with baseline serum concentrations higher than 122 µg/L (53). In our opinion, the measurement of serum selenium before supplementation might be useful in individual living in countries with high selenium intake, like the USA, to avoid the risk of potential disadvantages, but not individuals living in countries, like most European countries, where dietary selenium intake is low and high concentrations of serum selenium are very unlikely to be achieved with supplementation.

Selenium supplements contain sodium selenite or selenomethionine, and the main difference between them is that, after saturation of selenoproteins, selenite is excreted whereas selenomethionine can further increase serum selenium through its incorporation into proteins (1, 4). Consequently, the effect of selenite is strictly linked to the individual state of selenium, whereas using selenomethionine as a supplement, the concentration of plasma selenium increases also in subjects with a sufficient selenium concentration to begin with.

USE OF SELENIUM IN GH

Based on the above described role of oxidative stress in GH and the antioxidant activity of selenium, clinical studies were performed to investigate its effect in GH patients. The hypothesis is that selenium deficiency could enhance oxidative stress in thyrotoxicosis, by worsening the antioxidant machinery in response to ROS. To address this issue, a number of studies have tested selenium, either alone or within a mixture containing other antioxidant agents, in GH patients treated with ATD (2, 3, 55-57) (Table 1). These investigations did not provide clear-cut results. Vrca et al. performed a study in Croatia, an area characterized by moderate selenium deficiency, randomizing GH patients to therapy with methimazole (MMI) plus an antioxidant mixture containing β-carotene, Vitamin C and Vitamin E and a relatively low dose of selenium (60 µg), or MMI alone (55). The main difference between the two groups was that euthyroidism was more rapidly reached in the former group, likely reducing the period of exposure to oxidative stress. Moreover, a better response to treatment in terms of improvement of LDL-cholesterol levels was found in patients treated with the antioxidant mixture compared with those given MMI alone (55). Similar date on thyroid function were reported by Guerra et al. in patients given a mixture containing low doses of selenium (15 μ g/day), compared to MMI alone (57).

Three studies were performed using "pure" selenium rather than a mixture of antioxidants (2, 3, 56) (Table 1). In one of these studies (56), hyperthyroid, selenium-deficient patients were randomized to receive a block-and-replace regimen (MMI plus levothyroxine), plus 200 µg/day of selenium or placebo. A slightly better control of hyperthyroidism was observed in the selenium group (56). Two additional randomized clinical trials performed in GH patients treated with MMI alone did not show beneficial effects on thyroid function and peripheral manifestations of hyperthyroidism in patients given selenium compared with placebo (2, 3). The conflicting results between the first study and the subsequent two can be explained, at least in part, by the different levels of baseline selenium, which was insufficient only in the first one. These findings support the main hypothesis of a role of ROS in GH, which might be relevant especially in patients with selenium deficiency. Thus, a supplementation with selenium can be considered in these cases, whereas, in our opinion, there is no sufficient evidence to recommend the routine addition of selenium to ATD in the management of all patients with GH.

USE OF SELENIUM IN GO

Over the last few years, in addition to the most common treatment for the management of moderate-to-severe and active GO, namely high dose oral and intravenous glucocorticoids (GC), orbital irradiation and surgical procedures (58), other medications have been proven effective to various extents, including rituximab (59), the recently FDAapproved teprotumumab (60, 61), mycophenolate (62), and tocilizumab (63). Until recently, the treatment of mild GO was typically limited to local measures (19 but ~15% of patients GO progress to an extent that requires specific treatments (13, 58). In general, major treatments are not recommended for mild GO, unless there is a sufficient impairment in the quality of life, which justifies the risk of GC-related adverse events (58, 64-66). In view of the role of oxidative stress in GO, antioxidant supplements have been proposed as a possible therapeutic approach, and basic and clinical studies have investigated the effect of selenium, due to its antioxidant and immunomodulating actions.

In Vitro Studies

A couple of *in vitro* studies have provided evidence for a beneficial effect of selenium in primary cultures of orbital

fibroblasts (OFs) (38, 65, 66) (**Table 2**). In a first study (38), after induction of oxidative stress by treating OFs with H_2O_2 , GPX activity, glutathione disulfide (GSSG), cell proliferation, HA and pro-inflammatory cytokines were measured, in the presence or absence of selenium-(Methyl)-selenocysteine (SeMCys). H_2O_2 induced oxidative stress in OFs, reflected by a dose-dependent increase in GSSG, a known measure of cell response to ROS (**Figure 1**). SeMCys reduced the effects of H_2O_2 , providing evidence for an antioxidant action of selenium in OFs. The effect of selenium was observed in OFs from both GO patients and control subjects, which should not be seen as a limitation for its clinical use. Thus, in the same study the authors reported that proliferation of OFs, as well as the

production of HA, were significantly greater in GO than in control fibroblasts. Moreover, SeMCys significantly reduced cell proliferation and hyaluronic acid (HA) release in GO OFs, whereas no effects were observed in control OFs (38). These findings might suggest that fibroblasts from GO patients are somehow different, leading to a different response to oxidative stress and selenium activity.

In a subsequent study, the same authors reported a dual effect of H_2O_2 on cell proliferation, depending on dose used (67). At low concentrations, H_2O_2 promoted cell proliferation, whereas at high concentration progressively decreased cell vitality and cell proliferation. The effects of both high and low dose H_2O_2 were inhibited by SeMCys which, interestingly,

TABLE 1 | Selenium in Graves' Hyperthyroidism (GH).

| Reference | Journal and years | Type of study | Methods | Dosage of compounds used | Results | | |
|-----------------------------|--|---|--|---|---|--|--|
| Vrca VB. et al. (53) | Acta Pharm. 2012 | Prospective, placebo- controlled, clinical trial | Fifty-five patients with newly diagnosed GH were randomized to receive methimazole alone or methimazole plus a fixed combination of antioxidant agents | The mixture of antioxidants contained: | Beneficial effect on GH Better and faster normalization of thyroid function was observed in patients with GH treated with methimazole plus the mixture of antioxidant agents compared with control group. In addition, a better response to treatment, in terms of improvement of LDL-cholesterol levels, was found in patients treated with the antioxidant mixture compared with those given methimazole alone | | |
| Calissendorff J. et al. (2) | Eur Thyroid J 2015 | Prospective, placebo- controlled, clinical trial | Thirty-eight consecutive patients with newly diagnosed and untreated GH were randomized to receive "block and replace" treatment with methimazole and levothyroxine, plus Selenium or alone for 9 months | Selenium (200 μg daily orally) | Beneficial effect on GH Patients treated with selenium had a better control of hyperthyroidism: at 18 weeks, the serum levels of FT4 were lower in the seleniur group compared to the placebo (14 vs. 17 pmol/l group, p = 0.01). Similar results were observed also at 36 weeks (15 vs. 18 pmol/l, p = 0.01). In accordance, the TSH levels increased more in the selenium group at 18 weeks (0.05 vs. 0.02 mIU/l, p = 0.04) | | |
| Leo M. et al. (3) | J Endocrinol Invest 2017 | Prospective, placebo- controlled, clinical trial | Thirty consecutive patients with untreated GH were randomized to receive Methimazole plus Selenium vs methimazole alone for 3 months | L-seleno-methionine (166 μg daily orally) | No beneficial effect on GH. Administration of Methimazole leads to the normalization of FT3 and FT4, with no difference between groups. Serum levels of malondialdehyde, a marker of oxidative stress was similarly high in the two groups and decreased significantly after therapy, with no difference between groups. The results suggested that selenium had not significant effect on short-term control of hyperthyroidism | | |
| Kahaly GJ. et al. (54) | (54) Metab. 2017 placebo-consecutive controlled, with untrear clinical trial were rando receive Met plus Selenium o methimazol | | consecutive patients with untreated GH were randomized to receive Methimazole | Sodium selenite (300 μg daily orally) | No beneficial effect on GH. The response to treatment, in terms of thyroifunction normalization of thyroid hormones, was very similar in the two groups at week 2. During a 12-week follow-up, GH relapsed in 48% of patients included in the selenium group and in 44% of patients of the placebo group. Serum concentrations of Selenium an selenoprotein P were unrelated to response or recurrence rates. At week 36, 12 of 29 patients (41%) and 15 of 33 patients (45%) were responders and still in remission in the selenium and placebo groups, respectively. | | |

TABLE 2 | Selenium in Graves' Orbitopathy (GO).

| Reference | Journal and years | Type of study | Methods | Dosage of compounds used | Results | |
|--------------------------------|----------------------|--|---|---|--|--|
| Rotondo Dottore G. et al. (65) | Thyroid 2016 | In vitro study | Primary cultures of orbital fibroblasts from 6 GO patients and 6 control subjects were obtained. To induce oxidative stress, cells were incubated for 24 h with medium containing H ₂ O ₂ at various concentrations. The primary objective was to assess the effects of selenium in GO fibroblasts. | Cells were pre-incubated for 2 days with medium without compounds or containing Semethyl-selenocysteine hydrochloride (SeMCys) or, as control, methyl-cysteine at various concentrations. Cell proliferation, hyaluronic acid (HA) and pro-inflammatory cytokines production were measured | Selenium reduced proliferation and release of HA and cytokines in GO fibroblasts. H ₂ O ₂ promoted an increased release of glutathione disulfide (GSSG), a marker of oxidative stress, and of fibroblast proliferation, which were reduced by selenium. H ₂ O ₂ promoted the production of cytokines involved in GO pathogenesis, namely TNF- α , IL1- β and IFN- γ . The increase in TNF- α and IFN- γ was rescued by selenium. Whereas the effects of selenium were similar in GO and control fibroblasts concerning oxidative stress and cytokines, they were exclusive to GO fibroblasts concerning proliferation and HA production. | |
| Rotondo Dottore G. et al. (66) | Endocrine. 2017 | | Primary cultures of orbital fibroblasts from 6 patients with long lasting inactive GO and 6 control subjects were obtained. To induce oxidative stress, cells were incubated for 90 min at 37°C with medium containing H_2O_2 50 μ M. The aim of the study was to evaluate the effects of selenium in GO | Cells were preincubated for 2 days at 37°C with medium without compounds containing SeMCys or, as control, methyloysteine (MCys) at a 10 μM concentration. Cell vitality, lactate dehydrogenase production (as a measure of cell necrosis) and apoptosis were measured | Selenium reduced cell damage in GO fibroblasts SeMCys rescued from H_2O_2 -dependent cytotoxicity, by reducing necrosis and apoptosis, with no difference between GO and control fibroblasts. MCys had no effect. To determine whether the findings reflected the antioxidant actions of selenium, the assessment of GSSG was performed. H_2O_2 promoted the increased production of GSSG, which was counteracted by SeMCys, but not by MCys, with no differences between GO | |
| Marcocci C. et al. (4) | N Engl J Med 2011 | Prospective, multicenter, placebo- controlled clinical trial | fibroblasts. One hundred fifty-nine patients with mild GO were randomized to receive sodium selenite or compared with placebo for 6 months | Sodium selenite (100 μg twice daily orally), or pentoxifylline (600 mg twice daily orally) | and control fibroblasts. Beneficial effect of selenium in mild ${\bf GO}$. Compared with placebo, Selenium, but not pentoxifylline, leads to the improvement of quality of life (P < 0.001), eye overall outcome (P = 0.01) and slowed the progression of ${\bf GO}$ (P = 0.01), at the 6 months evaluation. | |

inhibited HA synthesis in GO, but not in control fibroblasts, even though $\rm H_2O_2$ did not affect HA release (67). This observation suggests that selenium may influence HA release, at least in part, regardless of the oxidative stress induced by $\rm H_2O_2$, through mechanisms that are still to be clarified. Furthermore, a reduction in the release of pro-inflammatory cytokines induced by low dose $\rm H_2O_2$, namely interferon- γ (IFN γ) and TNF α , was found in both GO and control OFs treated with SeMCys (67, 68), contributing the beneficial effects of selenium in GO OFs.

Clinical Studies

The first clinical, pilot study on antioxidants, showed an improvement of GO soft tissue involvement in patients treated with an antioxidant mixture containing allopurinol plus nicotinamide, compared with those given placebo (54). After these promising results, the European Group on Graves' Orbitopathy (EUGOGO) performed a randomized, placebo-

controlled, multicenter, clinical trial in European countries known to have marginal selenium-deficiency, to investigate the effect of selenium and pentoxifylline in mild GO (4). One hundred and fifty-nine patients with mild GO were randomized to receive sodium selenite (100 mcg twice/day, equivalent to 91.3 µg of selenium), pentoxifylline (600 mg twice/day), or placebo (twice/day) for 6 months, followed by a follow-up period of 6 months. The overall eye outcome—assessed at the end of treatment by a change in a composite score including exophthalmometry, clinical activity score (CAS), measurement of eyelid aperture, diplopia and visual acuity—as well as the quality of life were significantly better in the selenium group. GO improved in 61% of patients treated with selenium and in 36% of patients receiving placebo, whereas the eye disease worsened in 7 and 26% of patients in the selenium and placebo group, respectively (Figure 2). The effect of pentoxifylline was not relevant compared to placebo. Similarly, at 6 months, the scores of the quality of life of GO

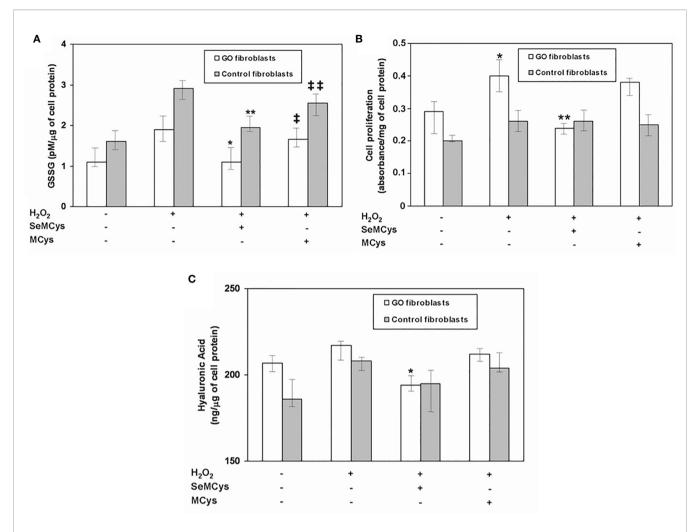


FIGURE 1 | **(A)** Effect of selenium (10 μMol) or of the negative control methylcysteine (MCys) (10 μMol) on Glutathione disulfide (GSSG), released after treatment with H_2O_2 (5 μMol), in fibroblasts from patients with Graves' orbitopathy (GO fibroblasts) or from control patients. P* and **P = 0.02 vs H_2O_2 ; \ddagger and \ddagger P = NS vs H_2O_2 ; P = NS between GO and control fibroblasts; **(B)** Effect of selenium (10 μMol) or methylcysteine (MCys) (10 μMol) on cell proliferation, induced by H_2O_2 treatment (5 μMol), in GO and control fibroblasts. *P = 0.02 vs untreated cells; **P = 0.02 vs H_2O_2 ; P = 0.003 between GO and control fibroblasts; **(C)** Effect of selenium (10 μMol) or methylcysteine (MCys) (10 μMol) on hyaluronic acid (HA) release in GO and control fibroblasts. *P = 0.02 vs H_2O_2 ; P = 0.02 between Graves' Orbitopathy and control fibroblasts. Reprinted with permission by Thyroid 2017, Volume 27, pp. 271–278, published by Mary Ann Liebert, Inc., New Rochelle, NY.

(GO-QOL) increased from baseline by 6 or more points for visual functioning in 33 patients (62%) and for appearance in 40 patients (75%). Interestingly, the majority of selenium-treated patients who had a positive change in eyelid aperture, soft tissue involvement, or both, also had an improvement of 6 points or more on the appearance subscale of GO-QOL [84%; 95% confidence interval (CI), 67 to 95] and on visual functioning subscale (72%; 95% CI, 53 to 86), as well as in the overall score (81%; 95% CI 63 to 93). The GO improvement following treatment with selenium was maintained also at 12 months, suggesting that the beneficial effect of selenium is persistent after treatment withdrawal. A limit of this study is that neither baseline not end of treatment selenium concentrations were not measured. Therefore, it remains unclear whether correction of

selenium deficiency accounted for the beneficial effects of selenium administration.

CONCLUSIONS

The redox state is a key point in cellular homeostasis and its unbalance leads to the disruption of intracellular reactions, thereby damaging cell structures. Oxidative stress plays an important role in both GH and GO but the question of whether the use of selenium in the management of patients with GH and GO can be truly effective is still to be clarified. Studies performed so far regarding the use of antioxidant agents, especially as

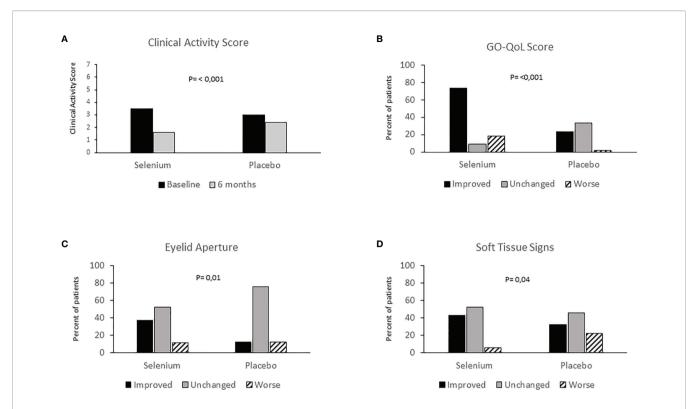


FIGURE 2 | (A) Clinical Activity Score (CAS) evaluation at baseline and 6 months in patients with mild Graves' Orbitopathy treated with selenium or placebo; (B) Graves' orbitopathy-specific quality-of-life questionnaire (GO-QOL) at baseline and 6 months in patients with mild Graves' Orbitopathy treated with selenium, placebo or Pentoxifylline; (C) Eyelid aperture at baseline and 6 months in patients with mild Graves' Orbitopathy treated with selenium, placebo or Pentoxifylline; (D) Soft tissue signs at baseline and 6 months in patients with mild Graves' Orbitopathy treated with selenium, placebo or Pentoxifylline. Redrawn from the Engl J Med 2011; 364: 1920-1931.

selenium, provided conflicting results on the possibility of a new therapeutic approach. The real effectiveness of selenium in patients with GH is very dubious and at this moment there is not enough evidence to use it. Current evidence on selenium supplementation in patients with mild GO provided promising results, suggesting that selenium may improve the quality of life and the course of GO and opening to the possibility of its clinical use. It is also possible that selenium may have beneficial effects in moderate-to-severe GO, especially when patients are selenium-deficient. The action of selenium for the long-term outcome

remains to be well cleared. Further clinical studies are needed to define these issues.

AUTHOR CONTRIBUTIONS

GL wrote the manuscript. CM and MM contributed to the conception of the manuscript and revised the paper critically. All authors contributed to the article and approved the submitted version.

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Simvastatin and ROCK Inhibitor Y-27632 Inhibit Myofibroblast Differentiation of Graves' Ophthalmopathy-Derived Orbital Fibroblasts *via* RhoA-Mediated ERK and p38 Signaling Pathways

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Transforming growth factor-β (TGF-β)-induced differentiation of orbital fibroblasts into myofibroblasts is an important pathogenesis of Graves' ophthalmopathy (GO) and leads to orbital tissue fibrosis. In the present study, we explored the antifibrotic effects of simvastatin and ROCK inhibitor Y-27632 in primary cultured GO orbital fibroblasts and tried to explain the molecular mechanisms behind these effects. Both simvastatin and Y-27632 inhibited TGF- β -induced α -smooth muscle actin (α -SMA) expression, which serves as a marker of fibrosis. The inhibitory effect of simvastatin on TGF- β -induced RhoA, ROCK1, and α-SMA expression could be reversed by geranylgeranyl pyrophosphate, an intermediate in the biosynthesis of cholesterol. This suggested that the mechanism of simvastatin-mediated antifibrotic effects may involve RhoA/ROCK signaling. Furthermore, simvastatin and Y-27632 suppressed TGF-β-induced phosphorylation of ERK and p38. The TGF- β -mediated α -SMA expression was suppressed by pharmacological inhibitors of p38 and ERK. These results suggested that simvastatin inhibits TGF-β-induced myofibroblast differentiation via suppression of the RhoA/ROCK/ERK and p38 MAPK signaling pathways. Thus, our study provides evidence that simvastatin and ROCK inhibitors may be potential therapeutic drugs for the prevention and treatment of orbital fibrosis in GO.

Keywords: simvastatin, Y-27632, Ras homolog family member A (RhoA), Rho-associated protein kinase (ROCK), myofibroblast, Graves' ophthalmopathy, ERK, p38

INTRODUCTION

Graves' ophthalmopathy (GO) is the ocular manifestation of Graves' disease (GD) and occurs in 25-50% of GD patients (1). It is characterized by inflammation and fibrosis of the orbital tissue, including the retrobulbar fat, connective tissue, and extraocular muscles (2). The expansion of the orbital tissue displaces the orbital globe by pushing it forward and results in clinical signs such as proptosis, lagophthalmos, and even sight-threatening exposure keratopathy or compressive optic neuropathy (1). Fibrosis of orbital tissue and enlargement of extraocular muscles may lead to strabismus and diplopia, which strongly affect the quality of life of patients suffering from GO (3). Orbital fibroblasts appear to play a central role in the pathogenesis of GO. The thyroid-stimulating hormone receptor and the insulin-like growth factor receptor on orbital fibroblasts are the major autoantigens responsible for GO (4). Once activated, orbital fibroblasts may generate numerous inflammatory molecules and differentiate into adipocytes or myofibroblasts, leading to tissue remolding in GO (5).

Orbital fibrosis is an important pathological change that occurs in GO (6). Transforming growth factor- β (TGF- β) plays a pivotal role in the pathophysiology of many fibrotic disorders, including GO (7). In previous studies, TGF- β has been used to induce extracellular matrix production and myofibroblast differentiation in GO orbital fibroblasts. The expression of fibrotic markers, including that of α -smooth muscle actin (α -SMA), connective tissue growth factor (CTGF), fibronectin, and collagen, was increased after the stimulation of TGF- β in GO orbital fibroblasts (7).

Statins are a class of drugs that are commonly used to lower serum cholesterol and prevent the development of cardiovascular diseases by inhibiting the enzyme hydroxymethylglutaryl-coenzyme A (HMG-CoA) reductase, which plays a crucial role in cholesterol synthesis. Statins also show pleiotropic effects e.g., anti-inflammatory, antifibrotic, and immune-modulatory (8, 9). In a recent database study among GD patients, the use of statins was associated with a reduced chance of developing GO (10). However, the precise molecular mechanism for it has not been fully established. Statins exert their antifibrotic effects on several organs, such as heart, lungs, and intestines (9, 11–13). In this study, we aimed to explore the antifibrotic effect of statins in GO to explain the possible molecular mechanisms through which statins reduce/inhibit GO development.

Ras homolog family member A (RhoA) plays an essential role in cell adhesion, migration, and transformation by organizing the actin cytoskeleton (14) and the Rho-associated protein kinase (ROCK) is

an important downstream effector of RhoA. There are two ROCK isoforms, ROCK1 and ROCK2, that are widely distributed in whole body tissues (15). The RhoA/ROCK signaling may play an important role in TGF-β signaling to control myofibroblast differentiation (16). Inhibition of the RhoA/ROCK pathway could reduce the TGF-β-induced expression of α-SMA, CTGF, and collagen I, as well as the myofibroblast differentiation of fibroblasts (17-19). Statins inhibit the synthesis of intermediates of cholesterol biosynthesis such as farnesyl pyrophosphate (FPP) and geranylgeranyl pyrophosphate (GGPP), which are essential for the post-translational modification of the Rho family proteins (8). Therefore, statins have a potential therapeutic role in fibrotic diseases owing to their inhibitory effects on RhoA/ROCK signaling (20, 21). Simvastatin, a type of statin, can suppress TGF-\(\beta\)-induced myofibroblast transformation of fibroblasts through RhoA/ROCK inhibition (20, 21).

In this study, we investigated the effects of simvastatin and ROCK inhibitor Y-27632 on TGF- $\beta1$ -induced $\alpha\text{-SMA}$ production in cultured GO orbital fibroblasts. We also explored the possible pathways through which simvastatin and ROCK inhibitor Y-27632 affect TGF- $\beta1$ signaling. We hypothesized that simvastatin and ROCK inhibitor Y-27632 could inhibit the myofibroblast differentiation of GO orbital fibroblasts, as well as the process of fibrosis in GO.

MATERIALS AND METHODS

Orbital Tissue Collection

Orbital tissue specimens were collected from the surgical waste produced during the orbital decompression surgeries of six GO patients. The clinical characteristics of the patients are listed in **Table 1**. All patients were euthyroid, with a clinical activity score of less than three for more than 6 months prior to the surgery. None of them had received radiotherapy or steroid pulse therapy in the past. Patient recruitment for the study was carried out at the Department of Ophthalmology, National Taiwan University Hospital, Taiwan. The protocol was approved by the Institutional Review Board of National Taiwan University Hospital, Taiwan (201908011RINA). Informed consent was obtained from all participants in accordance with the Declaration of Helsinki and Good Clinical Practice.

Orbital Fibroblast Culture and Stimulation

Orbital fibroblasts were cultivated as previously reported (22). Orbital tissues were minced and placed in plastic culture

 $\textbf{TABLE 1} \ | \ \text{Clinical characteristics of the patients included in this study}.$

| Age, years | Sex | Duration of GO, months | CAS | fT4 (ng/dL) | TSH (mU/L) | TBII (%) | Previous treatment for GO |
|------------|-----|------------------------|-----|-------------|------------|----------|---------------------------|
| 34 | F | 12 | 2 | 1.36 | 0.48 | 61.2 | None |
| 38 | M | 10 | 1 | 1.14 | 1.17 | 19.7 | None |
| 41 | F | 28 | 1 | 1.01 | 1.96 | 24.3 | None |
| 47 | F | 22 | 2 | 0.82 | 2.07 | 38.3 | Oral steroids |
| 57 | F | 20 | 1 | 1.09 | 1.39 | 54.2 | None |
| 58 | М | 36 | 2 | 0.91 | 0.57 | 44.5 | Oral steroids |

CAS, clinical activity score; fT4, free thyroxine; TSH, thyroid stimulating hormone; TBII, thyrotropin binding inhibitory immunoglobulin.

dishes with Dulbecco's modified Eagle's medium (DMEM) supplemented with 20% fetal bovine serum (FBS) and 50 IU/mL penicillin-streptomycin (Gibco[®]; Thermo Fisher Scientific, Waltham, MA, USA), allowing orbital fibroblasts to grow out. Monolayers were covered with DMEM supplemented with 10% FBS and serially passaged with gentle trypsin/ethylenediaminetetraacetic acid (EDTA) (Gibco[®]; Thermo Fisher Scientific, Waltham, MA, USA) treatment. These were incubated in a humidified incubator at 37°C and 5% carbon dioxide (CO₂).

The orbital fibroblasts were seeded into six-well plates at a density of 2×10^5 cells per well and stimulated at 90% confluence with 3 ng/mL of recombinant human TGF-β1 (Cat. No. 580701; BioLegend, San Diego, CA, USA) with or without pretreatment with simvastatin (TargetMol, Boston, MA, USA) or ROCK inhibitor Y-27632 (Sigma-Aldrich, St. Louis, MO, USA). In some experiments, fibroblasts were pretreated with PD98059 (ERK inhibitor) and SB203580 (p38 inhibitor) obtained from Cell Signaling Technology (Beverly, MA, USA) before stimulation with TGF-β1. We purchased FPP ammonium salt (F6892), GGPP ammonium salt (G6025), and mevalonate (90469) from Sigma (St. Louis, MO, USA) for cell stimulation in some experiments. GGTI-298 (Cat. No. 2430) and FTI-277 (Cat. No. 2407) were purchased from Tocris Bioscience (Bristol, United Kingdom). All experiments were performed with fibroblasts between the third and eighth passages from culture initiation. Six independent strains from different donors were used for the repeated experiments.

Immunofluorescence Staining

The orbital fibroblasts were cultured on six-well plates containing glass coverslips, and the cells were fixed and blocked for the immunofluorescence assay. The coverslips were incubated with anti-α-SMA primary antibody ab5694 (Abcam, Cambridge, MA, USA) overnight at 4°C. After three washes, the coverslips were processed with VectaFluor Excel Amplified Anti-Rabbit IgG Kit, DyLight[®] 488 Antibody Kit (DK-1488; Vector Laboratories, Burlingame, CA, USA). The nuclei were counterstained using H-1500 DAPI (Vector Laboratories, Burlingame, CA, USA). The glass slides were then observed with a fluorescence microscope coupled to a CCD camera (EVOS FLc; Life Technologies, Carlsbad, CA, USA). The fluorescence intensity quantification was performed using ImageJ.

Real-Time Polymerase Chain Reaction

Total RNA was extracted from cultured orbital fibroblasts using the TRIzol reagent (Invitrogen; Carlsbad, CA, USA). Complementary DNA (cDNA) was synthesized using 1 μ g of RNA according to the manufacturer's instructions for the iScript cDNA Synthesis Kit (Bio-Rad, Hercules, CA, USA). Real-time PCR was performed on a thermocycler (StepOne Real-Time PCR System; Applied Biosystems, Foster City, CA, USA) using SYBR Green Master Mix (Applied Biosystems, Foster City, CA, USA). All PCRs were performed in triplicate. The sequences of the primers used were as follows: α -SMA forward: 5'-CTC CCA GGG CTG TTT TCC CA-3', reverse: 5'-CCA TGT CGT CCC AGT TGG TG-3'; CTGF forward: 5'-TGT GTG ACG AGC CCA

AGG A-3', reverse: 5'-TCT GGG CCA AAC GTG TCT TC-3'; and Fibronectin 1 (FN1) forward: 5'-CCA AGA AGG GCT CGT GTG A-3', reverse: 5'-TGG CTG GAA CGG CAT CA-3'. The primers for glyceraldehyde 3-phosphate dehydrogenase (GAPDH) were obtained from Sino Biological, Inc. (Cat no. HP100003; Wayne, PA, USA). The messenger RNA (mRNA) levels of each target gene were normalized to GAPDH and represented as fold changes.

Western Blot Analysis

Proteins were extracted from tissue homogenates and cell lysates. After collecting the protein samples, equal amounts of protein (50 µg) were boiled in a sample buffer. The protein samples were separated in a 10% sodium dodecyl sulfate-polyacrylamide gel and transferred to a nitrocellulose membrane. The blots were blocked with 4% bovine serum albumin for 1 h at room temperature and then probed with primary antibodies at 4°C overnight. The ab5694 antibody for α -SMA (1:1000 dilution) was purchased from Abcam (Cambridge, MA, USA). The antibodies for RhoA (1:1000 dilution, no. 2117), ROCK1 (1:1000 dilution, no. 4035), Smad2/3 (1:1000 dilution, no. 5678), phospho-Smad2 (Ser465/467)/Smad3 (Ser423/425) (1:1000 dilution, no. 8828), ERK1/2 (1:1000 dilution, no. 9102), phospho-ERK1/2 (Thr202/Tyr204) (1:2000 dilution, no. 4370), p38 MAPK (1:1000 dilution, no. 8212), phospho-p38 MAPK (Thr180/Tyr182) (1:1000 dilution, no. 9211), JNK (1:1000 dilution, no. 9252), and phospho-JNK (Thr183/ Tyr185) (1:1000 dilution, no. 4668) were purchased from Cell Signaling Technology (Beverly, MA, USA). Immunoreactive bands were detected with horseradish peroxidase-conjugated secondary antibodies and developed using enhanced chemiluminescence detection (MilliporeSigma, Burlington, MA, USA) and exposure to X-ray film. The relative amount of each immunoreactive band was quantified using the ImageJ software and normalized to the levels of the reference molecules.

Statistical Analyses

At least three cell strains from different individuals were used in all experiments, and all sample assays were carried out in triplicate. The experimental results are shown as the mean \pm SD calculated from normalized measurements. Analysis of variance or the Student's t-test were used to determine statistical significance (p < 0.05) using GraphPad Prism 8.4.2 (GraphPad Software, San Diego, CA, USA).

RESULTS

Simvastatin and ROCK Inhibitor Y-27632 Inhibited TGF- β -Induced α -SMA Expression in GO Orbital Fibroblasts

To determine the effect of simvastatin and ROCK inhibitor Y-27632 on TGF- β -induced myofibroblast differentiation, immunofluorescence staining for α -SMA (marker for myofibroblast) was performed in cultured GO orbital fibroblasts. The expression of α -SMA and actin filament

formation were markedly increased after TGF- $\beta 1$ (3 ng/mL) stimulation for 48 h (**Figure 1**). However, pretreatment of the cells with different concentrations of simvastatin (1, 5, 10 μ M) (**Figure 1A**) or ROCK inhibitor Y-27632 (1, 10, 30 μ M) (**Figure 1B**) diminished TGF- β -induced α -SMA expression in orbital fibroblasts.

To further confirm the results, we used real-time PCR and western blot analysis to assess the mRNA and protein expression levels of α-SMA, respectively. As shown in Figure 2, TGF-β1 significantly induced an increase in mRNA and protein expression levels of α -SMA in GO orbital fibroblasts. Pretreatment with different concentrations of simvastatin (1, 5, 10 μ M) (Figures 2A, C) or ROCK inhibitor Y-27632 (1, 10, 30 μM) (Figures 2B, D) significantly inhibited TGF-β-induced α-SMA mRNA expression and protein production. The inhibitory effect of simvastatin and Y-27632 appeared to be dosedependent. Therefore, 10 and 30 µM concentrations of simvastatin and Y-27632, respectively, were used in the subsequent experiments to achieve a more prominent effect. As α-SMA is the most common marker for fibroblast-tomyofibroblast differentiation, these results suggested that simvastatin and ROCK-inhibitor Y-27632 have an inhibitory effect on the TGF-B-induced differentiation of orbital fibroblasts into myofibroblasts.

Inhibitory Effect of Simvastatin on TGF- β -Induced RhoA, ROCK1, and α -SMA Expression Was Reversed by GGPP

Simvastatin, an HMG-CoA reductase, inhibits the synthesis of mevalonate and other precursors of cholesterol such as FPP and GGPP, which are important for farnesvlation and geranylgeranylation of Rho proteins. To investigate the mechanism of simvastatin-mediated inhibition of myofibroblast differentiation, we studied the effects of the HMG-CoA downstream intermediates, mevalonate, FPP, and GGPP, on the expressions of RhoA, ROCK1, and α-SMA. As shown in Figures 3A-D, simvastatin inhibited TGF-β1-induced expression of RhoA, ROCK1, and α-SMA in GO orbital fibroblasts. The addition of 10 µM of GGPP reversed the inhibitory effects of simvastatin on RhoA, ROCK1, and α-SMA expression. However, the addition of FPP or mevalonate failed to prevent the suppressive effects of simvastatin. Moreover, the geranylgeranyl transferase inhibitor (GGTI-298), not the farnesyl transferase inhibitor (FTI-227), showed simvastatin-like inhibition of TGF- β 1-induced α -SMA (**Figure 3E**). Based on these results, we propose that the mechanism of simvastatin-mediated inhibition of myofibroblast differentiation may involve RhoA/ ROCK signaling. Simvastatin may inhibit the TGF-β1-induced RhoA/ROCK signaling by blocking Rho geranylgeranylation, but not Rho farnesylation.

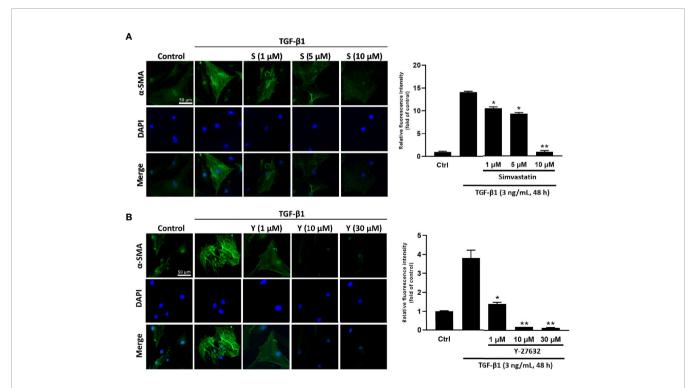


FIGURE 1 | Immunocytochemical characterization of transforming growth factor-β (TGF-β)-induced α-smooth muscle actin (α-SMA) expression in Graves' ophthalmopathy (GO) orbital fibroblasts. Primary cultured GO orbital fibroblasts were stimulated with 3 ng/mL TGF-β1 for 48 h with or without a 1-h pretreatment with different concentrations of simvastatin (1, 5, 10 μM) (A) or ROCK inhibitor Y-27632 (1, 10, 30 μM) (B). α-SMA expression was detected using immunofluorescence staining (green). The bar charts show mean data of relative fluorescence intensity of α-SMA presented as the fold of control. *p < 0.05, **p < 0.01 compared to cells treated with TGF-β1 alone. S = simvastatin: Y = Y-27632.

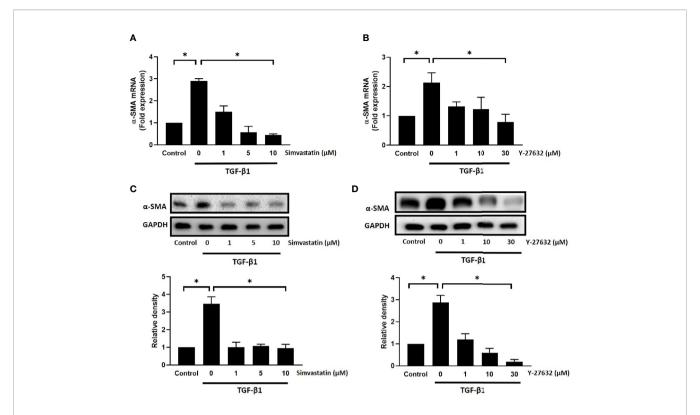


FIGURE 2 | Effect of simvastatin and ROCK inhibitor Y-27632 on transforming growth factor- β (TGF- β)-induced α -smooth muscle actin (α -SMA) messenger RNA (mRNA) and protein expression in Graves' ophthalmopathy (GO) orbital fibroblasts. Primary cultured GO orbital fibroblasts were stimulated with TGF- β 1 (3 ng/mL) for 48 h with or without a 1-h pretreatment with different concentrations of simvastatin (1, 5, 10 μM) (**A, C**) or ROCK inhibitor Y-27632 (1, 10, 30 μM) (**B, D**). The α -SMA mRNA expression was examined using real-time PCR (**A, B**). The α -SMA protein production was determined using western blot analysis (**C, D**). Data are presented as mean ± SD from at least three independent experiments. *p < 0.05.

Simvastatin and ROCK Inhibitor Y-27632 Inhibited TGF-β-Induced Phosphorylation of ERK and p38, But Not That of Smad

The signaling pathways implicated in TGF-β-induced myofibroblast differentiation include the Smad and non-Smad pathways, which incorporate the different branches of MAPK signaling (16). To explore the effects of simvastatin and ROCK inhibitor Y-27632 on TGF-β signaling in GO orbital fibroblasts, we investigated whether simvastatin and Y-27632 could inhibit TGF-βinduced phosphorylation of Smad2/3, ERK1/2, p38, and JNK. As shown in **Figures 4A–C**, 48 h of 3 ng/mL TGF-β treatment induced phosphorylation of Smad2/3, ERK1/2, and p38 in GO orbital fibroblasts. Simvastatin and Y-27632 successfully abrogated TGFβ-induced phosphorylation of ERK1/2 and p38. In contrast, they had no significant impact on TGF-β-induced phosphorylation of Smad2/3. Since the Smad and MAPK pathway could usually be activated in several minutes, we also assessed their phosphorylated states at early time-points (30, 60, 120 min) after TGF-β stimulation (Figure 4D). The results consistently demonstrated that simvastatin and Y-27632 inhibited TGF-β-induced phosphorylation of ERK1/2 and p38, but not Smad2/3. Our data suggested that simvastatin and Y-27632 could both inhibit the early- and late-phase activation of TGF-β-induced ERK and p38 signaling, but not Smad pathway in GO orbital fibroblasts.

Simvastatin and ROCK Inhibitor Y-27632 Inhibited TGF-β-Induced α-SMA Expression by Blocking ERK and p38 Signaling

To determine the potential involvement of ERK and p38 signaling in the mechanism by which simvastatin and Y-27632 inhibit myofibroblast differentiation, we investigated the effects of PD98059 (ERK inhibitor) and SB203580 (p38 inhibitor) on TGF-β-induced α-SMA expression. The results of western blot analysis showed that TGF-β-induced α-SMA expression in GO orbital fibroblasts was suppressed by PD98059 (**Figures 5A, C**) and SB203580 (**Figures 5B, D**). Based on previous data, simvastatin and Y-27632 have similar effects of ERK inhibitor and p38 inhibitor to block TGF-β-induced ERK and p38 signaling. These results suggested that simvastatin and Y-27632 inhibited TGF-β-induced α-SMA expression by blocking ERK and p38 signaling as well.

DISCUSSION

Orbital connective tissue remodeling and fibrosis are characteristic processes that appear relatively late in the clinical course of GO. The TGF- β produced by GO orbital fibroblasts may stimulate

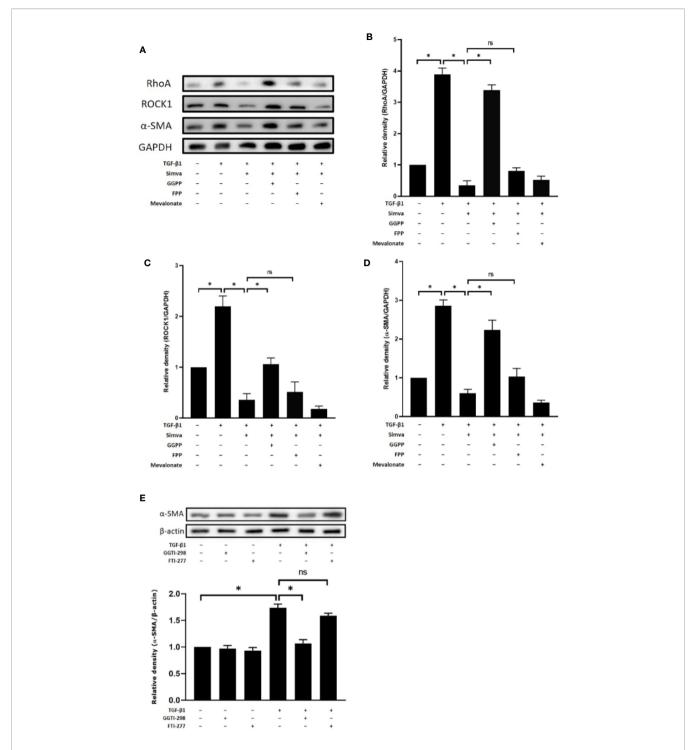


FIGURE 3 | Effects of hydroxymethylglutaryl-coenzyme A (HMG-CoA) downstream intermediates on simvastatin-mediated inhibition of transforming growth factor- β (TGF- β)-induced RhoA, Rho-associated protein kinase 1 (ROCK1), and α -smooth muscle actin (α -SMA) expression. (A) Graves' ophthalmopathy (GO) orbital fibroblasts were stimulated with TGF- β 1 (3 ng/mL) for 48 h with or without a 1-h pretreatment with simvastatin (10 μM) and addition of geranylgeranyl pyrophosphate (GGPP) (10 μM), farnesyl pyrophosphate (FPP) (10 μM), or mevalonate (200 μM). The protein levels of RhoA, ROCK1, and α -SMA were determined using western blot analysis. (B–D) The densities of RhoA (B), ROCK1 (C), and α -SMA (D) protein bands were quantified and normalized to GAPDH. (E) GO orbital fibroblasts were stimulated with TGF- β 1 (3 ng/mL) for 48 h with or without a 1-h pretreatment with geranylgeranyl transferase inhibitor (GGTI-298) (10 μM) or farnesyl transferase inhibitor (FTI-227) (10 μM). The α -SMA protein production was determined using western blot analysis. Data are presented as mean ± SD of at least three independent experiments. *p < 0.05.

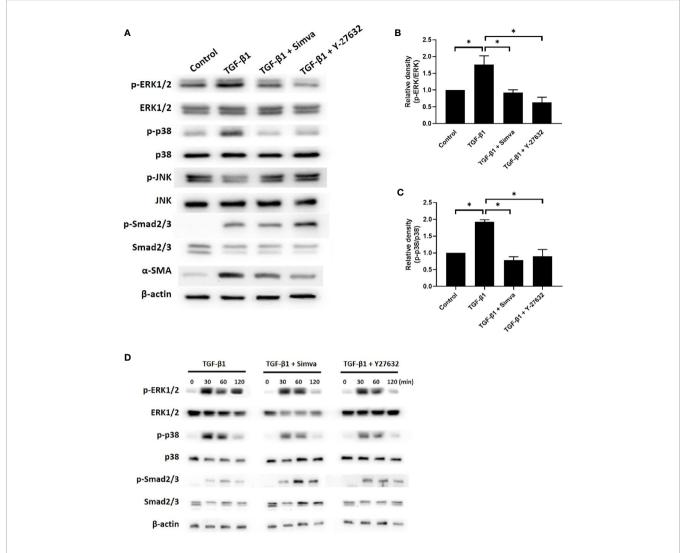


FIGURE 4 | Effects of simvastatin and ROCK inhibitor Y-27632 on transforming growth factor-β (TGF-β)-induced signaling. (A) Primary cultured Graves' ophthalmopathy (GO) orbital fibroblasts were stimulated with TGF-β1 (3 ng/mL) for 48 h with or without a 1-h pretreatment with simvastatin (10 μM) or ROCK inhibitor Y-27632 (30 μM). The expression and phosphorylation levels of Smad2/3, ERK1/2, p38, and JNK were determined using western blot analysis. (B, C) The densities of ERK1/2 (B) and p38 (C) protein bands were quantified and normalized to β-actin. (D) GO orbital fibroblasts were stimulated with TGF-β1 (3 ng/mL) for 30, 60, 120 min with or without a 1-h pretreatment with simvastatin (10 μM) or ROCK inhibitor Y-27632 (30 μM). The expression and phosphorylation levels of Smad2/3, ERK1/2, and p38 were determined using western blot analysis. Data are presented as mean ± SD of at least three independent experiments. *p < 0.05.

excessive production of extracellular matrix and differentiation of Thy-1-positive orbital fibroblasts into myofibroblasts, which express strong $\alpha\text{-SMA}$ immunoreactivity (22, 23). In this study, we showed that simvastatin and ROCK inhibitor Y-27632 inhibited the expression of TGF- β -induced $\alpha\text{-SMA}$, which serves as a marker for fibrosis and indicates myofibroblast differentiation.

Some studies have demonstrated the inhibitory effects of simvastatin and Y-27632 on TGF- β -induced myofibroblast differentiation from fibroblasts derived from different disease specimens, such as nasal polyps (24), keloids (20), and penile tunica albuginea from Peyronie's disease (21). To the best of our knowledge, this is the first study to investigate the antifibrotic

effects of simvastatin and Y-27632 in GO-derived fibroblasts. Further investigations on any synergistic effects of these two components will be helpful to understand their antifibrotic properties and potential therapeutic roles in GO.

Statins are commonly used to prevent coronary artery disease and stroke by reducing low-density lipoprotein cholesterol levels. Recently, the pleiotropic antifibrotic effects of statins have been proposed in various organ systems, such as heart (9, 11), liver (25, 26), and lungs (27, 28). Also, the possible protective role of statins in GO has been proposed (29). Statin usage is believed to be associated with a reduced risk of developing GO among patients with GD in a large cohort study (10). They found that

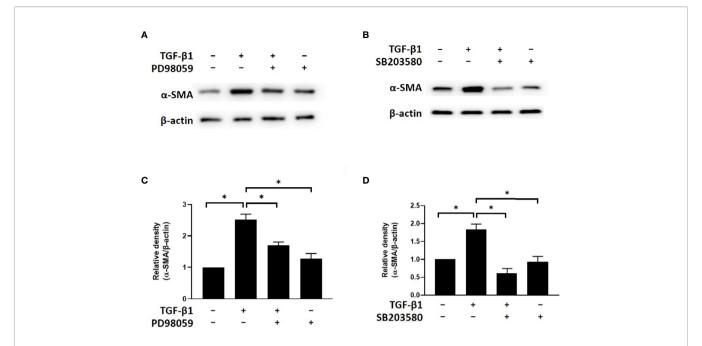


FIGURE 5 | Effects of ERK and p38 inhibitors on TGF- β -induced α-SMA expression in Graves's ophthalmopathy (GO) orbital fibroblasts. Primary cultured GO orbital fibroblasts were stimulated with transforming growth factor- β (TGF- β 1) (3 ng/mL) for 48 h with or without a 1-h pretreatment with 10 μM of PD98059 (ERK inhibitor) (**A, C**) and 10 μM of SB203580 (p38 inhibitor) (**B, D**). The protein levels of α-SMA were determined using western blot analysis. The densities of α-SMA protein bands were quantified and normalized to β -actin. Data are presented as mean ± SD of at least three independent experiments. *p < 0.05.

patients who used statins for at least 60 d during the period of observation had a 40% lower risk of developing GO. The precise molecular mechanisms by which statins reduce GO risk are not fully established. Some evidence suggests that statins may modulate both apoptosis and autophagic activities in patients with GD (29). This elucidation was based on the involvement of cellular apoptosis and autophagy in the pathogenesis of GO (30, 31). In addition, a recent study revealed that simvastatin may inhibit adipogenesis, as well as the expression of early and late adipogenic genes in human orbital fibroblasts (32). Our results provided *in vitro* evidence of simvastatin-mediated antifibrotic effects in GO that may also explain the possible protective effect of simvastatin against the development of GO.

The possible mechanism underlying statin-mediated antifibrotic effects is the inhibition of geranylgeranylated Rho protein, which in turn inhibits the Rho/ROCK signaling pathway (33, 34). Statins inhibit HMG-CoA reductase, the catalyst for the synthesis of mevalonate from HMG-CoA. This inhibition leads to a reduction of downstream intermediate compounds, including the isoprenoid GGPP and FPP. These molecules are necessary for the posttranslational modification of the Rho proteins, which is crucial for the Rho proteins to play their proper functions. In human keloid fibroblasts, simvastatin inhibited TGF-β-induced RhoA activation and RhoA/ROCK signaling by interfering with posttranslational geranylgeranylation of RhoA (20). In human airway fibroblasts, the inhibitory effects of simvastatin on TGF-βinduced fibronectin could be reversed by the addition of either GGPP or FPP (28). The study of human tenon fibroblasts suggested that the inhibition of Rho-geranylgeranylation, not Rhofarnesylation, was the mechanism for lovastatin to inhibit myofibroblast differentiation (35). In our study, we found that only GGPP, and not FPP, could reverse the simvastatin-mediated inhibition of TGF- β -induced α -SMA.

Consistently, only the geranylgeranyl transferase inhibitor (GGTI-298), not the farnesyl transferase inhibitor (FTI-227), showed simvastatin-like inhibition of TGF- β 1-induced α -SMA. These findings suggested that geranylgeranylation rather than farnesylation of RhoA is crucial for TGF- β -induced α -SMA expression in GO orbital fibroblasts. And simvastatin may inhibit the TGF- β 1-induced RhoA/ROCK signaling by blocking Rho geranylgeranylation, but not Rho farnesylation.

The RhoA/ROCK signaling pathway is known to regulate numerous cellular functions, including cell proliferation, migration, contraction, and adhesion (36). The profound involvement of the RhoA/ROCK pathway in various disease processes has made ROCK a potential therapeutic target in many kinds of diseases, e.g., cardiovascular (37), neoplastic (38), and neurologic (39). Moreover, ROCK inhibitors have been used as potential therapeutic drugs for several ophthalmic diseases, including glaucoma, corneal endothelial diseases, age-related macular degeneration, and diabetic retinopathy (40, 41). However, there are few studies that have investigated the role of Rho/ROCK signaling or that of ROCK inhibitors in GO. Using an in vitro model of GO, our study demonstrated the involvement of RhoA/ROCK signaling in TGF-\beta-induced myofibroblast differentiation and the antifibrotic effects of the ROCK inhibitor Y-27632. In many studies Y-27632 is a commonly used ROCK inhibitor that inhibits both ROCK1 and ROCK2 (42). Further investigations are necessary to

explore the possible applications of ROCK inhibitors in the treatment of GO.

Transforming growth factor-β is the most potent inducer of myofibroblast differentiation and acts by activating the canonical Smad pathway or the non-Smad pathways, including Rho/ROCK signaling and different branches of the MAPK pathway (16, 43). The ROCK inhibitors were reported to inhibit TGF-β-induced myofibroblast differentiation by regulating the Smad or MAPK signaling pathways (19, 44). For example, Y-27632 suppressed TGF-\(\beta\)-induced phosphorylation of Smad3, but not that of Smad2, in ocular Tenon's capsule fibroblasts (44). It also inhibited TGF-β-induced phosphorylation of ERK and JNK, but not that of p38, in renal mesangial cells (19). In our study, Y-27632 abrogated TGF-B-induced phosphorylation of ERK and p38, but not that of JNK or Smad2/3. These results indicate that ROCK mediates MAPK signaling, but each MAPK signaling is regulated distinctively in different cells or by different stimuli. In our study, simvastatin showed an effect similar to that of Y-27632 in TGF-β-induced myofibroblast differentiation. These results suggest that, although simvastatin does not regulate the TGF-β-mediated Smad pathway in GO orbital fibroblasts, it regulates the TGF-β-mediated ERK/p38 MAPK

pathways, probably in a ROCK-dependent manner. In summary, we propose that simvastatin can inhibit TGF-β-induced myofibroblast differentiation *via* suppression of the RhoA/ROCK/ERK and p38 MAPK signaling pathways (**Figure 6**).

Contrary to our findings, previous studies showed that simvastatin could inhibit TGF- β -mediated Smad phosphorylation in human ventricular (11) and intestinal fibroblasts (45). We believe that the cellular mechanisms of simvastatin's antifibrotic effect are diverse and complex in different cell types. Our data could not elucidate whether RhoA/ROCK/ERK and p38 MAPK signaling has any interaction with Smad2/3 signaling. Recently, Fang *et al.* reported that interleukin-17A can activate the JNK signaling in CD90⁺ orbital fibroblasts to promote GO fibrosis initiated by TGF- β -mediated Smad transcription (46). Further investigation is needed to understand the possible interactions between the TGF- β -induced RhoA, MAPK, and Smad pathways in GO orbital fibroblasts.

Our study also had some limitations. First, the GO orbital fibroblasts were obtained from patients with inactive GO because of the surgical indication. Further studies must be conducted using specimens from patients with active GO to compare the results between different disease stages. Second, we only managed to obtain orbital tissues from a small cohort of

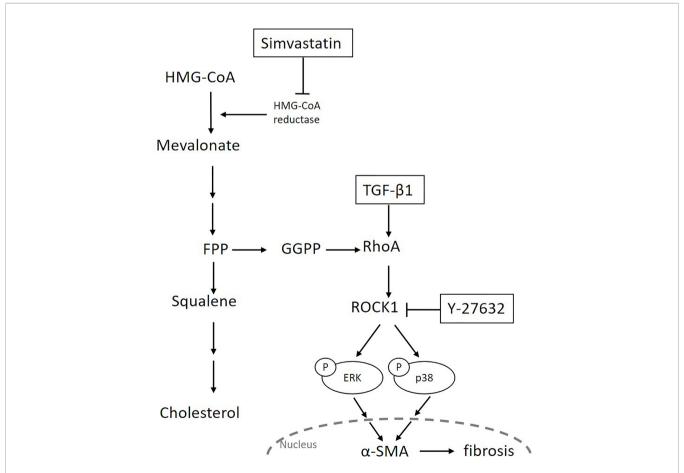


FIGURE 6 | Schematic diagram of the possible signaling mechanisms underlying simvastatin- and Y-27632-mediated inhibition of transforming growth factor-β (TGF-β1)-induced orbital tissue fibrosis in Graves' ophthalmopathy.

patients. We may need a larger sample size to validate our results in the future

In conclusion, our results provide preliminary evidence that simvastatin and ROCK inhibitor Y-27632 inhibit TGF- β -induced differentiation of GO-derived fibroblasts into myofibroblasts. We propose that simvastatin and ROCK inhibitors may be potential candidates for the prevention and treatment of orbital tissue fibrosis in GO.

DATA AVAILABILITY STATEMENT

The raw data supporting the conclusions of this article will be made available by the authors, without undue reservation.

ETHICS STATEMENT

The studies involving human participants were reviewed and approved by the Institutional Review Board of National Taiwan

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University Hospital. The patients/participants provided their written informed consent to participate in this study. No animal studies are presented in this manuscript. No potentially identifiable human images or data is presented in this study.

AUTHOR CONTRIBUTIONS

Y-HW, S-LL and C-CW conceptualized the experiments, designed the study, and secured funding. S-LL provided the surgical specimens. S-HW and C-CW performed the experiments and helped analyze the data. Y-HW interpreted the data and wrote the manuscript. C-HY supervised the research activity and revised the manuscript. All authors contributed to the article and approved the submitted version.

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Stimulatory Thyrotropin Receptor Antibodies Are a Biomarker for Graves' Orbitopathy

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INTRODUCTION

Graves' Orbitopathy (GO) or thyroid eye disease (TED) is the most common and prominent extrathyroidal manifestation in patients diagnosed with Graves' disease (GD). Approximately 50% of patients diagnosed with GD develop GO, and with further orbital imaging, up to 80% of patients test positive for TED (1). Physical signs and symptoms (2), as well as the impact on the psychological well-being (3, 4) and quality of life (5, 6), underline the importance of effective disease management. Furthermore, the incidence of TED is linked to autoimmune gastritis and coeliac disease (7), implicating the importance of diagnosis of GO in patients, to identify risk factors for gastrointestinal autoimmunity.

Current evidence indicates that stimulating thyrotropin receptor IgG antibodies (stim. TSH-R-Ab or TSAb) are the main causative agent of GO, making its measurement a useful tool to predict and assess both clinical disease severity and activity.

TERMINOLOGY

Numerous terms have been established to describe TSH-R-Ab, the different nomenclature refers to various types of immunoassays (8). Total TSH-R-Ab also referred to as TRAb, are Ab that interact distinctively with the TSH-R. Routinely, these Ab are gauged by competitive immune binding assays and therefore are termed TSH-R-binding inhibitory immunoglobulins (TBII). Since binding assays only assess the binding of Ab to the TSH-R, it cannot display the function of the Ab it tests. Cell-based bioassays however differentiate between Ab that block or stimulate the TSH-R. Stimulating Ab are termed thyroid-stimulating Ab (TSAb) or thyroid-stimulating immunoglobulins (TSI), while blocking Ab are referred to as thyroid blocking Ab (TBAb) or thyroid blocking immunoglobulins (TBI). Further alternative terms for TBAb are TSH-R-stimulating-blocking Ab and TSH-R-blocking Ab or TRBAb.

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DISTINCTION BETWEEN BLOCKING AND STIMULATING ANTIBODIES

TSH-R-Ab are found in patients with autoimmune thyroid disease (AITD) and play a major role in their pathogenesis and clinical presentation (9). TSH-R-Ab are divided into three groups since they have different ways to interact with the TSH-R. TSAb bind to the large extracellular amino-terminal and cause a stimulation of the TSH-R. This results in increased cyclic adenosine 3', 5'monophosphate (cAMP), which increases the transcription of proteins necessary for the synthesis of thyroxine (T4), and triiodothyronine (T3). The outcome of this pathway is higher synthesis rates of thyroid-related hormones T3 and T4, and increased proliferation of the thyroid follicular endothelial cell. On the other hand, TBAb inhibit the function of the TSH-R, decreasing the synthesis rate of thyroid hormones, and inhibiting the proliferation and growth of the thyroid cell (10, 11). Other TSH-R-Ab neither block nor stimulate but rather have a neutral effect on the TSH-R. These Ab are referred to as neutral Ab or "cleavage" Ab and are not fully understood in their respective effect in vivo. Neutral Ab are able to induce various signaling cascades including some of which are initiated by TSAb, as well. However, neutral Ab can induce unique downstream signaling cascades including the activation of protein kinase C/mitogen activated protein kinase (MAPK), mammalian target of rapamycin (mTOR), nuclear factor 'kappa light chain enhancer' of activated B-cells (NF-κB), different cytokines, and reactive oxygen species (ROS). The initiation of these pathways is yet unclear, a recruitment of multiple G-proteins is possible. In vitro experiments show the expression of heat shock Proteins (p27, p107), endoplasmic reticulum stress protein (grp98), various oncogenes (p53, p73, retinoblastoma protein), and apoptosis of rat thyrocytes after a period of exposure to neutral Ab. It is unclear if ROS alone, or other signaling cascades play a role in the initiation of apoptosis (12, 13).

ROLE OF FUNCTIONAL TSH-R AUTOANTIBODIES IN GO

GO/TED is characterized by a protrusion of the eyes, upper lid retraction, diplopia, and irritation of the periorbital tissue and conjunctiva. In a study including 101 consecutive patients with TED, none tested positive for TBAb while 91 (90%) showed presence of TSAb of whom 90 were diagnosed with GD (14), concluding that not TBAb but rather TSAb are greatly prevalent in patients with TED. Current evidence suggests that TED is primarily caused by TSAb, which are present in patients with GD and Hashimoto's thyroiditis (HT). Animal models support the hypothesis of TSAb and TBAb, being synthesized as a result of immunization with the TSH-R, detected as an auto-antigen (15). The TSH-R is physiologically expressed in the soft tissue of the orbit, which was demonstrated in experiments where TSH-R mRNA was detected in affected tissue of the eye socket (16). Another study independently noted an overexpression of the TSH-R and human leukocyte antigen-DR (HLA-DR) in patients with GO (17). Produced by B- cells, TSAb enter the bloodstream

and stimulate the TSH-R in various tissues e.g., the thyroid gland, orbital soft tissue, skin, and heart. In orbital fibroblasts, the Ab induce differentiation into pre-adipocytes and secretion of hydrophilic glycosaminoglycans (GAG), resulting in edema and later fibrosis (18, 19) (**Figure 1**).

Inflammatory cytokines play a major role in the pathogenesis of GO. By binding to the TSH-R, TSAb induces inflammation through inflammatory cytokines such as tumor necrosis factor- α (TNF- α), interferon- γ , and interleukin-1 which perpetuate the synthesis of collagen, GAG, and differentiation of orbital fibroblasts into adipocytes (20). Furthermore, by inducing the expression of heat shock proteins, prostaglandins, adhesion molecules, CD40, and major histocompatibility complex protein class two, cytokines amplify the inflammation cascade (21). The importance of inflammation explains the success of anti-inflammatory drugs like mycophenolate and glucocorticoids in the treatment of GO (22–24). The pathogenesis of GO is further supported by evidence, as high TSAb titers are associated with GO in patients with GD (25).

CLINICAL RELEVANCE OF TSAB AS A BIOMARKER FOR GO

Since the clinical presentation of GD and GO, especially the eye protrusion, is caused by stimulating TSH-R-Ab, measurement of TSAb is an excellent tool to manage patients with GD/GO. As there is a need to assess the function of the Ab, a cell-based bioassay rather than a binding assay is preferred (26, 27). In cell-based bioassays, IgG Ab are gauged through the signal transduction of cAMP, mediated through genetically engineered TSH-R on intact cells. Previously, the concentration of cAMP was measured through radiolabeled immunoassays (RIA). More recently, the amount of cAMP is assessed with cAMP-inducible reporter genes that express the luciferase enzyme. The enzyme concentration is then gauged with a luminometer after the luciferase substrate is added. TBAb are measured in an analogous bioassay while measuring their ability to competitively block the TSH-R against bovine TSH. In contrast, clinically employed immune binding assays assess the amount of TRAb, by measuring the displacement of a tracer, which is either radioiodine labeled bovine TSH or a monoclonal mouse antibody (MAb) with affinity to the TSH-R.

Previously, cell-based bioassays were toilsome and complicated procedures with varying specificity and sensitivity between laboratories, making it an unreliable way to measure Ab in comparison to binding assays. However, through major improvements in genetic engineering and molecular cloning, cell-based bioassays have become much more reliable and easier to perform. To improve reproducibility, the use of transfected cell lines of the Chinese ovarian hamster (CHO) was established, making the procedure more convenient with an improvement of assay sensitivity and specificity. Even though binding TSH-R-Ab or TBII can be used for differential diagnosis of GO (28), cell-based bioassays significantly outperform binding assays in terms of sensitivity and accuracy (27, 29–31), making it a better choice for management of GO.

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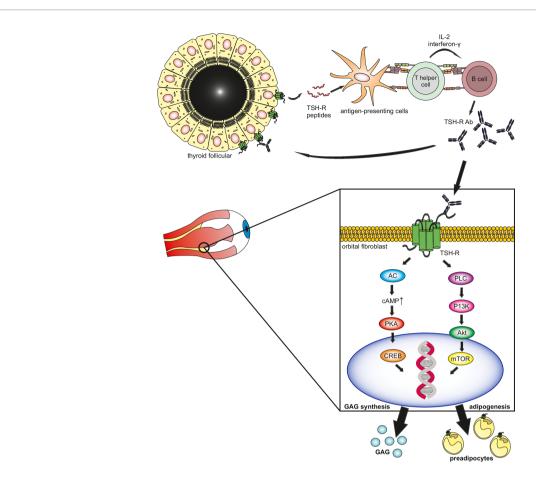


FIGURE 1 | The thyrotropin receptor (TSH-R) is the main autoantigen in Graves' hyperthyroidism and associated eye disease. TSH-R peptides are ingested by antigen-presenting cells (APC) through phagocytosis and expressed through MHC class II. T-helper cells recognize the antigen, by binding with the T-cell receptor, along with CD154 to its ligand CD40 on the surface of the APC. The activated T-helper cells bind to B-cells, transforming them into TSH-R antibody-secreting plasma cells through inflammatory cytokines interleukin II and gamma interferon. The synthesized TSAb bind to the TSH-R expressed by thyrocytes and orbital target cells (fibroblasts, pre-adipocytes), activating the Gαs adenylyl cyclase (AC) pathway. This stimulates protein kinase A which induces gene activation through the cAMP responsive element binding (CREB) protein. Additionally, the Gαq protein kinase C (PKC) pathway activates protein kinase B (Akt) inducing the mammalian target of rapamycin (mTOR) further inducing gene expression. The induction of gene expression lead to differentiation into pre-adipocytes and synthesis of hydrophilic mucopolysaccharides (glycosaminoglycans), hence leading to edema and later fibrosis in the orbital space, resulting in the clinical phenotype of thyroid eye disease.

Several studies were conducted to assess the reliability of TSAb as predictor for GO. A report investigating the relation of TSH-R-Ab and GO underlines the benefits of using functional Ab in the clinical routine (32). A total of 155 GD patients were tested for TSAb, using two different types of cell lines. All hyperthyroid patients with GD and TED tested positive for TSAb/TSI and TSAb were detected in 150 of 155 patients with TED. None of the 40 controls was Ab positive. TSAb levels in patients with GO were 3-fold and 8-fold higher than in GD patients and healthy controls, respectively.

To evaluate the relation between TSAb and the clinical presentation of TED, two standardized classifications were utilized. A clinical activity score (CAS), determined through an author unaware of the measured lab data, was used. To gauge the severity of TED, a clinical severity score (CSS) was utilized, based on the NOSPECS classification (33). TSAb titer correlated with both severity (r = 0.87, p < 0.001) and activity (r = 0.87, p < 0.001)

0.001). In contrast, radiolabeled assays (RIA) showed weaker correlations at $r=0.17,\ p<0.015$ and $r=0.54,\ p<0.001,$ respectively. Further, a cross-sectional trial assessing the clinical relevance of TSI in regards to TED, showed a similar correlation between severity (r = 0.83, p<0.001) and activity (r = 0.81, p<0.001) (34). A more significant association of TSAb with clinical features of GO was observed than TBII and thus TSAb may be regarded as functional biomarker for GO.

A multicenter cross-sectional study evaluated the relation of TSAb and TED in children with GD (35). 422 samples from 157 children with GD, 101 samples from non-thyroidal autoimmune diseases, and 50 healthy controls were tested. All patients with GD+GO tested positive for TSAb, compared to 96% in the binding assay (both p < 0.001). Further, in euthyroid children with GO, TBII were positive in 24 of 31 (77%) children only, while TSI were detected in all subjects (both p = 0.016). Additionally, in a different trial (36), children with GD+GO

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displayed the highest titers of TSAb (SRR% 417±135) in comparison to GD only (SRR% 320±157). Additionally, hyperthyroid children with GD+GO displayed higher TSI levels compared to children with GD only (median SRR%, 481 vs. 395%, both p < 0.001). TBII however, failed to differ between the two groups (p < 0.125). With the classification of disease severity, children were distinguished in moderate-to-severe and mild GO according to the classification of the European Group on Graves' Orbitopathy (EUGOGO). As in adults, children with moderate-to-severe GO exhibited higher TSAb titers compared to cases with mild GO (median SRR% 536 vs. 259, p < 0.001). Following the subjects during an average 3-year antithyroid drug (ATD) treatment, TSAb decreased by 69% and 20% in patients with GD and GD+GO, respectively. In contrast, TBII titers did not significantly differ between the two groups (90 vs. 89% in GD vs. GD+GO). Hence, TSAb is a strong indicator and predictor of GO in children (37), and supports the use of TSI/TSAb for management of pediatric GD/GO.

Furthermore, a 2-year prospective trial was conducted to evaluate the success of ATD treatment of one hundred consecutive hyperthyroid adults with GD, by measuring functional Ab and TBII (38). Forty-four of one hundred patients responded to ATD with Methimazole (MMI), of whom 43% suffered from GO. In the group with 56 non-responders, 66% of patients were diagnosed with GO. TSAb mirrored the activity of GD and was able to differentiate responders from non-responders. Additionally, and in contrast to TBII, TSAb titers were higher in adults with GO+GD, compared to GD alone. Finally, studies from Japan and Singapore also support the fact that TSAb correlate with the severity of GO. Indeed, increased TSAb titers are found in adults with higher GO scores (39), and in patients with GD+GO vs. GD only (40). As shown previously, GO activity did not correlate with serum TBII levels.

CLINICAL UTILITY OF FUNCTIONAL TSH-R-AB

Cell-based bioassays exclusively and solely differentiate between blocking and stimulating TSH-R-Ab and can measure TSH-R-Ab at very low concentrations (27). The presence of TSI/TSAb is predictive for GO (39). In contrast, TBII is not associated with GO. The utilization of TSAb facilitates rapid diagnosis of GD and allows differential diagnosis of thyrotoxicosis, being the only biomarker able to reliably differentiate GO+GD from patients with GD only. In the clinical routine, using early *in vitro* testing of TSAb showed a 46% faster time to diagnosis of GD and 47% cost savings since costly procedures and consultation of

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specialists was reduced (41). During specific treatment, TSAb detects patients with GO and can predict their responsiveness to therapy. Therefore, TSAb is the superior tool to manage GO in pediatric and adult patients (9, 20, 37, 42). Recently, an immunoassay has been developed to measure TSH-R-Ab using the bridge technology (43). It has been assumed that this assay detects TSAb only. The assay utilizes a pair of recombinant TSH-R and TSH-R-Ab are measured by binding one antibody arm to a capture receptor on the solid phase and bridging with the other arm to a detection receptor generating a signal. However, numerous comparative studies have proved that the bridge immunoassay is not able to differentiate between functional TSH-R-Ab (stimulating or blocking). Indeed, twenty hypothyroid Hashimoto's thyroiditis (HT) patients with high titers of TBAb measured in a blocking TSH-R-Ab cell-based bioassay, tested all positive in the bridge immunoassay (30). In another study, ten samples from TBAb-positive/TSAb-negative (both measured in bioassays) patients with GD or HT were positive in the bridge immunoassay (31). Further, various mixtures of monoclonal antibodies (MAb) of M22 and K1-70 were positive detected in the bridge assay. A recent study revealed that TBAb were present in one patient with HT and in two patients with GD and these patients were also positive in the bridge immunoassay but negative in the TSAb bioassay (29). Furthermore, all TBAb positive samples in a Graves' disease animal model were positive in the bridge assay without exception (15). Therefore, the bridge assay is a purely binding immunoassay. In conclusion, although not standardized vet and requiring more time and experienced lab personal, functional TSH-R-Ab in general and stimulatory Ab in particular are clinically useful and have been demonstrated to be a reliable and accurate biomarker for the diagnosis, differential diagnosis and monitoring of patients with GO.

AUTHOR CONTRIBUTIONS

Literature search, writing, text revision, layout: AG. Primary concept, writing, editing, critical evaluation, supervision: GK. Critical evaluation and editorial assistance: TD and JL. All authors contributed to the article and approved the submitted version.

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George et al. TSH-R-Ab and GO

Conflict of Interest: GK consults for Immunovant, Mediomics, Merck, Novartis, and Quidel.

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Microarray Data of Lacrimal Gland Implicates Dysregulated Protein Processing in Endoplasmic Reticulum in Graves' Ophthalmopathy

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Graves' ophthalmopathy (GO) has become one of the most common orbital diseases. Although some evidences announced the potential mechanism of pathological changes in extraocular muscle and orbital adipose tissue, little is known about that in lacrimal enlargement of GO patients. Thus, gene expression profiles of lacrimal gland derived from GO patients and normal controls were investigated using the microarray datasets of GSE105149 and GSE58331. The raw data and annotation files of GSE105149 and GSE58331 were downloaded from Gene Expression Omnibus (GEO) database. Bioinformatics including differentially expressed genes (DEGs), Gene Ontology, Kyoto Encyclopedia of Gene and Genome (KEGG) pathway, protein-protein interaction (PPI) network construction, hub gene identification, and gene set variation analysis (GSVA) were successively performed. A total of 173 overlapping DEGs in GSE105149 and GSE58331 were screened out, including 20 up-regulated and 153 down-regulated genes. Gene Ontology, KEGG and GSVA analyses of these DEGs showed that the most significant mechanism was closely associated with endoplasmic reticulum (ER). Moreover, we identified 40 module genes and 13 hub genes which were also enriched in the ER-associated terms and pathways. Among the hub genes, five genes including HSP90AA1, HSP90B1, DNAJC10, HSPA5, and CANX may be involved in the dysfunction of protein processing in ER. Taken together, our observations revealed a dysregulated gene network which is essential for protein processing in ER in GO patients. These findings provided a potential mechanism in the progression of lacrimal enlargement in GO patients, as a new insight into GO pathogenesis.

Keywords: Graves' ophthalmopathy, lacrimal gland, protein processing in endoplasmic reticulum, hub gene, microarray data

INTRODUCTION

Graves' ophthalmopathy (GO), also known as thyroid eye disease (TED), is an orbital disease that is uniquely linked to Graves' disease (GD), generally present in 25%-50% of GD patients (1). The main clinical features of GO include proptosis, lacrimal gland enlargement, eyelid retraction, diplopia, and exposure keratopathy, even developing into irreversible vision loss in severe cases (2). It is generally agreed that GO is an autoimmune disease that results in orbital remodeling, enlargement and fibrosis (3). These pathological changes involves three distinct but related immune processes: inflammation, adipogenesis and glycosaminoglycan accumulation (4). Most investigators have hypothesized that orbital fibroblasts are the main targets of inflammatory cytokines released by infiltrated immune cells in the extraocular muscle and orbital adipose tissue (5). Moreover, activated orbital fibroblasts could secrete many cytokines, such as interleukins 1β, 6, 8, 16, TNF-α, RANTES and CD154 (6). These cytokines promote orbital trafficking of monocytes and macrophages, facilitate differentiation of orbital fibroblasts and stimulate accumulation of hyaluronic acid-rich stroma, leading to uncontrolled immune responses (4-6). Although several mechanisms have been proposed for the development of GO, the exact pathogenesis of GO has not yet been illustrated.

Recently, high-throughput technologies provided ample evidence on the potential biomarkers and molecular mechanisms of GO. For examples, microarray analysis revealed the potential role of CASO2 to trigger autoimmunity events (7), and identified several adipogenesis-related genes and some genes involved in Wnt and IGF-1 signaling as being potentially implicated in pathogenesis (8-11). RNA-Seq also identified some meaningful genes (such as PTX3, HOXB2, HOXB3, CCL2, and SERPINA1) as potential biomarkers of GO (12-14). In addition, microRNA and protein sequencing found some blood circulating biomarkers that have the potential to diagnose GD, predict GO disease status and optimize patient management, such as hsa-miR-27a-3p, hsa-miR-22-3p, zonulin, haptoglobin, and lumican (15). All these accumulating findings improve our knowledge of the mechanisms underlying pathological changes in extraocular muscle and orbital adipose tissue of GO. Nevertheless, our understanding of the molecular mechanisms of lacrimal gland enlargement in GO is still limited.

In the present study, microarray analysis was performed to systematically investigate the key genes and pathways in lacrimal gland by mining the microarray datasets (GSE105149 and GSE58331) of GO and normal lacrimal samples. GSE105149 and GSE58331 were obtained from the public database GEO. These results would help to uncover a potential mechanism of lacrimal enlargement and offer new information for the pathogenesis of GO.

MATERIALS AND METHODS

Characteristics of the Microarray Data Sets

The Gene Expression Omnibus (GEO, http://www.ncbi.nlm.nih. gov/geo/) is a public functional genomics data repository. GEO

has collected a large amount of omics data, such as highthroughput gene expression data, chips and microarrays. Microarray datasets of GSE105149 and GSE58331 were obtained by searching the keywords including Graves' ophthalmopathy, thyroid eye disease and thyroid-associated ophthalmopathy in GEO. The raw data and annotation files of GSE105149 and GSE58331 were downloaded from the GEO database. Rosenbaum JT et al. contributed to the generation of GSE105149 and GSE58331and provided many significant findings from different perspectives (16-21). GSE105149 was the microarray data of various lacrimal gland tissues, including sarcoidosis, granulomatosis with polyangiitis, thyroid eye disease, nonspecific orbital inflammation and healthy controls (16). Normal lacrimal samples from healthy individuals were obtained at the time of cosmetic surgery or blepharoplasty (16). GSE58331 was the microarray data of anterior orbit and lacrimal gland tissues, which were obtained from subjects with inflammatory diseases such as nonspecific orbital inflammation, sarcoidosis, granulomatosis with polyangiitis, IgG4-associated disease and thyroid eye disease as wells as from normal controls (17-21). Normal control tissues were obtained during surgeries such as blepharoplasty and enucleation on eyes with non-inflamed orbits (17-21). The diagnoses of all samples were based on the clinical and histopathological information submitted by orbital disease specialists and ocular pathologists (16-21). On account of GO or normal lacrimal gland as the inclusion criteria, the following samples were chosen in the present study: four TED lacrimal samples (GSM2823348, GSM2823349, GSM2823350, GSM2823351) and seven normal lacrimal samples (GSM2823306, GSM2823307, GSM2823308, GSM2823309, GSM2823310, GSM2823311, GSM2823312) were derived from GSE105149, and no sample was repeated; eight TED lacrimal samples (GSM1407195, GSM1407196, GSM1407199, GSM1407200, GSM1407202, GSM1407203, GSM1407204, GSM1407206) and seven normal lacrimal samples (GSM1407233, GSM1407235, GSM1407236, GSM1407237, GSM1407238, GSM1407240, GSM1407243) were obtained from GSE58331, in which eight TED samples were from four patients. Both microarrays have the same platform of GPL570 [HG-U133_Plus_2] Affymetrix Human Genome U133 Plus 2.0 Array.

Screening of the Differentially Expressed Genes

According to the raw data and annotation files, microarray probes were converted into corresponding gene symbols, and the expression value of one gene symbol corresponding to multiple probes was calculated by taking the average value. In addition, Z-score was chosen to normalize the converted data to extend the range of differential expression. The Limma R package (http://www.bioconductor.org/packages/release/bioc/html/limma.html) was used to detect differentially expressed genes (DEGs) between TED and normal lacrimal group. The cut-off criterion of DEGs was P< 0.05 and $|\log$ (fold change) $| \ge 1$. The false discovery rate (FDR) was not used to identify DEGs in

the present study. Nonetheless, we calculated FDR and published it alongside the p-values to give a better idea of the robustness of DEGs.

Gene Ontology and Kyoto Encyclopedia of Genes and Genomes Pathway Analyses

Using the Database for Annotation, Visualization and Integrated Discovery (DAVID, https://david.ncifcrf.gov/), DEGs were mapped to Gene Ontology analysis to detect their possible functional processes from three terms of biological process (BP), molecular function (MF) and cellular component (CC). Moreover, Kyoto Encyclopedia of Genes and Genomes (KEGG) pathway analysis of DEGs was carried out to investigate their potential biological pathways by the KEGG Orthology-based Annotation System 2.0 (KOBAS 2.0, http://kobas.cbi.pku.edu. cn). P < 0.05 was considered as statistically significant difference.

Construction of Protein-Protein Interaction Network

DEGs were entered into the Search Tool for the Retrieval of Interacting Genes (STRING) database (https://string-db.org/) to construct a functional protein-protein interaction (PPI) network. An interaction with the composite score > 0.4 was considered statistically significant. Then, PPI network complex was visualized with Cytoscape software. The central node genes were screened out with the cut-off criteria of connectivity degree ≥10. Whereafter, the Molecular Complex Detection (MCODE) in Cytoscape software was utilized to identify densely connected regions from PPI network complex. The connected region with MCODE score >3 and node number >4 was considered to be the significant module.

Gene Set Variation Analysis

Gene set variation analysis (GSVA) is a nonparametric and unsupervised clustering approach to estimate the score of gene-set enriched pathways based on the expression profile of each sample (22). For GSVA analysis, we chose HALLMARK gene sets as the reference, t value > 2 and p value < 0.05 as the cut-off to identify significantly altered pathways. The GSVA R package (http://www.bioconductor.org/packages/release/bioc/html/GSVA.html) was used to calculate a pathway enrichment score between TED and normal lacrimal group. GSVA analysis for GSE105149, GSE58331, the combined data of GSE105149 and GSE58331 were performed, respectively.

RESULTS

Screening of the Common Molecular Origins Associated with GO Lacrimal Enlargement

Using P< 0.05 and |log (fold change) | \geq 1 as the cut-off criterion, data from GSE105149 and GSE58331 were separately analyzed by the limma R package to identify DEGs. Representative heat maps showed different expression of all DEGs in the TED

lacrimal samples relative to normal lacrimal samples (**Figure 1A**). 80 up-regulated and 440 down-regulated genes were present in the TED lacrimal samples of GSE105149, 868 up-regulated and 252 down-regulated genes were in the TED lacrimal samples of GSE58331. The overlap between the two microarray datasets contained 173 genes, including 20 up-regulated and 153 down-regulated genes in the TED lacrimal glands (**Figure 1B**, **Table S1**). Collectively, the common DEGs in two different gene expression profiles provided evidence on the common molecular origins of lacrimal gland enlargement in GO patients.

Investigation of the Potential Functions and Pathways in GO Lacrimal Enlargement

To investigate the mechanism, the overlapping DEGs were further measured through Gene Ontology and KEGG pathway analyses to explore the possible functions and pathways in the pathogenesis of GO lacrimal enlargement. Based on DAVID software, we observed a total of 81 Gene Ontology assignments, including 31 BPs, 9 MFs, and 41 CCs. The top five significant terms of Gene Ontology classification were illustrated in Figures 2A-C. In the BP category, most DEGs were involved in RNA splicing, protein folding in ER, cell-cell adhesion, mRNA processing and ER to Golgi vesicle-mediated transport. In the MF category, most DEGs may play roles in RNA binding, unfolded protein binding, peptidyl-prolyl cis-trans isomerase activity, cis-trans isomerase activity and nucleotide binding. In the CC category, a large percentage of the DEGs were associated with intracellular organelle lumen, nuclear lumen, cytosol, Golgi apparatus and ER. Furthermore, the KEGG analysis showed the dysfunction of protein processing in ER, phagosome and antigen processing and presentation pathways in GO lacrimal samples (Figure 2D). Taken together, these findings suggested that multiple functions and pathways were significantly associated with GO lacrimal enlargement.

Identification of the Most Significant Pathway and Hub Genes in GO Lacrimal Enlargement

To identify the most significant DEGs "hub genes" associated with GO lacrimal enlargement, we performed a PPI network construction of all the overlapping DEGs on the basis of the STRING database. Finally, 157 nodes and 282 edges were present in the PPI network complex (Figure 3A), in which 16 central node genes were selected with the cut-off criteria of connectivity degree ≥ 10 (Figure 3B). Subsequently, module analysis was performed by the plug-in MCODE in Cytoscape software in order to identify the significant modules from the PPI network complex. Then, 6 modules successfully emerged from the above PPI network complex, including 40 genes. According to the selection criteria of MCODE score > 3 and node number > 4, module a, module b, module c and module d were regarded as significant modules (Figure 3C). Among them, module A with the highest MCODE score and the maximum node number was considered as the most important module. Therefore, the common genes present in 16 central nodes and module A were

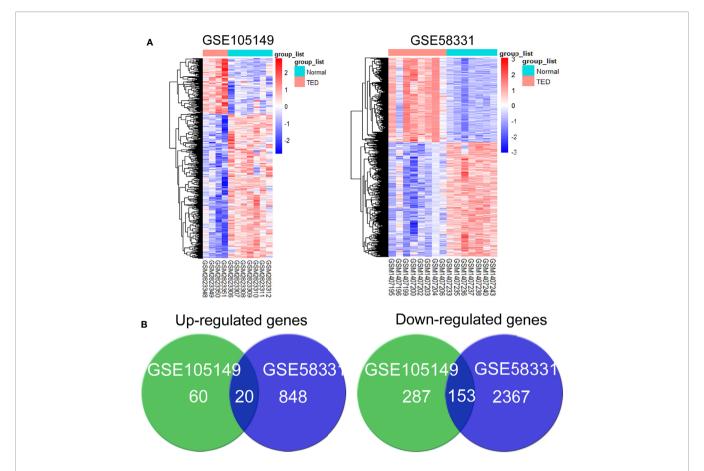


FIGURE 1 | Screening of the common differentially expressed genes associated with lacrimal enlargement in thyroid eye disease (TED). (A) hierarchical clustering heat maps showed different expression of all differentially expressed genes (DEGs) in the TED lacrimal samples relative to normal lacrimal samples. Red indicates that the expression of genes is relatively upregulated, and blue indicates that the expression of genes is relatively downregulated. (B) The DEGs were identified with the P-value <0.05 and |log (fold change)≥1 between thyroid eye disease (TED) and normal lacrimal samples from the mRNA expression profiling sets GSE105149 and GSE58331. Venn diagram showed the respective frequency of up-regulated and down-regulated genes in the TED lacrimal samples.

selected as hub genes, including HSP90AA1, DNAJC10, HNRNPM, HNRNPD, HNRNPC, CANX, HSPA5, HSP90B1, SRSF11, DDX5, RBM25, HNRNPA2B1, and HSPA9 (Figure **3C**, **Table 1**). Besides, function and pathway analyses of DEGs in the modules or hub genes were further carried out. The functional annotations of the genes involved in the modules were protein folding in ER, poly (A) RNA binding, focal adhesion (Figures 4A-C). Pathway analysis showed that the module genes mainly had a close relationship with protein processing in ER, spliceosome and pathogenic Escherichia coli infection (Figure 4D). Moreover, the 13 hub genes were significantly enriched in multiple functions such as protein folding in ER, poly (A) RNA binding and extracellular matrix (Figures 5A-C). And KEGG revealed that these hub genes were enriched in several pathways, especially protein processing in ER (Figure 5D).

According to Gene Ontology and KEGG pathway analyses for the entire DEGs set, 40 module genes and 13 hub genes, we observed that protein processing in ER was the most significant pathway, in which five hub genes including *HSP90AA1*, *HSP90B1*, *DNAJC10*, *HSPA5* and *CANX* were involved.

Differences in Pathway Activities between GO and Normal Lacrimal Gland

KEGG analysis requires relatively arbitrary selection of DEGs as input, only a small number of pathways were enriched in GO. In order to correct this selection to some extent, we applied GSVA analysis to investigate the critical pathways of GO. Since GSVA transforms data from a gene to a gene set based on the sample matrix in the coordinate frame, the enrichment pathway of each sample can be evaluated. As a result, GSVA analysis of TED versus normal lacrimal samples significantly revealed four upregulated and five down-regulated signatures in the TED lacrimal samples of GSE105149 (Figure 6A), five up-regulated and 19 down-regulated signatures in the TED lacrimal samples of GSE58331 (Figure 6B). The common enriched signals in both GSE105149 and GSE58331 were KRAS_SIGNALING_DN, MYOGENESIS, MYC TARGETS V1, UNFOLDED PROTEIN RESPONSE, MTORC1_SIGNALING, PROTEIN_SECRETION and ANDROGEN_RESPONSE. In addition, the GSVA analysis of the combined data of GSE105149 and GSE58331 further supported the above results (Figure 6C). These results further supported the differences in pathway activities between GO and

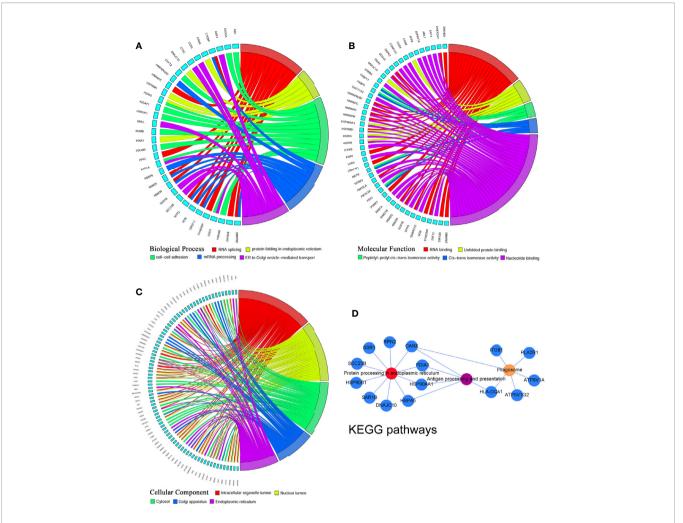


FIGURE 2 | Gene Ontology terms and Kyoto Encyclopedia of Genes and Genomes (KEGG) pathways of the common differentially expressed genes (DEGs) set. Circos plot represents the relationship between genes and the top 5 significant terms including biological process (A) cellular component (B) and molecular function (C). Different colors from red, yellow, green, blue to purple indicate that the P values are sorted from minimum to maximum. (D) KEGG pathway analysis showed that DEGs were mainly enriched in pathways related to protein processing in endoplasmic reticulum, phagosome, and antigen processing and presentation. Different colors from red, yellow to purple indicate that the P values are sorted from minimum to maximum.

normal lacrimal gland, and provided novel or related pathways compared with KEGG pathway analysis.

DISCUSSION

Currently, our understanding of GO pathogenesis is mainly focused on the extraocular muscle and orbital adipose tissue. The molecular origins of lacrimal gland enlargement in GO has not yet been elucidated. In this work, two gene expression profiles from different cohort studies were analyzed to identify DEGs with common changes between lacrimal gland from patients with GO and that from normal groups. Further bioinformatics analysis was performed to investigate the significance of these DEGs, including Gene ontology analysis, KEGG pathway analysis, PPI network construction, hub gene identification and GSVA analysis.

We identified a total of 173 common DEGs which may be involved in the most significant function and pathway in ER, including BPs "Protein folding in ER" "ER to Golgi vesiclemediated transport", MFs "Peptidyl-prolyl cis-trans isomerase activity" "Cis-trans isomerase activity" and CCs "Golgi apparatus" "ER" and pathways "Protein processing in ER" "Phagosome". As we known, ER is like a special factory for eukaryotic cells, in which at least one-third of molecules are synthesized, folded and transported (23). It maintains proper physiological proteostasis that the correctly folded molecules are modified and transported into the Golgi complexes, while the misfolded or unfolded molecules are trapped in the ER together with the molecular chaperone complex. When a large number of misfolded and unfolded molecules are accumulated, ER stress occurs and stimulates the unfolded protein response (UPR). Significantly, UPR intersects with different networks of inflammation, stress signals, and oxidative stress, in which NF-

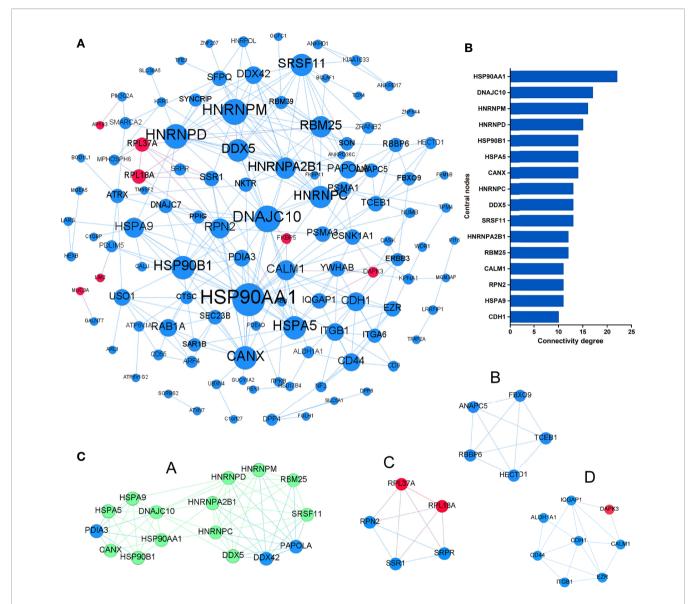


FIGURE 3 | Protein-protein interaction network of the common differentially expressed genes (DEGs) set. (A) The protein-protein interaction (PPI) network was visualized using Cytoscape software. The node size was proportional to the connectivity degree. The genes with no connectivity were not present in the network.

(B) A total of 16 central node genes were screened out with the criteria of filtering degree ≥ 10 from the PPI network complex. (C) Module a, module b, module c, and module d were regarded as significant modules with the selection criteria of MCODE score > 3 and node number > 4. Red indicates that the expression of genes is relatively upregulated, and blue indicates that the expression of genes is relatively downregulated. In the module a, green indicates that the common genes present in 16 central nodes and module a were selected as hub genes.

kB and JNK-AP1 pathways are involved (24). Inflammation, as an important pathological change of GO, directly affects the severity and activity of GO. Clinically, glucocorticoids can effectively alleviate the condition of some active moderate to severe GO patients by controlling the inflammatory response (25). In addition, previous reports have confirmed the dysfunctions of NF-kB pathway and oxidative stress in the progression of GO (26–29). In this study, GSVA analysis also showed the dysregulation of protein secretion, oxidative phosphorylation, UPR and inflammatory response in lacrimal

samples of GO. Therefore, these above evidences supported the ER and related signaling networks as potential mechanisms of GO lacrimal enlargement.

Protein processing in ER was identified as the most significant pathway associated with the pathogenesis of GO lacrimal enlargement, in which 5 hub genes including *HSP90AA1*, *HSP90B1*, *DNAJC10*, *HSPA5*, and *CANX* were involved. These proteins respectively encoding by *HSP90B1*, *DNAJC10*, *HSPA5*, and *CANX* are all located in the ER. Of them, HSP90AA1 has the highest connectivity in the PPI construction of DEGs, which

TABLE 1 | The hub genes associated with lacrimal gland enlargement in Graves' ophthalmopathy.

| Gene symbol | GSE105149 | | | GS | Regulation | | |
|-------------|---------------------------|----------|-------------------|---------------------------|------------|-------------------|------|
| | False discoveryrate (FDR) | P value | Log (fold change) | False discoveryrate (FDR) | P value | Log (fold change) | |
| CANX | 5.56E-02 | 2.59E-04 | -1.02 | 3.25E-02 | 2.35E-03 | -1.00 | down |
| HSPA9 | 4.77E-02 | 1.22E-04 | -1.07 | 3.24E-02 | 2.33E-03 | -1.35 | down |
| RBM25 | 1.11E-01 | 5.34E-03 | -1.08 | 2.42E-02 | 8.83E-04 | -1.58 | down |
| HNRNPC | 7.67E-02 | 1.53E-03 | -1.11 | 3.46E-02 | 2.78E-03 | -1.18 | down |
| DNAJC10 | 5.80E-02 | 3.59E-04 | -1.13 | 3.09E-02 | 2.03E-03 | -1.57 | down |
| DDX5 | 6.31E-02 | 5.59E-04 | -1.14 | 4.36E-02 | 5.04E-03 | -1.78 | down |
| HNRNPM | 3.95E-02 | 9.20E-06 | -1.16 | 1.77E-02 | 1.57E-04 | -1.45 | down |
| HNRNPD | 7.82E-02 | 1.65E-03 | -1.26 | 3.61E-02 | 3.14E-03 | -1.34 | down |
| HSP90AA1 | 7.42E-02 | 1.11E-03 | -1.27 | 2.51E-02 | 1.01E-03 | -1.53 | down |
| HSP90B1 | 5.43E-02 | 2.25E-04 | -1.27 | 2.90E-02 | 1.68E-03 | -1.16 | down |
| HSPA5 | 7.43E-02 | 1.21E-03 | -1.43 | 3.22E-02 | 2.30E-03 | -1.34 | down |
| HNRNPA2B1 | 5.01E-02 | 1.73E-04 | -1.43 | 6.57E-02 | 1.17E-02 | -1.29 | down |
| SRSF11 | 6.10E-02 | 4.20E-04 | -1.54 | 2.47E-02 | 9.66E-04 | -1.42 | down |

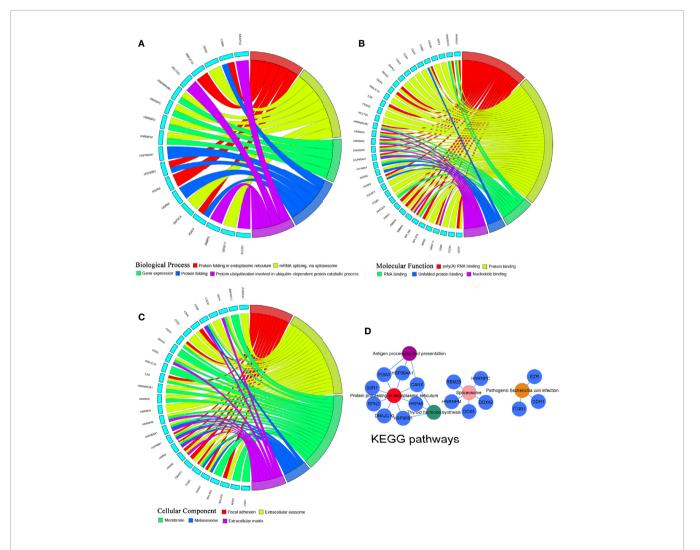


FIGURE 4 | Gene Ontology terms and Kyoto Encyclopedia of Genes and Genomes (KEGG) pathways of the module differentially expressed genes (DEGs) set.

(A) Biological process, (B) Cellular component, (C) Molecular function, (D) KEGG pathways: Colors from red, pink, yellow, green to purple indicate that the P values are sorted from minimum to maximum.

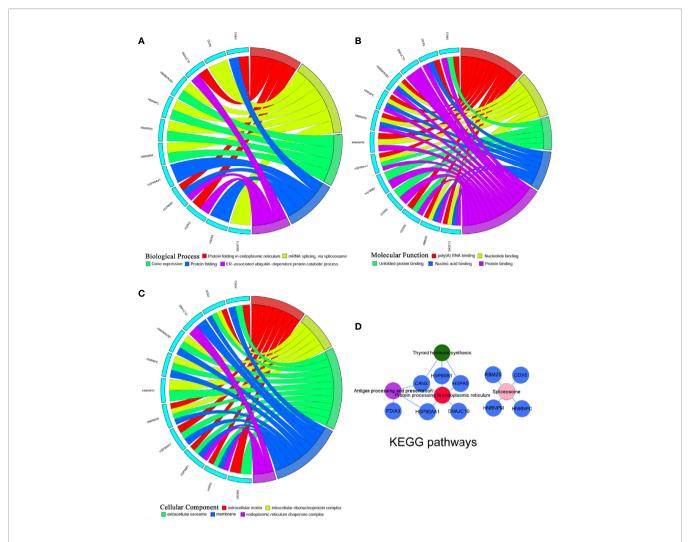


FIGURE 5 | Gene Ontology terms and Kyoto Encyclopedia of Genes and Genomes (KEGG) pathways of the hub differentially expressed genes (DEGs) set. **(A)** Biological process, **(B)** Cellular component, **(C)** Molecular function, **(D)** KEGG pathways: Colors from red, pink, green to purple indicate that the P values are sorted from minimum to maximum.

encodes the well-known cytoplasmic Hsp90a, a highly conserved molecular chaperone that facilitates folding, maturation, structural maintenance and appropriate modification of target molecules (known as clients) (30). These clients are enriched in vesicle-mediated transport, telomere maintenance, mitotic signal transduction, apoptosis, cell-cycle progression, innate immunity and targeted protein degradation (31). For example, the binding of Hsp90α to lipopolysaccharide (LPS) can mediate LPS-induced inflammation, including the secretion of TNF by monocytes (32). Interestingly, LPS-induced inflammation has been proved to affect the differentiation of orbital pre-adipocytes in GO (33). HSP90B1 encodes another Hsp90 isoform GRP94, an ERresident molecular chaperone, which plays a vital role in the processing and transportation of secreting proteins (34). GRP94 shows many biochemical characteristics with other HSP90 family members, but also exhibits different activities, such as calcium binding, which is required for conditions in the ER (35).

Studies have shown that GRP94 functions as a key molecular chaperone for various Toll-like receptors and integrins (36, 37), and also regulates innate and adaptive immunity (38). The absence of GRP94 in the hematopoietic system of mice leads to thrombocytopenia, prolonged bleeding, and massive platelets, which are clinically difficult to distinguish from human Bernardsoulier Syndrome (39). In addition, DNACJ10 and HSPA5 respectively encode the ER-resident proteins ERdj5 and BiP, which are crucial parts of ER-associated degradation (ERAD) complex involved in recognizing and degrading misfolded or unfolded molecules (40). ERAD is an essential quality-control system for ER molecules, in which the correctly folded molecules are transported to the secretion pathway and finally misfolded or unfolded molecules are translocated into the cytosol for proteasome degradation (41). Ushioda et al. confirmed that ERdj5 acts as a reductase that breaks the disulfide bonds of misfolded or unfolded molecules and facilitates ERAD by the

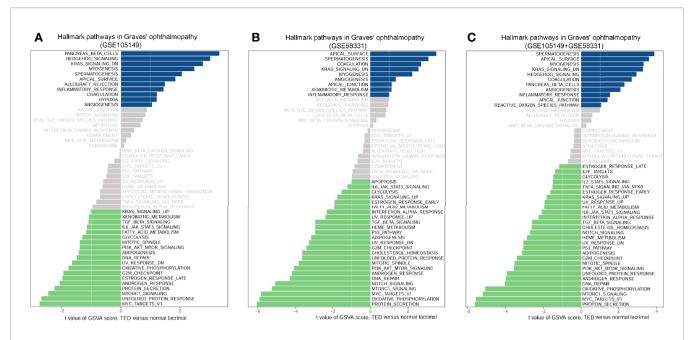


FIGURE 6 | Scores of gene set variation analysis (GSVA) for pathway activities between thyroid eye disease (TED) and normal lacrimal gland. **(A)** GSVA analysis of four TED versus seven normal lacrimal samples of GSE105149. **(B)** GSVA analysis of seven TED versus seven normal lacrimal samples of GSE58331. **(C)** The common GSVA analysis of 11 TED versus 14 normal lacrimal samples of GSE105149 and GSE58331. Shown are t values from a linear model, corrected for patient of origin. DN, down; v1, version.

physical and functional connections with BiP (42). Furthermore, studies suggested ERdj5 and BiP as the master regulator of calcium homeostasis and redox homeostasis in the ER (43). Similarly, Calnexin (CANX) is a key ER-resident chaperone to help folding and quality control, thus ensuring that only correctly folded and assembled molecules can further enter into the secretion pathway (44). All these findings further deepen our understanding of the important roles of HSP90AA1, HSP90B1, DNAJC10, HSPA5 and CANX in protein processing in ER. Meanwhile, the observation that the down-regulated expression of these 5 hub genes in GO lacrimal samples, further indicated that ER dysfunction may contribute to the enlargement of GO lacrimal gland, and may extend to other pathological manifestations of GO.

In summary, our study presented a comprehensive bioinformatics analysis to identify potential key genes and pathways that may help to uncover the molecular mechanisms of the lacrimal gland enlargement in GO. Our findings firstly revealed that down-regulated expression of HSP90AA1, HSP90B1, DNAJC10, HSPA5, and CANX might generate the dysfunction of protein processing in ER, providing a novel explanation for the etiology and molecular events of the lacrimal enlargement in GO. When further molecular experiments confirmed the findings of the identified candidate genes and pathways in GO, the underlying molecular mechanisms of lacrimal gland enlargement in GO would be elucidated. All of these would contribute to a novel insight into GO pathogenesis.

DATA AVAILABILITY STATEMENT

The data sets (GSE105149 and GSE58331) for this study can be found in Gene Expression Omnibus (http://www.ncbi.nlm.nih.gov/geo/).

AUTHOR CONTRIBUTIONS

WT and YS conceived the experiments. JY, ZM, and XJ analyzed the data. WT drafted the manuscript. All authors contributed to the article and approved the submitted version.

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SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at: https://www.frontiersin.org/articles/10.3389/fendo.2020. 571151/full#supplementary-material

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Conflict of Interest: Author Jia Yao was employed by company Chengdu SuAn Technology Co., Ltd.

The remaining authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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Immunological Features of Paranasal Sinus Mucosa in Patients with Graves' Orbitopathy

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Lu Y, Wu Y, Huang Y, Fang S, Li Y, Sun J and Zhou H (2021) Immunological Features of Paranasal Sinus Mucosa in Patients with Graves' Orbitopathy. Front. Endocrinol. 11:621321. **Background:** Previous studies showed that patients with Graves' orbitopathy (GO) had concomitant mucosal abnormality within the paranasal sinuses. It remains unknown whether the immunological reactions in sinus mucosa affect the orbit inflammation in GO.

Methods: Patients with GO underwent sinus computed tomography (CT) scans for sinus mucosal disease by two independent reviewers using the Lund-MacKay systems. Ethmoid mucosal samples were collected during orbital decompression surgeries for patients with GO and correction surgeries for patients with old orbital fractures as controls. Histological analysis and immunofluorescence were performed in all sinus mucosa tissues. Flow cytometry analysis was used to examine the immunological features of sinus mucosa in both GO and control groups.

Results: Immunohistochemistry showed that the paranasal sinus mucosa of patients with GO grew swelling, with goblet cell and small vessel proliferation, endothelial cell swelling, and inflammatory cell infiltration. The number of T helper (Th)1, Th17, and gamma-delta T cells in nasal sinus mucosa of patients with GO increased significantly compared with those from controls. Further, the proportion of Th1 cells was significantly correlated with clinical activity score. In addition, there was a decreased number of regulatory T cells in patients with GO. The number of Th2 cells showed no significant difference between the two groups. Finally, the proportion of interleukin-22-producing cell subsets in gammadelta T cells of patients with GO was significantly increased compared with those from controls.

Conclusions: Our observations illustrated a potential pathogenic role of mucosal-infiltrating T cells, which may have the possibility to aggravate inflammatory responses in GO.

Keywords: immune, T cells, paranasal sinus, mucosa, Graves' orbitopathy

INTRODUCTION

Graves' orbitopathy (GO) is an autoimmune disorder ranking first in orbital diseases. GO is also the most common extrathyroidal manifestation of Graves' disease (GD) (1). Orbital tissue remodeling is a major pathological feature of GO, which includes extraocular muscle swelling, adipogenesis, fibrosis, and extracellular matrix deposition such as glycosaminoglycan (2). It is generally accepted that the bony orbit confines the orbital soft tissues on all sides except anteriorly. Intriguingly, the two most frequently enlarged extraocular muscles are the inferior and medial rectus muscles, which are next to ethmoid and maxillary sinus, respectively. Both muscles overlie the thinnest bones of the orbit (3). Gouveris et al. reported that patients with GD had concomitant mucosal thickening within the paranasal sinuses. Patients with GO who underwent orbital decompression surgeries exhibited a higher prevalence (29.4%) of sinus mucosal thickening on CT scan. Histological exams showed chronic rhinosinusitis in 45.5% of GO patients (4). These findings may be explained by circulating autoantibodies targeting autoantigens, namely thyrotropin receptor and insulin-like growth factor-1 receptor, associated with GD and found to be expressed in sinus mucosa (5).

Until today, the pathogenesis of GO is not fully understand. Many studies have indicated that T cells are critical in GO autoimmune responses. They have confirmed that at least three T helper (Th) cell groups are involved in the development of GO, which are Th1 cells, Th2 cells, and Th17 cells. Early studies mainly focused on Th1/Th2 subsets and their related cytokines in the inflammation of GO. It is recognized that Th1 cells play a significant role in inducing cytokine release in early active GO, and the chronic fibrotic phase of GO is dominated by Th2-type immune responses (6-9). Furthermore, Th17 cells have been proved to be an important proinflammatory and profibrotic cell subsets. In GO patients, the level of interleukin (IL)-17A and the number of IL-17A-producing T cells in the peripheral blood were higher than those of healthy controls. IL-17A was shown to promote inflammation and fibrosis of orbital fibroblasts (OF) derived from GO patients (10-12). Moreover, Th17 cells could stimulate the expression of proinflammatory cytokines such as IL-6, IL-8, monocyte chemoattractant protein-1, tumor necrosis factor-α, etc., and costimulatory molecules to regulate the balance of fibrosis and adipogenesis in different OF subsets (13). In this current study, we raise a hypothesis that there is a relationship between nasal sinuses inflammation and GO pathology. We sought to investigate why there is a higher prevalence of sinus mucosa disorder in GO patients and to unravel the potential immunological features of paranasal sinus mucosa in GO.

MATERIALS AND METHODS

Subjects and Samples

A collection of 58 ethmoid sinus mucosa tissues (38 samples of patients with GO undergoing orbital decompression surgery and 20 samples of patients with old orbital fracture undergoing

orbital repair surgery) were included in the study. Informed consent was provided by all participants. Use of these samples was approved by the Ethical Committee of Shanghai Ninth People's Hospital, Shanghai JiaoTong University School of Medicine. All 38 GO patients were in inactive stage. Steroid or immunosuppressive treatment was discontinued for at least 3 months before surgery, and all patients were clinically euthyroid at the time of surgery. Exclusion criteria included age less than 18 years old, and patients with history of sinus surgery. Patients with rheumatic or hemolymph diseases were also excluded. Based on the findings of pre-operative computer tomography (CT) scans, scores based on the Lund-Mackay staging systems (Table 1) were calculated for each patient and control subject.

Histology and Immunofluorescence

For histological analysis, nasal sinus mucosa tissues were fixed in 4% formalin and paraffin-embedded. The specimens were cut in 4-µm-thick sections before staining with hematoxylin and eosin (HE). The sections were then examined with a light microscope. The expression of CD4 and CD8 was determined by immunofluorescence analysis. Cryostat sections (4 μ m thick) were fixed with 4% paraformaldehyde fixative at room temperature for 20 min and washed in PBS for three times. The sections were then incubated with a blocking solution (normal goat serum) at room temperature for another 20 min. Next, they were incubated with rabbit anti-CD4 and mouse anti-CD8 (Abcam, USA) primary antibodies overnight at 4°C. After being washed in PBS for three times, the sections were incubated with secondary antibodies for 1 h at room temperature. Finally, coverslipmounted slides were observed and photographed under a fluorescence microscope.

Flow Cytometry Analysis

The ethmoid sinus mucosa tissues were digested with 5% Type IV collagenase into single cell suspensions, and were stimulated with phorbol 12-myristate 13-acetate (50ng/ml; Sigma-Aldrich, USA) and ionomycin (1μ g/ml; Sigma-Aldrich, USA) in the presence of

TABLE 1 | Lund-Mackay Scoring System.

| Anatomic Location | Scoring Criteria* |
|---------------------------|--------------------------|
| Maxillary sinuses | 0: No abnormality |
| | 1: Partial opacification |
| | 2: Total opacification |
| Anterior ethmoid sinuses | 0: No abnormality |
| | 1: Partial opacification |
| | 2: Total opacification |
| Posterior ethmoid sinuses | 0: No abnormality |
| | 1: Partial opacification |
| | 2: Total opacification |
| Sphenoid sinuses | 0: No abnormality |
| | 1: Partial opacification |
| | 2: Total opacification |
| Frontal sinuses | 0: No abnormality |
| | 1: Partial opacification |
| | 2: Total opacification |
| Ostiomeatal complex (OMC) | 0: Not occluded |
| . , , | 2: Occluded |

^{*}Total score can range from 0 to 24.

Golgi Plug (1 μl/ml; BD Biosciences, USA) at 37°C for 6 h before flow cytometry. To analyze T cell subsets in sinus mucosa, cells were incubated with APC-Cy7-Fixable Viability Dye (eBioscience, USA) before staining for surface markers (FITC-anti-CD3, AF700-anti-CD8, BV650-anti-γδTCR) (All from BD Biosciences, USA). Then, cells were treated with Fixation/Permeabilization reagents (eBioscience, USA) and stained with intracellular markers (BV711-anti-IFN-γ, BV421-anti-IL-13; BD Biosciences, USA; PE-anti-IL-17A, APC-anti-FOXP3, PE-Cy7-anti-IL-22; eBioscience, USA). A BD LSRFortessa X-20 was applied for sample sorting. The data were analyzed after adjusting the fluorescence compensation with FlowJoV10.

Statistical Analysis

All values are expressed as the mean ± standard deviation (SD). Statistical analyses were performed by one-way ANOVA, analysis of Spearman's correlation and t-test followed by the appropriate tests with the statistical analysis program GraphPad Prism 6.0 (GraphPad Prism Software, USA) and SPSS 25 (IBM SPSS Software, USA). A *P* value less than 0.05 was considered to indicate a statistically significant difference.

RESULTS

Clinical Characteristic and Radiographic Findings of GO Patients

Clinical characteristics of patients and controls were described in **Table 2**. As expected, the GO group had more proptosis reflected in higher exophthalmometry readings than controls (23.37 \pm 2.44 vs. 15.24 \pm 2.73, P < 0.0001) mm. CT scans demonstrated a higher score in patients with GO compared with controls for Lund-MacKay system (4.00 \pm 1.94 vs. 1.67 \pm 0.58, P < 0.0001). Among GO patients, the most common location for mucosal thickening was in the ethmoid sinuses (69.2%), followed by the

TABLE 2 | Clinical Characteristic and Radiographic Finding of GO and normal control.

| | GO | Control | P value |
|---------------------------------|-------------------|------------------|----------|
| Number, n | 38 | 20 | |
| Male/Female, n/n | 16/22 | 11/9 | 0.41 |
| Age(years), mean ± SD | 44.80 ± 11.64 | 40.91 ± 12.21 | 0.38 |
| Smoker, n (%) | 13 (34.2%) | 11 (55.5%) | 0.16 |
| Proptosis(mm), mean ± SD | 23.65 ± 2.31 | 15.24 ± 2.73 | < 0.0001 |
| Course of disease (months), | 23.84 ± 20.62 | / | |
| mean ± SD | | | |
| Sinus mucosal thickening, n (%) | 15 (39.5%) | 3 (15%) | 0.08 |
| Lund-Mackay Score, mean ± SD | 3.84 ± 1.83 | 1.67 ± 0.58 | < 0.0001 |
| Involved nasal sinuses, n (%) | | | |
| Ethmoid | 11 (73.3%) | 1 (33.3%) | 0.25 |
| Maxillary | 9 (60.0%) | 3 (100%) | 0.51 |
| OMC | 6(40.0%) | 0 | 0.51 |
| Frontal | 0 | 0 | 1 |
| Sphenoid | 4 (26.7%) | 1 (33.3%) | 1 |

Mucosal thickening defined as Lund-MacKay score \geq 1.

GO, Graves' orbitopathy; CT, computed tomography; SD, standard deviation; OMC, osteomeatal complex.

maxillary (53.8%), OMC (46.2%), and sphenoid sinuses (30.8%). The representative CT scanning was showed in **Figure 1A**. When comparing the extent of sinus mucosal thickening to the degree of orbital proptosis as measured by exophthalmometry, higher Lund-MacKay scores did not correlate with greater orbital proptosis (r=-0.1386, P=0.582).

Histological Changes of Paranasal Sinus Mucosa in GO Patients

Histological changes of the sinus mucosa were found in 21 out of the 38 GO patients (60%). The mucosa showed thickening in GO patients. Compared with normal nasal sinus, typical histological changes include mucosal swelling and epithelial cell proliferation. Some cases show goblet cell proliferation and mucus over-secretion; chronic inflammatory cell infiltration and small vessel proliferation (**Figure 1B**). $CD4^+$ and $CD8^+$ T cells infiltration can be found in GO patients' mucosa, which was detected by immunofluorescence staining (**Figure 1C**). There was a significant increase of $CD4^+$ (29.78 \pm 15.16 vs. 10.60 \pm 3.78, P=0.024)/HPF and $CD8^+$ T cells (29.00 \pm 10.33 vs. 15.33 \pm 4.76, P=0.013)/HPF in GO mucosa compared with controls (**Figure 1D**).

Immunological Characteristics of Paranasal Sinuses Mucosa in GO

T cell populations present in the nasal mucosa from GO patients and normal controls were identified as CD4/CD8/γδ T cells by using flow cytometric analysis. The proportion (%) of CD3⁺ to total living cells (Figure 2A) was higher in GO patients compared with normal controls. (33.69 \pm 15.84% vs.17.15 \pm 9.04%, P<0.001). Th1 cells were defined as CD3⁺CD8⁻IFN- γ ⁺ subsets and Th2 cells as CD3+CD8-IL-13+ subsets. Our data indicated that Th1 cells increased significantly in GO patients' mucosa (7.19 \pm 4.69% vs. 1.74 \pm 1.67%, P<0.01), while Th2 cells showed no difference between GO patients and normal controls $(9.37 \pm 6.70\% \text{ vs. } 6.90 \pm 5.10\%, P=0.16)$ (**Figure 2B**). In patients with GO, the mucosa was also characterized by significantly increased Th17 cells (CD3⁺CD8⁻IL-17A⁺) (1.91 ± 0.99% vs. $0.84 \pm 0.63\%$, P<0.001) (Figure 2C) and decreased Treg cells $(CD3^{+}CD8^{-}FOXP3^{+})$ (1.38 ± 1.39% vs. 3.04 ± 1.62%, P<0.001) (Figure 3A), which suggested a similar Th-driven inflammation to orbital connective tissue. In addition to the changes of Th cells in the mucosa, we further investigated into the difference of $\gamma\delta$ T cells between the two groups, which play an important role in mucosal immunity. Our results showed that γδT cells was significantly elevated in GO patients' mucosa (1.47 \pm 1.05% vs. 0.60 ± 0.38 , P<0.01) (**Figure 3B**). Additionally, we examined IL-17A expression in those sinus mucosal $\gamma \delta T$ cells and found that there was no difference between the two groups. Intriguingly, we observed a significantly augmented amount of IL-22-secreting $\gamma \delta T$ cells in the mucosa of GO patients compare with controls $(24.32 \pm 14.76\% \text{ vs. } 8.42 \pm 6.30\%, P=0.04)$ (Figure 3C).

The Relationship Between Th1, Th2, Th17 Cells in Nasal Sinus Mucosa and Clinical Characteristics of GO

The proportion of different subtypes of T cells was further analyzed with GO clinical activity score (CAS). Our data

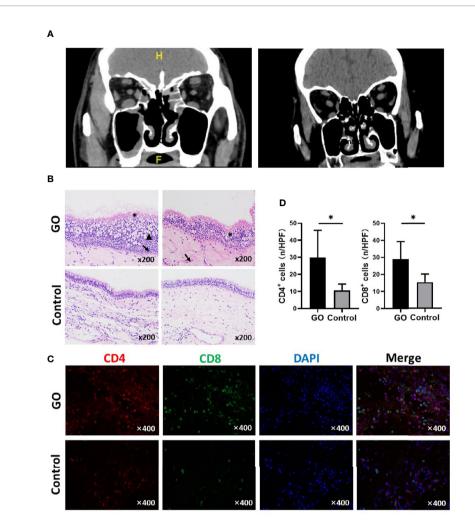


FIGURE 1 | Orbital computed tomography (CT) images and nasal sinus mucosa histological presentation of Graves' orbitopathy (GO) patients and controls. (A) Left: A GO patient shows mucosal thickening of maxillary sinus and ethmoid sinus on both sides; Right: Normal control with right orbital fracture. (B) Typical changes include mucosal swelling and epithelial cells proliferation. *Disorder of ciliary columnar epithelial; †epithelial cell proliferation; Mucus over-secretion. (HE staining, ×200). (C) Immunofluorescence staining results of CD4 and CD8 positive cells in GO and control mucosa. (D) Statistical analyses of CD4 and CD8 positive cells, the mucosa of GO has a significant increase compared with control. Significance (P) values after t-test was represented by *P < 0.05.

indicated that the proportion of Th1 cells was significantly correlated with disease activity ($R^2 = 0.11$, P = 0.0474), while there was no correlation between the number of Th2 cells ($R^2 = 0.07$, P = 0.136) or Th17cells ($R^2 = 0.02$, P = 0.446) and CAS (**Figure 4A**). The study cohort was then divided into two groups according to the time course of GO. One group was longer than 2 years, and the other was less than 2 years. The subtypes of T cells were reanalyzed on the basis of disease course. Our data indicated that the average proportion of Th1 cells ($8.23 \pm 5.03\%$ vs. $5.62 \pm 3.40\%$) and Th17 cells ($2.04 \pm 1.16\%$ vs. $1.74 \pm 0.62\%$) declined with the extension of disease course, while Th2 cells ($8.27 \pm 5.77\%$ vs. $11.11 \pm 8.02\%$) slightly increased with GO duration. However, there was no significant difference in any of the T cell subtypes between these two groups (P = 0.11, 0.40, 0.24) (**Figure 4B**), which may attribute to the limited number of cases.

DISCUSSION

Previous studies suggested that disorders of the paranasal sinus mucosa were associated with a number of systemic autoimmune diseases. Patients with GD were reported to have significantly greater paranasal sinus mucosal thickening on CT scan than normal controls. The presence of autoantigens TSHR and IGF-1R were also identified in paranasal sinus mucosa (5). These studies hold the opinion that autoantibodies circulating in the blood of patients with GD would be expected to bind to either TSHR or IGF-1R in the sinus mucosa, as occurs in the thyroids and orbital connective tissues, resulting in the localized mucosal edema and inflammatory cell infiltration.

T cell immunity plays an important role in the pathogenesis of GO (14). The frequency of Th1 cells and the Th1/Th2 ratio

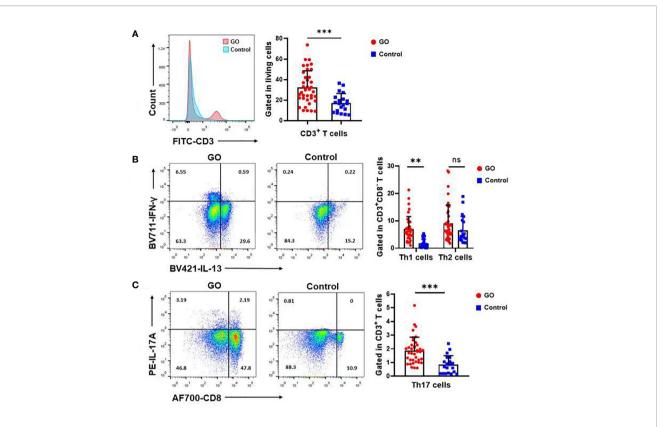


FIGURE 2 | Th1, Th17 but not Th2 cells varied in Graves' orbitopathy (GO) patients compared with controls. **(A)** The percentage (%) of CD3⁺ cells to total living cells was higher in GO patients compared with controls. **(B)** Flow cytometry showed that Th1 cells increased significantly in GO patients' mucosa, while Th2 cells showed no significant difference between GO patients and controls. Significance (*P*) values after t-test was represented by **P < 0.01, ***P < 0.001. **(C)** Flow cytometry showed that Th17 cells increased significantly in GO patients' mucosa. Significance (*P*) values after t-test was represented by ***P < 0.001.

were positively correlated with the inflammatory activity score of GO (8). Wakelkamp et al. (15) found that active GO patients were characterized by Th1 type cytokines and there was no direct correlation between Th2 type cytokines and disease progression. Han et al. further demonstrated that IFN-γ secreted by Th1 cells and IL-4 secreted by Th2 cells can promote the production of hyaluronic acid by orbital fibroblasts. However, our research did find that there was no difference between GO and control groups in Th2 population. Some studies have revealed that Th1 cells may predominate in the orbit in early GO and Th2 cells might play a greater role in later stages of the disease (16, 17). Yet, others studies have indicated that the active phase is characterized by the presence of proinflammatory and Th1derived cytokines, whereas other cytokines, including Th2derived cytokines, do not seem to be linked to a specific stage of GO (18, 19). However, we found most of the studies outdated. Notably, these studies defined Th2 cells by means of establishing T cell clones or examining bulk RNA expression of cytokines within orbital connective tissues due to technical limitation at that time, which made their results not convincing enough. In order to explain the phenomenon, the characteristics of chronic sinusitis should be considered. Chronic rhinosinusitis (CRS) is a persistent inflammatory disease affecting paranasal sinuses. CRS

is categorized into two distinct subgroups defined as CRS with nasal polyps (CRSwNP) and CRS without nasal polyps (CRSsNP) (20). Th1 cytokines are mostly found in CRSsNP and Th2 cytokines in CRSwNP. The sinus mucosal abnormality in GO patients isn't usually accompanied by polyps, which might explain the Th1 but not Th2 dominance observed in our study cohort.

We and other groups reported that in GO patients the level of IL-17A and the number of IL-17A-producing T cells were higher than those of healthy controls in the periphery blood as well as in orbital connective tissues (10–12, 21, 22). These results indicate that the CD4⁺ T cells may contribute to the immunopathological process in GO. In this current study, we also found that there was a higher proportion of both Th1 cells and Th17 cells in the sinus mucosa of GO patients. Combined with previous studies, we suspect that these lymphocytes in nasal sinus may infiltrate into the orbit through the very thin bone walls and blood vessels.

Furthermore, we studied Treg cells in the sinus. Treg cells are a suppressive subset of CD4⁺ T subsets important for the regulation of immune responses in various autoimmune diseases. Treg cell dysfunction also contributes to the development of autoimmune thyroid disease (23–25). A study showed that patients with improved GO were more likely to have higher proportion of Treg

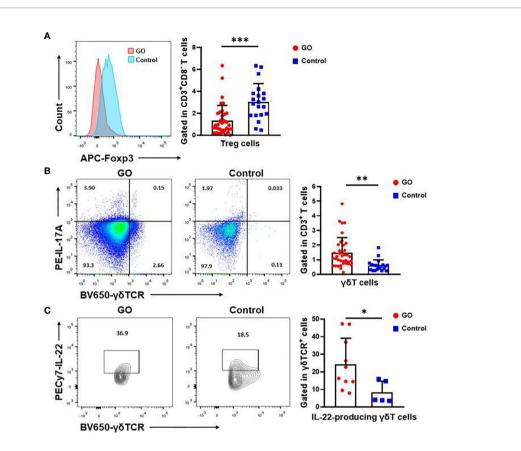


FIGURE 3 | The significant change of Treg cells and $\gamma\delta T$ cells in the nasal sinus mucosa of Graves' orbitopathy (GO) patients. **(A)** There was also a significant decrease in the percentage (%) of Treg cells in GO patients. Significance (*P*) values after t-test was represented by ***P < 0.001. **(B)** Flow cytometry showed that $\gamma\delta T$ cells increased significantly in GO patients' mucosa. Significance (*P*) values after t-test was represented by **P < 0.01. **(C)** There was also a significant increase in the percentage (%) of IL-22 secreting $\gamma\delta T$ cells in GO patients. Significance (*P*) values after t-test was represented by *P < 0.05.

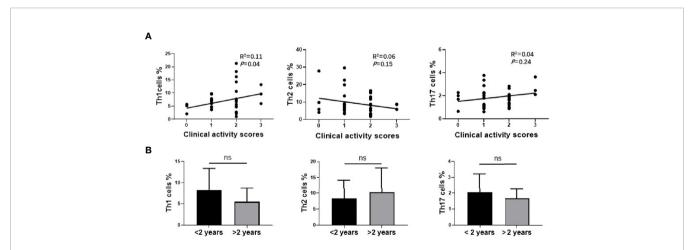


FIGURE 4 | The relationship between Th1, Th2, Th17 cells in nasal sinus mucosa and clinical characteristics of Graves' orbitopathy (GO). (A) The analysis of Spearman's correlation between Th1, Th2, Th17 cells and CAS of GO patients. Th1 cells amount was correlated with CAS of GO patients significantly. (B) There was no significant difference between two groups of different disease course in Th1, Th2 or Th17 cells.

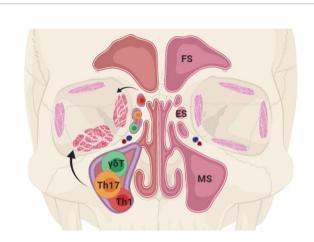


FIGURE 5 | Theoretic model for the relationship between immunological abnormality in nasal sinus of Graves' orbitopathy (GO) patients and orbital inflammation. FS, frontal sinus; ES, ethmoid sinus; MS, maxillary sinus (Declaration: this figure was designed by the first author Yi Lu with the help of drawing website "https://biorender.com").

cells than those with stable or deteriorated GO. Thus, the number of Treg cells in the peripheral blood of GO patients can be used as a predictor of clinical course (26). Kahaly et al. (27) demonstrated that the proportion of Treg cells in the peripheral blood leukocytes derived from GO patients increased after incubation with rabbit polyclonal anti-T lymphocyte globulin. Taken together, our findings reinforced the point of view that the decline of Treg cells might be responsible for the activation of inflammatory cells in GO.

Despite CD4⁺ and CD8⁺ T cells, γδT cells are a distinctive subset of T cells that were first recognized by Brenner in 1986. These cells primarily colonize in the mucosa and skin where they play an important role in immune regulation (28). In autoimmune disorders, γδT cells have immunoregulatory properties and secrete both IL-17A and IL-22 (29). A previous study showed that orbitalinfiltrating T cells were primarily $\gamma\delta T$ cells, while the classic $\alpha\beta T$ helper cells were rare (30). Intriguingly, we previously confirmed an elevated subset of IL-17A-producing $\gamma\delta T$ cells in the circulation of GO patients compared with healthy controls (12). However, in this current study, we did not find the up-regulation of IL-17Aproducing γδT cells, while IL-22-producing γδT cells were observed to be increased in GO sinus mucosa. IL-22 can affect the fibroblasts in the skins, intestines, and joints. Proliferation of synovial fibroblasts in rheumatoid arthritis patients as well as monocyte chemoattractant protein-1-dependent recruitment of monocytes to the sites of inflammation are driven by IL-22 (31). Furthermore, IL-22 can increase the production of antiinflammatory factors such as IL-11 as well as inflammatory mediators such as IL-6 and CXCLs chemokines in human colonic myofibroblast cells (32, 33). These results indicate that the infiltration of $\gamma \delta T$ cells in sinus mucosa might be associated with the complicated and refined regulation of orbital inflammation in patients with GO.

This research helps us build a preliminary understanding of the relationship between sinus disorder and GO. It could represent the local effects of sinus for the adjacent orbital inflammation. Inflammatory cells and cytokines may spread from the sinus to the orbit *via* natural dehiscence in adjacent bony walls or by hematogenous spread through common blood vessels (**Figure 5**). All in all, the immune abnormality of the nasal sinus mucosa in GO patients may not only be a clinical manifestation of Graves' disease, but also be involved in the pathogenesis of GO.

DATA AVAILABILITY STATEMENT

The original contributions presented in the study are included in the article/supplementary material. Further inquiries can be directed to the corresponding author.

ETHICS STATEMENT

The studies involving human participants were reviewed and approved by Ethic Committee of Shanghai Ninth People's Hospital. The patients/participants provided their written informed consent to participate in this study.

AUTHOR CONTRIBUTIONS

Research conception and design: YLu, YW, SF, HZ. Experiment implementation: YLu, YW, JS, YLi. Analysis and interpretation of data: YLu, YH, SF. Drafting of manuscript: YLu, YW. Critical revision: SF, HZ. All authors contributed to the article and approved the submitted version.

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Conflict of Interest: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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Integrative Analysis of Proteomics and DNA Methylation in Orbital Fibroblasts From Graves' Ophthalmopathy

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Virakul S, Sompam P, Pisitkun T, van der Spek PJ, Dalm VASH, Paridaens D, van Hagen PM, Hirankam N, Palaga T and Dik WA (2021) Integrative Analysis of Proteomics and DNA Methylation in Orbital Fibroblasts From Graves' Ophthalmopathy. Front. Endocrinol. 11:619989. doi: 10.3389/fendo.2020.619989 **Background:** Graves' ophthalmopathy (GO) is a frequent extrathyroidal complication of Graves' hyperthyroidism. Orbital fibroblasts contribute to both orbital tissue inflammation and remodeling in GO, and as such are crucial cellular elements in active GO and inactive GO. However, so far it is largely unknown whether GO disease progression is associated with functional reprogramming of the orbital fibroblast effector function. Therefore, the aim of this study was to compare both the proteome and global DNA methylation patterns between orbital fibroblasts isolated from active GO, inactive GO and healthy controls.

Methods: Orbital fibroblasts from inactive GO (n=5), active GO (n=4) and controls (n=5) were cultured and total protein and DNA was isolated. Labelled and fractionated proteins were analyzed with a liquid chromatography tandem-mass spectrometer (LC-MS/MS). Data are available *via* ProteomeXchange with identifier PXD022257. Furthermore, bisulphite-treated DNA was analyzed for methylation pattern with the Illumina Infinium Human Methylation 450K beadchip. In addition, RNA was isolated from the orbital fibroblasts for real-time quantitative (RQ)-PCR. Network and pathway analyses were performed.

Results: Orbital fibroblasts from active GO displayed overexpression of proteins that are typically involved in inflammation, cellular proliferation, hyaluronan synthesis and adipogenesis, while various proteins associated with extracellular matrix (ECM) biology and fibrotic disease, were typically overexpressed in orbital fibroblasts from inactive GO. Moreover, orbital fibroblasts from active GO displayed hypermethylation of genes that linked to inflammation and hypomethylated genes that linked to adipogenesis and autoimmunity. Further analysis revealed networks that contained molecules to which

both hypermethylated and hypomethylated genes were linked, including NF- κ B, ERK1/2, Alp, RNA polymerase II, Akt and IFN α . In addition, NF- κ B, Akt and IFN α were also identified in networks that were derived from the differentially expressed proteins. Generally, poor correlation between protein expression, DNA methylation and mRNA expression was observed.

Conclusions: Both the proteomics and DNA methylation data support that orbital fibroblasts from active GO are involved in inflammation, adipogenesis, and glycosaminoglycan production, while orbital fibroblasts from inactive disease are more skewed towards an active role in extracellular matrix remodeling. This switch in orbital fibroblast effector function may have therapeutic implications and further studies into the underlying mechanism are thus warranted.

Keywords: graves' ophthalmopathy, orbital fibroblast, proteomics, DNA methylation, epigenetics

INTRODUCTION

Fibroblasts are crucial for maintaining tissue homeostasis, and are major producers of important cellular mediators for inflammatory and tissue remodeling processes during normal healing responses, but also under pathological conditions, including chronic inflammatory and fibrotic diseases (1). Chronic tissue inflammation and fibrosis are characterized by excessive fibroblast accumulation in affected tissues (1). Fibroblast accumulation occurs through different mechanisms, including enhanced proliferation by tissue resident CD34 fibroblasts, recruitment of fibrocytes (a population of circulating cells with fibroblast-like properties that express CD34⁺ and extracellular matrix (ECM) molecules) and diminished apoptosis/prolonged survival (2-4). These fibroblasts can alter their phenotype and effector functions, as evidenced by their differentiation into myofibroblasts (5, 6). Myofibroblasts are considered to represent the end-effector cells in fibrotic reactions. Myofibroblasts express α-smooth muscle actin that facilitates cellular contraction and tissue distortion and actively produce high amounts of ECM molecules, such as collagen and hyaluronan (1).

Graves' ophthalmopathy (GO) is a disease of the orbital softtissues surrounding the eyes, and affects up to 50% of patients with Graves' disease (GD) (7). In general, GO patients first suffer from an initial phase of progressive disease (the 'active' phase) that is characterized by active inflammation (8). This active phase may last for months after which the activity subsides and progresses to a phase of slow spontaneous recovery. This 'chronic' phase may take months to years and is associated with pathological tissue alterations, including adipose tissue expansion, excessive hyaluronan accumulation and fibrosis (8). The orbital tissue alterations are largely responsible for several morbidities such as proptosis, chronic eye movement dysfunction, and eventually determines the 'severity' of GO (7). Previously, protein expression profiles of orbital tissues from GO with different disease activity and smoking status revealed differences in several proteins involved in inflammation and

adipogenesis (9, 10). However, those proteins could not be specifically linked to their pathogenic cellular source.

Orbital fibroblasts represent the main effector cells in GO as they are crucially involved in regulation of both the local inflammatory and tissue remodeling responses. The effector functions of the orbital fibroblasts are in turn affected by several mediators, including the thyrotropin receptor (TSHR) stimulatory autoantibodies, cytokines, growth factors and physical cellular interactions (8). Importantly, orbital fibroblasts from GO patients may react in a different way to such stimuli than orbital fibroblasts isolated from healthy orbital tissue. For instance a stronger upregulation of CD40, Thy1, IGF1R and BAFF by GO orbital fibroblasts was reported (11-14). Moreover, orbital fibroblasts isolated from active GO display enhanced secretion of certain pro-inflammatory mediators in comparison to orbital fibroblasts from control tissue, even in the absence of further in vitro stimulation (15, 16). Furthermore, differentiation of orbital fibroblasts into adipocytes and profibrotic myofibroblasts appears to be more restricted to the late inactive stage of disease and is associated with increased TSHR expression by these cells (17, 18).

Gradually, epigenetic regulation (e.g. histone modification, DNA methylation and non-coding RNA) has become the subject of studies in primary fibroblasts isolated from different fibroproliferative diseases, including idiopathic pulmonary fibrosis (IPF) and systemic sclerosis (SSc) (4, 19-22). Fibroblasts from IPF and SSc have altered DNA methylation profiles, which contribute to the regulation of gene transcription in these cells (19, 20). In GO, differential gene expression has been demonstrated in orbital tissue, suggesting a role for epigenetics in the pathogenesis of GO (23). However, data on DNA methylation in orbital fibroblasts from GO is lacking to date. Insight in this could help to develop novel therapeutic options that target aberrant epigenetic modifications in GO. Here we conducted a study to compare the proteome of orbital fibroblasts isolated from active GO, inactive GO and control orbital tissue and integrated these data with global DNA methylation analysis performed on the same orbital fibroblasts.

MATERIAL AND METHODS

Orbital Fibroblast Isolation From Orbital Tissue

Orbital fibroblasts were cultured from four patients with GO in an active stage (clinical activity score (CAS) $\geq 3/7$) and five patients with GO in an inactive stage of disease (CAS < 3/7) (24), who underwent orbital decompression surgery. In addition, orbital fibroblasts were cultured from five controls without thyroid or inflammatory disease that underwent orbital surgery for other reasons, as described previously (25). GO patients were euthyroid and had not received immunosuppressive treatment for at least three months prior to orbital decompression surgery. Further patient characteristics are given in Table 1. All orbital tissues were obtained at the Rotterdam Eye Hospital (Rotterdam, the Netherlands), after informed consent and in accordance with the principles of the Declaration of Helsinki. Approval was given by the local medical ethics committee (protocol ID-2007-01). Orbital fibroblasts were cultured in Dulbecco's modified Eagle's medium (DMEM) supplemented with 10% fetal calf serum (FCS) and antibiotics (penicillin and streptomycin; Cambrex BioWhittaker, Verviers, Belgium) (25). Orbital fibroblasts were serially passaged with gentle treatment of trypsin/EDTA and used for experiments between the 2nd and 6th passage.

Proteome Analysis

Protein Extraction, Peptide Preparation and Labeling

Orbital fibroblasts from four patients with GO at an active stage and five GO patients at an inactive stage of disease, and five controls were included in this experiment. Orbital fibroblasts (5 x 10⁶ cells) were obtained and pelleted after which HaltTM Protease Inhibitor Cocktail and Phosphatase Inhibitor Cocktail (Thermo Fisher Scientific, Rockford, IL) was added before digestion and homogenization in 5% sodium deoxycholate with a sonicator. Protein concentration was determined with PierceTM BCA Protein Assay Kit (Thermo Fisher Scientific, Rockford, IL). For peptide preparation, 100 µg of protein was digested with 2 µg trypsin and labelled with 10-plex tandem mass tag (TMT) reagent according to the manufacturer's protocol (Thermo Fisher Scientific, Rockford, IL). Afterwards, the peptide mixture was passed

TABLE 1 | Patients and controls data.

| Participants | Cell ID | Age | Gender |
|-----------------|---------|-----|--------|
| Healthy control | COF-3 | 48 | Female |
| | COF-4 | 82 | Female |
| | COF-7 | 80 | Female |
| | COF-9 | 59 | Female |
| | COF-10 | 37 | Female |
| Inactive GO | GO-11 | 43 | Female |
| | GO-27 | 69 | Female |
| | GO-31 | 46 | Female |
| | GO-56 | 51 | Female |
| | GO-89 | 32 | Female |
| Active GO | GO-19 | 31 | Female |
| | GO-37 | 62 | Female |
| | GO-88 | 77 | Male |
| | GO-90 | 84 | Female |

through the Pierce High pH Reversed-Phase Peptide Fractionation Kit (Thermo Fisher Scientific, Rockford, IL).

Peptide Analysis

The labelled fractionated peptides were analyzed with a liquid chromatography tandem-mass spectrometer (LC-MS/MS) (Q Exactive Plus mass spectrometer; Thermo Scientific, San Jose, CA). The LC-MS/MS methods included a full MS scan at a resolution of 70,000 followed by 10 data-dependent MS2 scans at a resolution of 37,500. The normalized collision energy of higher-energy collisional dissociation (HCD) fragmentation was set at 28%. MS scan range of 400 to 1,600 m/z was selected and precursor ions with unassigned charge states, a charge state of +1 and a charge state of greater than +8 were excluded. Proteome discoverer 2.1 software (Thermo Scientific, San Jose, CA) was used to analyze the MS raw data files with database containing forward and reverse peptide sequences from the human Uniprot Database. The search parameters were set for the fix modifications of carbamidomethylation of cysteine (+57.02146 Da) and TMT modifications (+229.2634 Da) at N-terminal and lysine, while oxidation of methionine (+15.99491 Da) was set for the variable modification. A maximum of four modifications and two missed cleavages per peptide were allowed. Parent and fragment monoisotopic mass errors were set at 10 ppm and 0.2 Da, respectively. A target-decoy approach was used to limit the false discovery rate (FDR) of the identified peptides to less than 1%. Differential protein expression was compared between orbital fibroblasts isolated from active and inactive GO by Student's ttest. The mass spectrometry proteomics data have been deposited to the ProteomeXchange Consortium via the PRIDE [1] partner repository with the dataset identifier PXD022257. Network, pathway and functional analysis were further performed on these differential expressed proteins using Ingenuity (Qiagen) filtering only for those networks and pathways that are based on experimentally obtained information.

DNA Methylation Analysis

Total DNA was extracted from orbital fibroblasts of active GO (n = 4), inactive GO (n = 4) and controls (n = 4). Global DNA methylation was measured using the Illumina Infinium Human Methylation 450K beadchip on bisulphite-treated DNA. DNA methylation analysis was performed by GenomeStudio methylation analysis package. Methylation level ranges from 0 (unmethylated) to 1 (fully methylated). Differential DNA methylation levels/patterns were analyzed by two approaches; 1) genes with differential DNA methylation using FDR < 0.05, and 2) differences in DNA methylation level of at least 2-fold between active and inactive GO. This was followed by network, pathway and functional analysis using Ingenuity (Qiagen) filtering only for those networks and pathways that are based on experimentally obtained information.

mRNA Expression by Orbital Fibroblasts

Messenger RNA was isolated (GenElute Mammalian Total RNA Miniprep Kit; Sigma-Aldrich, St. Louis, MO, USA) from orbital fibroblasts from the same culture experiment as used for the

proteome and DNA methylation studies. The isolated mRNA was converted into cDNA and gene expression was determined by real-time quantitative (RQ)-PCR (QuantStudio 5 Real-Time PCR System; Applied Biosystems, Waltham, MA) and normalized to the control gene *ABL*, as described previously (25). Primer-probe combinations (*ABL*) and TaqMan gene expression assays (Life technologies, Foster, CA) used are listed in **Supplementary Table 1**. Data from mRNA expression was analyzed using ANOVA and subsequently analyzed with the Mann Whitney U test. A P-value < 0.05 was considered statistically significant.

RESULTS

Proteome Analysis in Orbital Fibroblasts

Twenty-five proteins differed significantly in their expression level between orbital fibroblasts from patients with active GO versus inactive GO (**Supplementary Table 2**). Cluster analysis revealed that the differentially expressed proteins clustered into three main groups (**Figure 1A**, indicated as cluster 1, cluster 2 and cluster 3). Proteins (n=6) in cluster 2 displayed higher expression in orbital fibroblasts from inactive GO compared to orbital fibroblasts from active GO and controls (**Figure 1B**). The proteins (n=16) in cluster 3 were higher expressed in orbital fibroblasts from active GO compared to inactive GO and controls (**Figure 1C**). The proteins (n=3) in cluster 1 were expressed at slightly higher level in orbital fibroblasts from active GO than orbital fibroblasts from inactive GO and controls.

Validation of Differential Protein Expression by RQ-PCR

For eleven of the proteins found to be differentially expressed (4 up-regulated and 7 down-regulated in inactive GO orbital fibroblasts compared to active GO) the expression levels of the corresponding mRNA molecules were determined by RQ-PCR (Figures 1B, C). From the 4 proteins (PSMB4, FBN2, COL6A1 and NCAM2) upregulated, only NCAM2 mRNA was significantly higher expressed (P<0.05) in orbital fibroblasts from inactive GO compared to active GO (Figure 1B), while the mRNA levels of PSMB4, FBN2 and COL6A1 only showed a trend to higher expression in inactive GO orbital fibroblasts compared with active GO orbital fibroblasts (Figure 1B). Of the 7 proteins (PACSIN3, NFKB1, SMC3, GFER, GSDMD, UGDH and MT1X) that were expressed at significantly lower levels in orbital fibroblasts from inactive GO, none of the corresponding mRNAs was significantly decreased in inactive GO (Figure 1C). Unexpectedly, NFKB1 mRNA expression was significantly upregulated in inactive GO orbital fibroblasts compared to active GO (Figure 1C).

Orbital Fibroblast Expressed Proteins in Active GO Relate to Inflammation, Hyaluronan and Adipogenesis

Proteins overexpressed by orbital fibroblasts from active GO related to inflammation, hyaluronan synthesis and adipogenesis.

One of the inflammation related proteins highly expressed in active GO orbital fibroblasts was NFKB1 (**Figure 1C**). The NFKB1 105 kD protein can be processed by the 26S proteasome into the DNA binding p50 subunit of the transcription factor NF- κ B (26). NF- κ B stimulates expression of genes involved in inflammation in many different cell types, including orbital fibroblasts (27, 28). The NFKB1 subunit exerts a protective role by dampening inflammation (29). Therefore, upregulation of NFKB1 may represent a counter regulatory mechanism to control the inflammatory response in orbital fibroblasts from active GO (8, 11).

ECM and fibrosis related proteins that were overexpressed in orbital fibroblasts from active GO included structural maintenance of chromosomes 3 (SMC3), growth factor, augmenter of liver regeneration (GFER) and uridine diphosphate-glucose dehydrogenase (UGDH) (Figure 1C). UGDH is involved in the biosynthesis of glycosaminoglycans (GAG), such as hyaluronan, chondroitin sulfate and heparan sulfate and was previously found elevated in urine from patients with active GO (30). Moreover, UGDH was found to be expressed at higher levels in orbital fibroblasts than in fibroblasts from other anatomical regions (31). This indicates that orbital fibroblasts in active GO acquire a phenotype very well equipped for GAG synthesis (32). Orbital fibroblasts from active GO also produced more protein kinase c and casein kinase substrate in neurons (PACSIN) 3, a protein involved in clathrin-mediated endocytosis that regulates glucose uptake by adipocytes (33, 34). The up-regulation of PACSIN3 in the active orbital fibroblasts may therefore relate to formation of adipocytes that finally accumulate in the inactive phase of GO (35).

Orbital Fibroblast Expressed Proteins in Inactive GO Relate to Extracellular Matrix Biology

Orbital fibroblasts from inactive GO overexpressed several proteins linked to ECM biology and fibrotic diseases. This included fibrillin-2 (FBN2), collagen type VI alpha 1 chain (COL6A1) and the neural adhesion molecule 2 (NCAM2) (Figure 1B) (36-39). FBN2 is involved in elastic fiber formation and is abundantly present in embryonic tissues, but is also elevated along with elastin in fibrotic tissue, for instance in systemic sclerosis (36, 40). Yet, data on elastin accumulation in orbital tissue from GO patients is lacking so far. The non-fibrillar collagen type VI is involved in structural organization of the ECM, for instance by interacting with a multitude of other key ECM components, including the fibrillary type-1 and type 3 collagens and hyaluronan (39, 41, 42). Increased COL6A1 tissue levels are present in lung fibrosis, liver fibrosis and keloid scarring (37-39). Furthermore, collagen type VI can bind a variety of different growth factors implicated in fibrosis, and as such serves as a reservoir that can regulate growth factor activity in the vicinity of fibroblasts and contribute to fibrosis (39, 42, 43). Although data on collagen type VI in GO orbital tissue is lacking so far, several growth factors that bind collagen type VI, including PDGF-AB, PDGF-

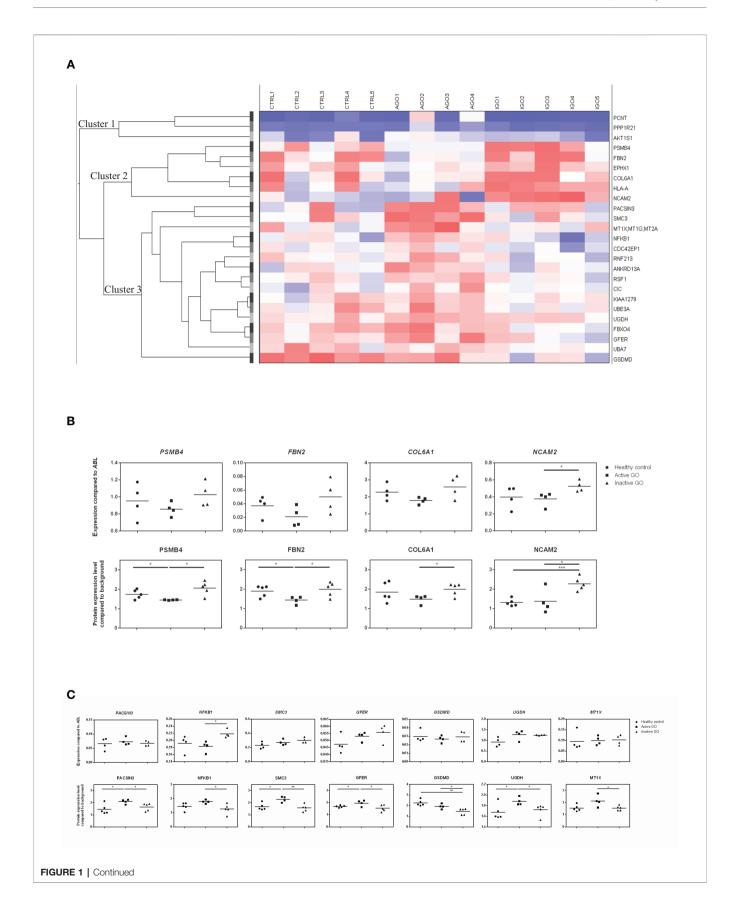


FIGURE 1 | Proteomic profiles of orbital fibroblasts from inactive GO, active GO and healthy controls. **(A)** Cluster analysis of the differential protein expression between orbital fibroblasts from patients with inactive GO (n = 5), active GO (n = 4) and healthy controls (n = 5). Differentially expressed proteins grouped in three main clusters. Proteins within cluster 1 proteins were expressed at slightly higher level in orbital fibroblasts from active GO compared with orbital fibroblasts from inactive GO and controls. Proteins within cluster 2 displayed higher expression in orbital fibroblasts from inactive GO compared with orbital fibroblasts from active GO and controls. Proteins within cluster 3 were expressed at a higher level in orbital fibroblasts from active GO compared to inactive GO and controls. **(B)** Gene and protein expression level from proteins within cluster 2. Protein expression levels were compared to high abundant protein expression level. Four proteins (PSMB4, FBN2, COL6A1 and NCAM2) up-regulated in inactive GO orbital fibroblasts compared to active GO were related to extracellular matrix biology and further determined by RQ-PCR. * and *** indicate p-value < 0.05 and < 0.001, respectively. Individual symbols represent orbital fibroblast cultures from individual patients. Horizontal bar depicts the median. **(C)** Gene and protein expression level Seven proteins (PACSIN3, NFKB1, SMC3, GFER, GSDMD, UGDH and MT1X) down-regulated in inactive GO orbital fibroblasts compared to active GO related to inflammation, hyaluronan and adipogenesis were determined by RQ-PCR. * and ** indicate p-value < 0.05 and < 0.01, respectively. Individual symbols represent orbital fibroblasts cultures from individual patients. Horizontal bar depicts the median.

BB, and HGF have been linked to GO pathogenesis (44–46). Neural adhesion molecule 2 (NCAM2) was also higher expressed by orbital fibroblasts from inactive GO. NCAM2 can activate fibroblast growth factor receptor (FGFR), and recent studies demonstrated that FGFR activation in orbital fibroblasts stimulates hyaluronan synthesis and adipogenesis (47–49). Therefore, it can be postulated that in late inactive GO, orbital fibroblast-derived NCAM2 might be involved in adipogenesis and hyaluronan synthesis through FGFR activation. Clearly further detailed study on the role of the molecules we here found to be expressed at higher levels in orbital fibroblasts from inactive GO is required. Yet, our data strongly suggest that in the inactive stage of GO the orbital

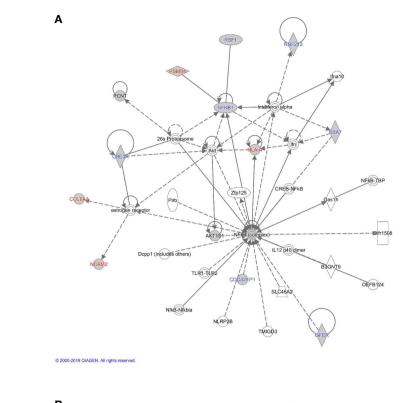
fibroblasts acquire effector functions that are strongly related to ECM remodeling.

Proteome Pathway, Network and Functional Analysis

The proteins differentially expressed between orbital fibroblasts from active GO and inactive GO were fed into Ingenuity with the aim to link these proteins to networks and pathways of interest to GO pathogenesis. The top two identified canonical pathways were UDP-D-xylose and UDP-D-glucuronate biosynthesis (p-value = 0.00217) and protein ubiquitination pathway (p-value = 0.00305) (**Table 2**). Moreover, all these proteins, except MT1X, were linked to 2 networks (**Figure 2**). The first network is associated with

TABLE 2 | Top 5 canonical pathways and diseases and bio-functions from the differentially expressed proteins.

| Top Canonical Pathways | | |
|--|---------------------|---|
| Name | p-value | Overlap |
| UDP-D-xylose and UDP-D-glucuronate Biosynthesis | 0.00217 | 50.0% |
| Protein Ubiquitination Pathway | 0.00305 | 1.1% |
| Crosstalk between Dendritic Cells and Natural | 0.00303 | 2.2% |
| Killer Cells | 0.00416 | 2.2/0 |
| Altered T Cell and B Cell Signaling in | 0.00427 | 2.2% |
| Rheumatoid Arthritis | | |
| OX40 Signaling Pathway | 0.00427 | 2.2% |
| Top Diseases and Bio Functions | | |
| Diseases and Disorders | | |
| Name | p-value | #Molecules |
| Cancer | 0.00000287 - 0.049 | 24 |
| Endocrine System Disorders | 0.00000287 - 0.0436 | 24 |
| Organismal Injury and Abnormalities | 0.00000087 - 0.0498 | 24 |
| Reproductive System Disease | 0.0000428 - 0.0436 | 19 |
| Respiratory Disease | 0.0000551 - 0.0394 | 13 |
| Molecular and Cellular Functions | | |
| Name | p-value | Proteins (listing from the lowest p-value) |
| Cell-To-Cell Signaling and Interaction | 0.000469 - 0.0434 | HLA-A, NFKB1, UBE3A, CDC42EP1, NCAM2, SMC3 |
| Carbohydrate Metabolism | 0.00111 - 0.0445 | UGDH, EPHX1, HLA-A, NFKB1 |
| Cell Death and Survival | 0.00111 - 0.0487 | HLA-A, NFKB1, FBN2, PSMB4, GSDMD, SMC3, COL6A1, GFER, RSF1, AKT1S1 EPHX1, UBA7, UBE3A |
| Cell Morphology | 0.00111 - 0.0434 | NFKB1, PCNT, UBE3A, NCAM2, GFER, GSDMD, HLA-A, CDC42EP1 |
| Cellular Assembly and Organization | 0.00111 - 0.0424 | KIF1BP, ANKRD13A, CDC42EP1, NCAM2, NFKB1, UBE3A, GFER, PCNT, RSF1, FBXO4, HLA-A |



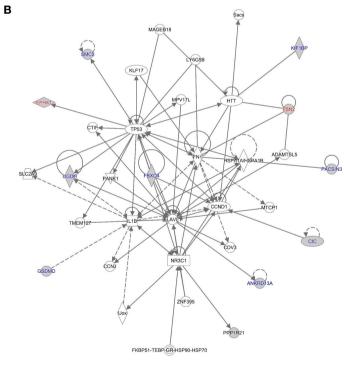


FIGURE 2 | Proteome based network analysis. (A) Network 1: linked to cellular development and connective tissue disorders. Differentially expressed proteins are highlighted in grey. The differentially expressed proteins labelled in red were the up-regulated while differentially expressed proteins labelled in blue were down-regulated in inactive GO comparing to active GO orbital fibroblasts. (B) Network 2: linked to free radical scavenging, DNA replication, recombination and repair and cellular assembly and organization. Differentially expressed proteins are highlighted in grey. The differentially expressed proteins labelled in blue were down-regulated while differentially expressed proteins labelled in blue were down-regulated in inactive GO comparing to active GO orbital fibroblasts.

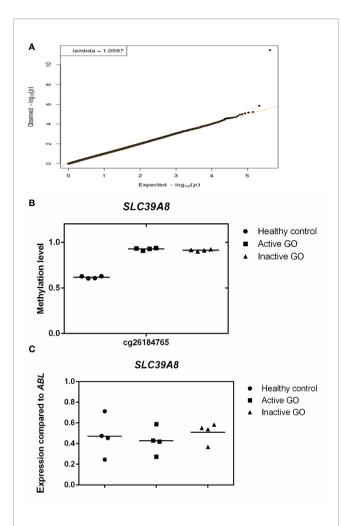


FIGURE 3 | Comparison of orbital fibroblast global DNA methylation profiles with FDR <0.05. **(A)** DNA methylation in active GO orbital fibroblasts (n = 4), inactive GO orbital fibroblasts (n = 4) and healthy control orbital fibroblasts (n = 4) were compared. Global DNA methylation was measured using the Illumina Infinium Human Methylation 450K beadchip on bisulphite-treated DNA and analyzed by GenomeStudio methylation analysis package. Genes with differential DNA methylation were further corrected for multiple testing using FDR < 0.05. **(B)** Methylation level of SLC39A8. Methylation level ranges from 0 (unmethylated) to 1 (fully methylated). Individual symbols represent orbital fibroblast cultures from individual patients. Horizontal bar depicts the median. **(C)** SLC39A8 expression was determined by real-time quantitative (RQ)-PCR and normalized to the control gene ABL. Data from mRNA expression was analyzed using ANOVA and subsequently analyzed with the Mann Whitney U test. Individual symbols represent orbital fibroblast cultures from individual patients. Horizontal bar depicts the median.

cellular development and connective tissue disorders (**Figure 2A**), while the second network is linked to free radical scavenging, DNA replication, recombination and repair and cellular assembly and organization (**Figure 2B**). In network 1, the differentially expressed proteins were linked with NF-κB, Akt, 26s proteasome, interferon alpha and estrogen receptor (**Figure 2A**). On the other hand, the differentially expressed proteins in network 2 were linked to several transcriptional regulators, such as cellular tumor antigen p53 (TP53), huntingtin (HTT), cyclin D1 (CCND1), and also other

(pathogenic) proteins including fibronectin 1 (FN1) and Hsp70-binding protein 1 (HSPA) (**Figure 2B**).

Global DNA Methylation Profile in Orbital Fibroblasts

After adjustment for multiple testing using false discovery rate (FDR < 0.05), none of the genes were differentially methylated when comparing the orbital fibroblasts isolated from active GO (n = 4) and inactive GO patients (n = 4), although hypermethylated GNAS (cg09885502) showed a trend to reach statistical significance (FDR = 0.0774) in inactive GO (Supplementary Figure 1). However, GNAS was not found differentially expressed at the protein level with the current technique used (Supplementary Figure 1). SLC39A8 was the only gene that showed statistically significant difference when comparing control orbital fibroblasts (n = 4) with all GO patients (active and inactive GO; n = 8) (FDR = 1.404 x 10^{-6}) (Figure 3A), being hypermethylated in the 3'UTR region in the orbital fibroblasts from GO patients comparing to healthy control (Figure 3B). Although this could potentially be associated with decreased SLC39A8 gene expression on GO orbital fibroblasts, no difference in mRNA expression was detected between the orbital fibroblast from inactive GO, active GO and controls (Figure 3C). Neither was SLC39A8 protein detected by our current proteomics approach. Gene expression level of enzymes regulating DNA methylation, the writers, DNA methyltransferases (DNMTs), and the erasers, ten eleven translocation (TET), were further examined; however, the expression level of all DNMT and TET were not significantly different in orbital fibroblasts from GO and healthy controls (Supplementary Figure 2).

Hypermethylated Genes in Orbital Fibroblasts From Active GO Patients

We further analyzed differences in methylation level in a less stringent manner by applying a cut-off difference of more than 2-fold between active and inactive GO patients. This resulted in a total of 142 hits, corresponding to 115 coding genes, that were hypermethylated in the orbital fibroblasts from active GO patients (**Figure 4** and **Supplementary Table 3**). Among the top 10 genes (with the fold differences ranging approximately from 3.1 to 6.9-fold), the highest fold difference was detected by a single probe located in the gene body of WDR8 (**Supplementary Figure 3** and **Supplementary Table 3**). Several other genes were detected by more than one probe including Mir548F5, MAB21L1, HTATIP2, TTC12, NIPAL2, TRIM2, PAQR5, OR2L13, RPH3AL, GPR6, MYOM2, DGKQ, ZNF234 and SPAG1 (**Supplementary Figure 3**).

Hypomethylated Genes in Orbital Fibroblasts From Active GO Patients

Applying the cut-off difference of more than 2-fold between active and inactive GO patients yielded 66 hits, corresponding for 63 genes, that were hypomethylated in the orbital fibroblasts from active GO compared with orbital fibroblasts from inactive GO patients (**Figure 5** and **Supplementary Table 4**). The gene set hypomethylated in active GO orbital fibroblasts comprised a.o.: *RNF168* (with the highest fold change of 5.08-fold, as

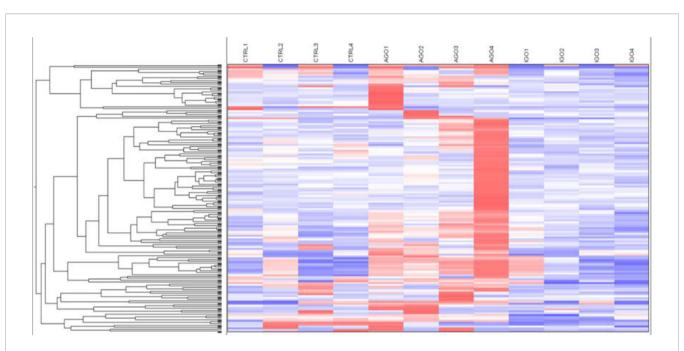


FIGURE 4 | Hypermethylated genes in orbital fibroblasts from active GO in comparison to orbital fibroblasts from inactive GO and healthy controls with a fold difference ≥ 2. DNA methylation in active GO orbital fibroblasts (n = 4) was compared with inactive GO orbital fibroblasts (n = 4). Global DNA methylation was measured using the Illumina Infinium Human Methylation 450K beadchip on bisulphite-treated DNA and analyzed by GenomeStudio methylation analysis package. Hypermethylated genes with differences in DNA methylation level of at least 2-fold in active GO orbital fibroblasts were clustered as shown in the heatmap.

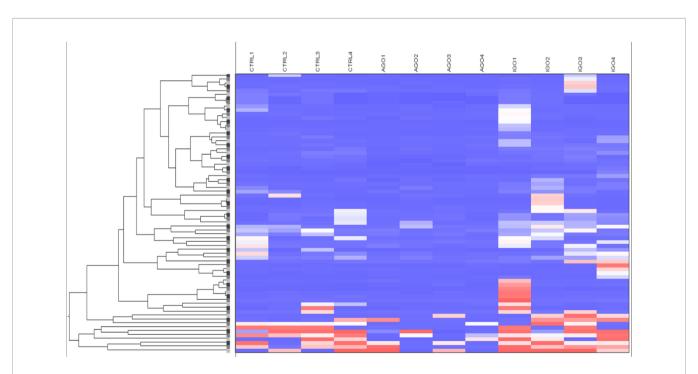


FIGURE 5 | Hypomethylated genes in orbital fibroblasts from active GO in comparison to orbital fibroblasts from inactive GO and healthy controls with a fold difference ≥ 2. DNA methylation in active GO orbital fibroblasts (n = 4) were compared with inactive GO orbital fibroblasts (n = 4). Global DNA methylation was measured using the Illumina Infinium Human Methylation 450K beadchip on bisulphite-treated DNA and analyzed by GenomeStudio methylation analysis package. Hypomethylated genes with differences in DNA methylation level of at least 2-fold in active GO orbital fibroblasts were clustered as shown in the heatmap.

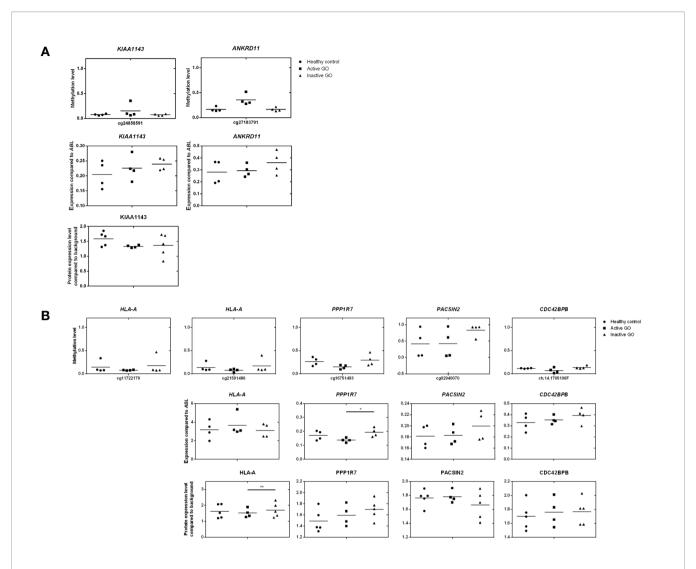


FIGURE 6 | Comparison of DNA methylation, mRNA expression and proteomic data. **(A)** Comparison of hypermethylated genes, mRNA expression and proteomic data. Gene expression was determined by real-time quantitative (RQ)-PCR and normalized to the control gene *ABL*. Data from mRNA expression was analyzed using ANOVA and subsequently analyzed with the Mann Whitney U test. Individual symbols represent orbital fibroblast cultures from individual patients. Horizontal bar depicts the median. **(B)** Comparison of hypomethylated genes, mRNA expression and proteomic data. Gene expression was determined by real-time quantitative (RQ)-PCR and normalized to the control gene *ABL*. Data from mRNA expression was analyzed using ANOVA and subsequently analyzed with the Mann Whitney U test. Individual symbols represent orbital fibroblast cultures from individual patients. Horizontal bar depicts the median.

detected by one probe in the gene body), *SIM2* (detected with 2 different probes located in the gene body: 2.19-fold differences at cg15316660 and 2.03-fold differences at cg23286646 in the gene body), *AIRE* (detected with 2 different probes: 2.35-fold difference at cg27251412 and 2.34-fold difference at cg09510531 200 nucleotides upstream of the transcriptional start site; TSS200) and *HLA-A* (detected with 2 different probes: 2.27-fold difference at cg11722179 and 2.22-fold difference at cg21591486 in the gene body) (**Figure 6B** and **Supplementary Figure 4**).

To investigate the effects of differential methylation at the TSS200 in AIRE and gene-body in HLA-A, mRNA expression levels were determined by RQ-PCR. AIRE gene expression was

not detectable by RQ-PCR in any of the orbital fibroblasts (data not shown), while *HLA-A* was not differentially expressed at the mRNA level between fibroblasts from active GO and inactive GO (**Figure 6B**). AIRE protein expression was not detected by the current proteomics approach (data not shown), while HLA-A protein expression was significantly down-regulated in active GO orbital fibroblasts (**Figure 6B**).

Integrative Analysis of Proteomics and Gene Methylation in Orbital Fibroblasts From Active GO Patients

Upon integration of the DNA hypermethylation data with the proteome data, *KIAA* and the *ANKRD* gene family emerged from

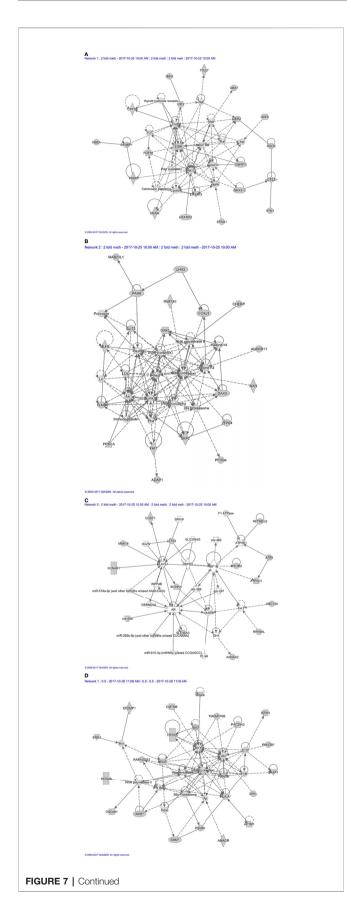


FIGURE 7 | Methylation based network analysis. (A) Network 1 derived from hypermethylated genes in orbital fibroblasts from active GO: linked to neurological disease, organismal injury and abnormalities, and nervous system development and function Network analysis was further performed on these differentially hypermethylated genes (highlighted in grey) using Ingenuity (Qiagen) filtering only for those networks and pathways that are based on experimentally obtained information. (B) Network 2 derived from hypermethylated genes in orbital fibroblasts from active GO: linked to embryogenic development, nervous system development and function, and organ development. Network analysis was further performed on these differentially hypermethylated genes (highlighted in grey) using Ingenuity (Qiagen) filtering only for those networks and pathways that are based on experimentally obtained information. (C) Network 3 derived from hypermethylated genes in orbital fibroblasts from active GO: linked to cardiovascular system development and function, organismal development, and cellular assembly and organization. Network analysis was further performed on these differentially hypermethylated genes (highlighted in grey) using Ingenuity (Qiagen) filtering only for those networks and pathways that are based on experimentally obtained information. (D) Network 4 derived from hypomethylated genes in orbital fibroblasts from active GO: linked to cellular growth and proliferation, nervous system development and function, and cell-to-cell signaling and interaction. Network analysis was further performed on these differentially hypomethylated genes (highlighted in grey) using Ingenuity (Qiagen) filtering only for those networks and pathways that are based on experimentally obtained information.

both types of analyses and were therefore further examined by RQ-PCR. Although hypermethylation of *KIAA1143* and *ANKRD11* genes was more than 2-fold, neither *KIAA1143* nor *ANKRD11* mRNA expression reached statistically significant difference in expression when comparing active and inactive GO orbital fibroblasts (**Figure 6A**). In addition, ANKRD11 protein was not detected by the current proteomics approach (data not shown). Neither the difference on gene expression data and proteome data reached statistical significance nor corresponded to the DNA methylation data.

Integration of the DNA hypermethylation data with the proteome data revealed overlap for PPP1R, PACSIN, and the CDC42 family, which were subsequently further validated by RQ-PCR. Although trends towards elevated mRNA expression for PACSIN2 and CDC42BPB were observed in inactive GO, no statistical significance existed among orbital fibroblast groups, while PPP1R7 mRNA expression was significantly (p < 0.05) higher in inactive GO orbital fibroblasts than in active GO orbital fibroblasts (P < 0.05) (**Figure 6B**). However, neither the mRNA expression data nor the proteome data corresponded to the DNA methylation data.

Network Analysis From Methylated Genes in Orbital Fibroblasts From GO Patients

Next, all the genes that displayed differential DNA methylation between orbital fibroblasts from active GO and inactive GO were fed into Ingenuity pathway analysis to identify potential critical molecular networks involved in GO pathogenesis. The top network with the highest score obtained for hypermethylated genes in active GO orbital fibroblasts genes included TSHR signaling, PDGF-BB signaling and the ERK1/2 signal transduction pathway, as depicted in network 1 (**Figure 7A**). Ingenuity linked this network mainly to neurological disease, organismal injury and abnormalities, and nervous system development and function. *EGR1*, one of the top 10 genes with the highest fold methylation difference, was also found in this network. The second network with the second highest

score (designated as network 2) contained genes that were associated with the PI3K, NF-κB, JNK and Akt signaling pathways (**Figure 7B**). Ingenuity linked this network mainly to embryogenic development, nervous system development and function, and organ development. Clearly, these signaling cascades exert several important roles in immune activation as well as fibrosis (27, 44, 50, 51). In addition, *Mir548F5* was found in another network (network 3) together with *MYOM2* and *RPH3AL* and were connected to IGF-1R, androgen receptor and various noncoding RNAs (**Figure 7C**). Ingenuity linked this network mainly to cardiovascular system development and function, organismal development, and cellular assembly and organization. Moreover, *LZTS2* and *MYOM2* found in this network are among the top 10 genes with the highest fold difference in methylation status (**Figure 7C**).

For hypomethylated genes in active GO orbital fibroblasts, 6 of the top 10 genes with the highest fold difference were found in network 4, including *ERN2*, *CMTM8*, *BZW1*, *PRKCB*, *PSMD2* and *AP1M1* (**Figure 7D**). Of the genes that were found hypomethylated in orbital fibroblasts from active GO, *AIRE* and *SIM2* linked to network (network 4) that also contained NF-κB, ERK1/2 and Akt signaling pathways (**Figure 7D**). *HLA-A* and *PSMD2* (proteasome 26S subunit, non-ATPase 2), that are associated with peptide degradation, antigen presentation and TCR signaling processes, also linked to network 4 (**Figure 7D**). Ingenuity associated network 4 mainly with cellular growth and proliferation, nervous system development and function, and cell-to-cell signaling and interaction.

TABLE 3 | Top 5 canonical pathways and diseases and bio-functions from the hypermethylated genes with the cut-off at more than 2-fold difference in the active GO orbital fibroblasts.

| Top Canonical Pathways | | |
|--|---------|---------|
| Name | p-value | Overlap |
| Tyrosine Biosynthesis IV | 0.0145 | 33.3% |
| Phenylalanine Degradation I (Aerobic) | 0.0193 | 25.0% |
| Chondroitin and Dermatan Biosynthesis | 0.0289 | 16.7% |
| Glutamate Receptor Signaling | 0.0315 | 3.5% |
| NAD Biosynthesis from 2-amino-3- carboxymuconate Semialdehyde | 0.0336 | 14.3% |
| Top Diseases and Bio Functions Diseases and Disorders | | |

| Name | p-value | #Molecules |
|-------------------------------------|--------------------------------|------------|
| Cancer | 1x10 ⁻⁶ - 0.0145 | 95 |
| Gastrointestinal Disease | 1x10 ⁻⁶ - 0.0145 | 90 |
| Organismal Injury and Abnormalities | 1x10 ⁻⁶ - 0.0145 | 97 |
| Developmental Disorder | 2.22x10 ⁻⁵ - 0.0145 | 26 |
| Ophthalmic Disease | 2.22x10 ⁻⁵ - 0.0145 | 9 |
| Molecular and Cellular Functions | | |

| Name | p-value | #Molecules |
|-----------------------------------|--------------------------------|------------|
| Gene Expression | 7.0x10 ⁻⁵ – 0.00209 | 30 |
| Cellular Development | 7.0x10 ⁻⁵ - 0.0145 | 28 |
| Cellular Growth and Proliferation | 7.0x10 ⁻⁵ – 0.0145 | 25 |
| Cell Cycle | 4.85x10 ⁻⁴ - 0.0145 | 18 |
| Cell Morphology | 4.85x10 ⁻⁴ - 0.0145 | 14 |

Pathway and Functional Analysis From Methylated Genes in the Orbital Fibroblasts From GO Patients

The top 5 canonical pathways obtained from genes hypermethylated in orbital fibroblasts from active GO are shown in Table 3, while for genes hypomethylated in orbital fibroblasts form active GO the top 5 canonical pathways are given in Table 4. When all the functions from genes hypermethylated and hypomethylated in active GO (Tables 3 and 4) were integrated, overlapping bio-functions and diseases became apparent. The top disease/disorder identified to be shared between the hypo- and hypermethylated genes was organismal injury and abnormalities, containing ninety-seven hypermethylated genes (p-value = $1x10^{-6} - 0.0145$) and sixty-two hypomethylated genes (p-value = $9.88 \times 10^{-5} - 0.0498$). For the molecular and cellular functions, cellular development and cellular growth and proliferation were among the top molecular and cellular functions that were shared between the hypo- and hypermethylated genes. Cellular development linked to 28 hypermethylated genes (p-value = $7.02 \times 10^{-5} - 0.0145$) (Table 3) and nineteen hypomethylated genes (p-value = 0.00132 - 0.0469) (Table 4), while cellular growth and proliferation linked to twenty-five hypermethylated genes (pvalue = $7.02 \times 10^{-5} - 0.0145$) (**Table 3**) and seventeen hypomethylated genes (p-value = $1.32 \times 10^{-4} - 0.0498$) (**Table 4**).

TABLE 4 | Top 5 canonical pathways and diseases and bio-functions from the hypomethylated genes with the cut-off at more than 2-fold difference in the active GO orbital fibroblasts.

| • | Overlap |
|---------|-----------------------------|
| 0.003 | 100.0% |
| 0.00598 | 50.0% |
| 0.0132 | 1.8% |
| 0.0136 | 3.4% |
| 0.0154 | 3.2% |
| | |
| | 0.00598 0.0132 0.0136 |

| Name | p-value | #Molecules |
|-------------------------------------|--------------------------------|------------|
| Cancer | 9.88x10 ⁻⁵ - 0.0498 | 61 |
| Organismal Injury and Abnormalities | 9.88x10 ⁻⁵ - 0.0498 | 62 |
| Dermatological Diseases and | 4.73x10 ⁻⁴ - 0.0496 | 43 |
| Conditions | | |
| Gastrointestinal Disease | 0.00106 - 0.044 | 55 |
| Metabolic Disease | 0.00172 - 0.0498 | 8 |
| Molecular and Cellular Functions | | |

| Name | p-value | #Molecules | |
|--|--------------------------------|------------|--|
| Cellular Growth and Proliferation | 1.32×10 ⁻⁴ - 0.0498 | 17 | |
| Cellular Movement | 0.00117 - 0.0498 | 12 | |
| Cell-To-Cell Signaling and Interaction | 0.00132 - 0.0498 | 13 | |
| Cellular Development | 0.00132 - 0.0469 | 19 | |
| Drug Metabolism | 0.00132 - 0.0452 | 4 | |

DISCUSSION

Disease activity and characteristics of GO alter with disease progression. Early active GO is characterized by active inflammation that progresses into a late inactive phase without apparent inflammation but characterized by extensive orbital tissue remodeling (increased adipose tissue formation, excessive hyaluronan deposition and fibrosis). Orbital fibroblasts represent the main effector cells in GO and contribute to both the early active and late inactive phases of disease (8). Here, we performed extensive proteomics and global DNA methylation analysis of orbital fibroblasts from active and inactive GO as well as control individuals. Our data clearly demonstrate distinctive proteome and DNA methylation profiles between orbital fibroblasts from patients with active and inactive GO. These profiles can be linked to pathologically relevant molecular networks specifically associated with active/inactive disease stage, including inflammation, cellular development, growth, proliferation and tissue remodeling. Therefore, this data clearly illustrates that with GO disease progression orbital fibroblasts exhibit specific characteristics linked to disease stage specific pathology, and that epigenetic regulation is involved in controlling this.

Orbital fibroblasts from active GO displayed overexpression of proteins typically involved in inflammation, cellular proliferation, hyaluronan synthesis and adipogenesis (**Figure 1C**). In contrast, various proteins linked to ECM biology and fibrotic disease were overexpressed in orbital fibroblasts from inactive GO (**Figure 1B**). These findings corroborated with the different types of networks identified, which included cellular development and connective tissue disorders, DNA replication, recombination and repair and cellular assembly and organization (**Figure 2**). Although our current protein findings fit the pathophysiological model of GO, further studies to unravel the

exact function of these proteins in orbital fibroblasts as well as their contribution to the different stages of GO are required.

To our knowledge, our study is the first that explored DNA methylation in orbital fibroblasts from GO patients. Comparison of DNA methylation with stringent analysis using FDR between healthy controls and all the GO orbital fibroblasts included in our study resulted in a significant difference in methylation only for SLC39A8 (hypermethylation in 3'UTR region in case of the GO orbital fibroblasts; Figure 3). SLC39A8 is a member of the major zinc transporter SLC39 (ZIP) family, and cellular zinc import by SLC39A8 is crucial in the control of immune activation and function (52, 53). SLC39A8 methylation status did not correlate with mRNA expression, while SLC39A8 protein was not detected at all. However, the SLC93A8 hypermethylation in GO orbital fibroblasts located to the 3'UTR region, a region in which hypermethylation does not necessarily relate to transcriptional repression as commonly reported in promoter regions (54-56). We can, however, not exclude that SLC39A8 expression is also controlled by other regulatory mechanism such as small non-coding RNAs. SLC39A8 is also thought to be the primary transporter of the toxic cation cadmium, which is found in cigarette smoke. In this study we did not detect SLC93A8 protein, which might be related to the sensitivity of the technique we applied. Yet, we propose that additional studies to understand the role of the SLC39A8 and other SLC39 family of solute carriers in the pathogenesis of GO are potentially of great interest considering that smoking is a major environmental risk factor for GO (57).

Additional analysis of global DNA methylation using a 2-fold difference cut-off was conducted with the intend to lower the stringency from FDR analysis. A drawback of this approach is however that individual outliers have a relatively large effect on the final outcome, which either may mask potential important

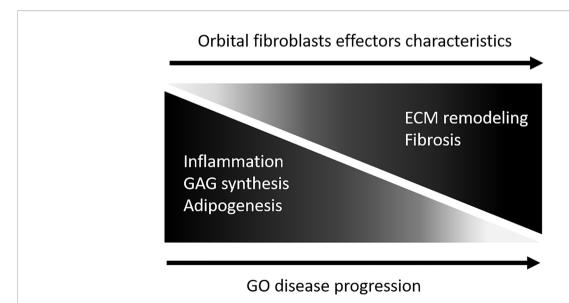


FIGURE 8 | Model depicting orbital fibroblast effector characteristics upon GO disease progression. During active disease orbital fibroblasts are skewed towards a functional phenotype involved in in inflammation, glycosaminoglycan synthesis and adipogenesis. With disease progression the orbital fibroblasts undergo a switch from this "inflammatory/pro-adipogenic primed" effector cell towards a "remodelling/pro-fibrotic" type of effector cell.

differences in methylation or identify false positive methylation differences, especially with the small groups (n = 4 per group) used in this study. Nonetheless, the overall DNA methylation patterns we observed suggest that orbital fibroblasts from inactive GO are more comparable to control orbital fibroblasts than to orbital fibroblasts from active GO (Figures 4 and 5). The global DNA methylation patterns and associated networks we found corroborate with the networks generated from the proteomics analysis, including NF-κB, Akt and IFNα (Figures 2 and 7), as well as previous reports that linked these pathways to GO pathogenesis (58-60). Other genes hypermethylated in inactive GO orbital fibroblasts also linked to other pathogenic pathways previously reported in GO, including PDGF (44), thyroid hormone receptor (61), NGF (62), AMPK (63) and ERK1/2 signaling pathways (64, 65) (Figure 7). In orbital fibroblasts from active GO hypermethylated genes clearly linked to inflammation, while hypomethylated genes linked to adipogenesis and autoimmune-related genes. This is in line with the association to inflammation and adipogenesis that we observed for the proteome of active GO orbital fibroblasts.

Genes hypomethylated in active GO orbital fibroblasts included AIRE and HLA-A (Figure 6B and Supplementary Figure 4). AIRE controls tissue specific antigen expression by medullary thymic epithelial cells to control thymic T-cell selection, and AIRE deficiency is associated with autoimmune disease (66, 67). CpG methylation in the promoter of the AIRE gene has previously been reported to control its tissue-specific expression pattern (68). Although hypomethylation of AIRE promoter was found in orbital fibroblasts from active GO (Supplementary Figure 4), AIRE expression was undetectable at mRNA and protein level (data not shown). Several studies showed AIRE expression in fibrocytes that typically infiltrate the orbital tissue form GO patients (69, 70). Potentially, fibrocyte numbers in our heterogeneous orbital fibroblast populations were low, which did however still allow detection of differences in AIRE methylation but not in AIRE mRNA or protein. HLA-A hypomethylation was detected in the gene body in orbital fibroblasts from active GO (Figure 6B). While HLA-A mRNA expression level did not differ between active and inactive GO, HLA-A protein expression was significantly down-regulated in active GO orbital fibroblasts (Figure 6B). Decreased HLA-A protein expression as observed in the orbital fibroblasts from active GO might regulate antigen presentation by these cells.

Fibroblasts isolated from fibrotic tissue display phenotypes that remain stable in *in vitro* culture for prolonged periods of time (71, 72). This allows investigation of epigenetic alterations and regulation, including histone modification, DNA methylation and non-coding RNA (71, 73). In general, we observed a poor correlation between protein expression, DNA methylation and mRNA expression. Poor correlations between DNA promotor methylation level and gene expression, as well as both positive and negative correlations between DNA gene body methylation and gene expression, have been reported (74–76). Our study thus clearly indicates that the proteome alterations we observed in orbital fibroblasts in relation to the specific GO disease stage are controlled by regulatory layers additional to DNA methylation (e.g.

histone modifications and small regulatory RNA molecules) that remain so-far hardly studied in GO. For example, several microRNAs, including miR-322, miR-508-3p, miR-9 and miR-16 have been described to target *NFKB1* mRNA (77–80). miR-16 was previously found to be upregulated in orbital tissue from inactive GO patients and might thus be involved in downregulating NFKB1 and resolving inflammation that occurs during this disease stage (81). Altogether, the current data support the role of NF- κ B signaling in regulating inflammation in GO orbital fibroblasts. Moreover, the data point towards existence of a complex epigenetic regulation machinery that controls NFKB1 expression and activity with GO disease progression.

There are several limitations associated with our study. First, the sample size is small, and it cannot be excluded that the extended orbital fibroblast isolation protocol and culture procedure altered the in vivo phenotypic/effector cell character of the orbital fibroblasts, although long term (epigenetic)stability of fibroblasts in culture is previously described (71, 72). Second, orbital fibroblasts are heterogeneous (8). Therefore, use of purified fibroblast subpopulations on the basis of cell surface marker expression, for instance Thy-1, CD34 and TSHR, could generate more in-depth insight into epigenetic regulation and fibroblast-subtype specific proteomes in GO. Despite these limitations, our proteomic and DNA methylation analysis identified known as well as novel molecules and molecular networks in relation to the pathogenesis of GO. In addition, both the proteomics and DNA methylation support that, with GO disease progression, orbital fibroblasts undergo a switch from an "inflammatory/pro-adipogenic primed" effector cell to a "remodeling/pro-fibrotic" type of effector cell, as depicted in Figure 8. We propose that detailed molecular understanding of this "inflammatory/pro-angiogenic-toremodeling/pro-fibrotic switch" should be an active field of future research as it can contribute to improvement of treatment protocols.

DATA AVAILABILITY STATEMENT

The mass spectrometry proteomics data have been deposited to the ProteomeXchange Consortium *via* the PRIDE partner repository with the dataset identifier PXD022257.

ETHICS STATEMENT

The studies involving human participants were reviewed and approved by protocol ID-2007-01. The patients/participants provided their written informed consent to participate in this study.

AUTHOR CONTRIBUTIONS

SV designed research, performed research, analyzed and interpreted data, wrote the paper. PS performed research, analyzed data, wrote the paper. TPi designed research, analyzed data, contributed analytic tools, prepared manuscript. PS analyzed data, contributed analytic tools. DP contributed patient materials. VD, PH, NH, and

TPa analyzed data, prepared manuscript. WD designed research, analyzed and interpreted data, prepared manuscript. All authors contributed to the article and approved the submitted version.

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SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at: https://www.frontiersin.org/articles/10.3389/fendo.2020. 619989/full#supplementary-material

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Supplementary Figure 1 | Comparison of GNAS DNA methylation and protein expression data.

Supplementary Figure 2 | Expression of *DNMT* and *TET* genes were determined by real-time quantitative (RQ)-PCR and normalized to the control gene *ABL*. Individual symbols represent orbital fibroblast cultures from individual patients. Horizontal bar depicts the median.

Supplementary Figure 3 | Methylation level of hypermethylated genes in orbital fibroblasts from active GO. Methylation level ranges from 0 (unmethylated) to 1 (fully methylated). Genes that were detected with the highest fold difference between active and inactive GO orbital fibroblasts, *WDR8*, and genes that were detected by more than one probe, *Mir548F5* and *MAB21L1*, are shown in this figure. Individual symbols represent orbital fibroblast cultures from individual patients. Horizontal bar depicts the median.

Supplementary Figure 4 | Methylation level of hypomethylated genes in orbital fibroblasts from active GO. Methylation level ranges from 0 (unmethylated) to 1 (fully methylated). Genes that were detected with the highest fold difference between active and inactive GO orbital fibroblasts, *RNF168*, and genes that were detected by more than one probe, *AIRE* and *SIM2*, are shown in this figure. Individual symbols represent orbital fibroblast cultures from individual patients. Horizontal bar depicts the median.

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Conflict of Interest: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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Association of Other Autoimmune Diseases With Thyroid Eye Disease

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Background: Thyroid eye disease (TED) is a potentially disfiguring and sight-threatening autoimmune (AI) orbitopathy, affecting up to 400,000 people in the UK. There are no accurate early predictors of TED severity. Although polyautoimmunity has been shown to affect AI disease severity, its influence on TED severity has never been investigated. The prevalence of polyautoimmunity among TED patients is also unclear, with discordant results reported in the literature. This study evaluates the prevalence of non-thyroid/ "other" AI (OAI) conditions in an ethnically diverse TED cohort and assesses how polyautoimmunity affects TED severity and activity.

Methods: A retrospective study of patients presenting to multidisciplinary TED clinics across three North-West London hospitals between 2011 and 2019. Data collected included: 1) demographics; 2) OAI conditions and management; 3) endocrine management of thyroid dysfunction; 4) details of TED and clinical activity score at presentation.

Results: Two hundred and sixty-seven patients with a median age of 46 (35–54) years were included, 79.4% were female and 55% were Black, Asian and minority ethnic (BAME). Thirty-seven patients (13.9%) had OAI conditions, with rheumatoid arthritis (3.7%), vitiligo (3.0%) and psoriasis (3.0%) among the most prevalent. Of patients with OAI conditions, 43.2% (16/37) required immunosuppression prior to TED onset. Non-immunosuppressed patients with OAI conditions had a significantly higher clinical activity score at presentation than TED-only and previously immunosuppressed patients (p=0.02). No significant differences were observed in thyroid receptor antibody titers between these groups.

Conclusions: This study finds a 13.9% prevalence of OAI conditions among TED patients. Patients with OAI conditions overall have a tendency for more severe and significantly more clinically active TED than those without OAI conditions. Larger, prospective studies are warranted to further evaluate polyautoimmunity as an early predictor of TED severity.

Keywords: thyroid eye disease, polyautoimmunity, disease severity, ethnically diverse, clinical activity score, immunosuppression, rheumatoid arthritis, psoriasis

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INTRODUCTION

Thyroid eye disease (TED), also known as Graves' ophthalmopathy/ orbitopathy, is a distressing, disfiguring, and potentially sight threatening, autoimmune disease (1). TED affects an estimated 400,000 people in the UK (2) and is the most common extrathyroidal manifestation of Graves' Disease (GD) (3–5). TED can cause a significant disease burden, causing pain, discomfort, double vision and considerable psychosocial distress to its sufferers (1, 6, 7). Around 5% of TED cases are estimated to progress to sight-threatening disease (8); however, there are currently no accurate early predictors of disease severity.

There is a well-established tendency for distinct autoimmune (AI) conditions to co-exist in patients who have previously been diagnosed with an AI condition, such as autoimmune thyroiditis (i.e. polyautoimmunity) (9–11). Although polyautoimmunity has been shown to affect AI disease severity (12–15), its association with TED severity has never been investigated.

There is also a paucity of medical literature investigating the prevalence of polyautoimmunity among TED patients. Only two studies have investigated the prevalence of non-thyroid AI, or "other AI" (OAI), conditions among TED patients. They estimate the prevalence of OAI conditions to be 8%–19% among TED patients (16, 17). However, both studies were carried out in monoethnic cohorts of Graves' hyperthyroidism patients only. Given that around 3% of TED patients have autoimmune hypothyroidism (Hashimoto's thyroiditis) and a further 2% never experience thyroid dysfunction (18), these results may not be applicable to all TED patients.

This study aims to evaluate the presence of non-thyroid AI conditions in an ethnically diverse, metropolitan TED cohort and to assess whether there is an association between the presence of polyautoimmunity and TED severity or clinical activity.

METHODOLOGY

This was a retrospective multi-center patient-cohort study based on patients attending three multidisciplinary thyroid eye disease (MDTED) clinics based in North West London, UK, at Imperial College Healthcare NHS Trust (ICHNT; Charing Cross Hospital CXH, Western Eye Hospital WEH) and Central Middlesex Hospital CMH, London North West Healthcare NHS Trust (LNWH). The audit was registered and approved by the relevant Trust departments.

Clinical records of all patients added to the joint MDT-TED clinic databases at CHX, WEH and CMH hospital sites from January 2011 to November 2019 were evaluated for eligibility. Patients were excluded if they had less than 3 months of follow-up, no TED diagnosis according to clinical/radiological criteria or incomplete clinic notes (i.e. no past medical history stated/no letters available for subsequent visits). Patients who had received ophthalmic or endocrine care outside Imperial or LNWH Trusts were also excluded as their hospital records were not accessible.

Data were collected from the electronic hospital records of eligible patients. Demographic data included: age at first

presentation at the MDT-TED clinic, gender, ethnicity, smoking status, and family history of thyroid disease (i.e. primary or secondary relative affected). Details of their endocrine care were collected to include: thyroid diagnosis (GD, Hashimoto's thyroiditis, Hashitoxicosis, hypothyroidism, euthyroid or other) at presentation, initial thyroid receptor antibody (TRAb) titer and thyroid dysfunction management [medication, thyroidectomy, and/or radioiodine (RAI) therapy].

Data collected on TED included: clinical activity score (CAS) at presentation; the patient's initial and most severe disease severity classification according to European Group on Graves' Orbitopathy (EUGOGO) criteria; management {selenium supplements, lubricating eyedrops, immunosuppression [first line treatment with intravenous methylprednisolone (IVMP) and/or second line immunosuppression, urgent or elective orbital decompression surgery, orbital radiotherapy (OR)]}.

Other autoimmune (OAI) conditions were only recorded for patients if the diagnosis had been confirmed by a specialist (for example, dermatologist, rheumatologist or gastroenterologist) following relevant clinical examinations and investigations. Steroid cover and immunomodulatory therapy for OAI were also recorded.

Statistical Analyses

All data were anonymized for statistical analyses using GraphPad Prism, version 8.0. The Shapiro-Wilks test was used to assess the normality of continuous data. Where data are not normally distributed, continuous variables have been expressed as medians (interquartile range). A Spearman rank correlation was used to evaluate the degree of association between two non-parametric variables (i.e. CAS score and TRAb titers). For groupwise analysis of non-parametric continuous data, a Kruskall-Wallis test was carried out, followed by Mann-Whitney U tests. Where more comparisons have been made than there are number of groups, alpha values have been adjusted using Bonferroni's correction to account for multiple comparisons. Chi squared tests were used to compare two categoric variables between groups.

RESULTS

Demographics

Two hundred and sixty-seven patients were included in the study. On average, patients were seen within 1.8 (0.8–2.9) months of referral from endocrine clinic to MDT-TED. Patients had a median age of 46 (35–54) years. Of these, 92.5% (247/267) had Graves hyperthyroidism, 3.7% (10/267) had Hashitoxicosis, 3.0% (8/267) had no thyroid hormonal dysfunction and 0.7% (2/267) had other thyroid pathologies, namely thyroid hormone resistance and follicular carcinoma of the thyroid. 79.4% were female and over half the participants (55%) were Black, Asian or minority ethnic (BAME). 13.9% (37/267) patients were noted to have OAI conditions, of whom 21.7% (8/37) had more than OAI condition. Nearly half [43% (16/37)] of patients with OAI conditions required immunosuppressive therapy prior to TED symptom onset. 42.6% (98/267) of patients were euthyroid at TED onset.

OAI Conditions

The most commonly observed AI condition was rheumatoid arthritis (RA) with the highest prevalence among Caucasian TED patients, (six compared to two BAME patients; **Table 1**). 3.0% (eight patients) had vitiligo and the same for psoriasis. Seven of the eight vitiligo patients were BAME, so this was the most prevalent OAI condition among BAME patients. 1.5% (4/267) had relapsing-remitting multiple sclerosis (RRMS). Overall, we did not detect statistically significant differences in the distribution of OAI conditions among Caucasian and BAME patients (p>0.05, post-hoc analysis not displayed).

In total, 6.0% patients (16/267) underwent immunosuppressive therapy for their OAI conditions before TED symptom onset with no statistically significant differences between the ethnic groups. On average, immunomodulatory therapy for OAI conditions was commenced 60 (2–120) months prior to TED symptom onset; only 18.8% (3/16) of patients were still receiving immunosuppressive therapy at the time of TED diagnosis. Treatment of TED included steroids, monoclonal antibodies and disease modifying drugs. 31.3% (5/16) of patients used prednisolone prior to TED symptom onset, 31.3% (5/16) had taken monoclonal antibodies, namely natalizumab, alemtuzumab or tocilizumab, and 50.0% (8/16) had commenced disease modifying therapies, such as methotrexate, prior to TED symptom onset.

Table 2 describes the characteristics of patients within three groups: those with TED only, those with TED + OAI conditions who had received immunosuppression for their AI conditions

prior to TED symptom onset and those with OAI who had not been immunosuppressed. Significant differences were observed in the median age of presentation to MDT-TED clinic between the three groups. The median age of presentation to MDT-TED clinic of previously immunosuppressed patients was 52 (46.3–57.8) years, which was significantly higher than that of patients with TED only, 45 (33.8–54.0) years (p = 0.04). No differences were observed in the distribution of sex, ethnicity, or smoking status in TED-only patients compared to patients with OAI conditions, regardless of prior immunosuppression.

A similar proportion of TED-only patients (49.6%) and patients with OAI conditions (43.2%) presented to MDT-TED clinic with hyperthyroidism, regardless of prior use of immunosuppression. The proportion of patients that underwent radioiodine therapy for thyroid dysfunction was also comparable across the three groups of patients.

Patients with OAI who had not received prior immunosuppressive therapy had the highest median CAS score at presentation (2, IQR: 1–4), compared to TED-only patients (p = 0.02; **Figure 1**) and previously immunosuppressed patients (p = 0.02; **Figure 1**).

Similar proportions of OAI (no prior immunosuppression) patients and OAI (prior immunosuppression) patients required first-line immunosuppressive therapy (IVMP) for TED (9.5% vs. 12.5%). However, only 6.3% of OAI (prior immunosuppression) patients required further (second-line) immunosuppressive TED therapy in the form of OR and/or MMF, while 28.6% of OAI (no prior immunosuppression) required first line (IVMP) and second line (OR and/or MMF) treatment.

TABLE 1 | Prevalence of non-thyroid autoimmune conditions in patients grouped by ethnicity.

| | | Ethnicity | | Total (n= 267) |
|---|--------------------|----------------|---------------------|----------------|
| | Caucasian (n = 74) | BAME (n = 147) | Not Stated (n = 46) | |
| Non-thyroid Autoimmune Conditions | | | | |
| Rheumatoid Arthritis | 6 (8.1%) | 2 (1.4%) | 2 (4.3%) | 10 (3.7%) |
| Vitiligo | 1 (1.3%) | 7 (4.8%) | 0 (0%) | 8 (3.0%) |
| Psoriasis | 2 (2.8%) | 3 (2.0%) | 3 (6.5%) | 8 (3.0%) |
| Relapsing-Remitting Multiple Sclerosis | 1 (1.3%) | 2 (1.4%) | 1 (2.2%) | 4 (1.5%) |
| Systemic Lupus Erythematosus | 0 (0%) | 2 (1.4%) | 0 (0%) | 2 (0.7%) |
| Crohn's Disease | 1 (1.3%) | 1 (0.7%) | 0 (0%) | 2 (0.7%) |
| Type 1 Diabetes Mellitus | 1 (1.3%) | 1 (0.7%) | 0 (0%) | 2 (0.7%) |
| Juvenile Idiopathic Arthritis | 0 (0%) | 1 (0.7%) | 0 (0%) | 1 (0.4%) |
| Alopecia Areata | 1 (1.3%) | 0 (0%) | 0 (0%) | 1 (0.4%) |
| Giant Cell Arteritis | 1 (1.3%) | 0 (0%) | 0 (0%) | 1 (0.4%) |
| Polymyalgia Rheumatica | 1 (1.3%) | 0 (0%) | 0 (0%) | 1 (0.4%) |
| Sjogren's Disease | 0 (0%) | 2 (1.4%) | 0 (0%) | 2 (0.7%) |
| Sarcoidosis | 1 (1.3%) | 0 (0%) | 0 (0%) | 1 (0.4%) |
| Al Pancreatitis | 0 (0%) | 1 (0.7%) | 0 (0%) | 1 (0.4%) |
| Al Neutropenia | 1 (1.3%) | 0 (0%) | 0 (0%) | 1 (0.4%) |
| Al Pernicious Anemia | 0 (0%) | 1 (0.7%) | 0 (0%) | 1 (0.4%) |
| pANCA Vasculitis | 0 (0%) | 0 (0%) | 1 (2.2%) | 1 (0.4%) |
| Coeliac | 0 (0%) | 0 (0%) | 1 (2.2%) | 1 (0.4%) |
| Urticarial Vasculitis | 0 (0%) | 0 (0%) | 1 (2.2%) | 1 (0.4%) |
| Myasthenia Gravis | 0 (0%) | 1 (0.7%) | 0 (0%) | 1 (0.4%) |
| Patients with >1 non-thyroid Al condition | 2 (2.8%) | 5 (3.4%) | 1 (2.2%) | 8 (3.0%) |
| Patients with one non-thyroid AI condition | 10 (13.6%) | 12 (8.2%) | 7 (15.2%) | 29 (10.9%) |
| Use of immunosuppressive therapy prior to TED symptom onset | 6 (8.1%) | 7 (4.8%) | 3 (6.5%) | 16 (6.0%) |

The presence of non-thyroid Al conditions was recorded if the condition had been diagnosed by a specialist following relevant examinations and investigations and reported in the patient's medical records. Data presented as "count (percentage of cohort)".

TABLE 2 | Summary of patient demographics, thyroid dysfunction and thyroid eye disease (TED) data categorized by the presence of non-thyroid autoimmune conditions and whether or not patients with non-thyroid autoimmune conditions had commenced immunosuppressive therapy prior to TED symptom onset.

| | | TED only (n = 230) | TED + OAI | conditions | p-value |
|--|---|---|---|---|---------|
| | | | No Previous Immunosuppression (n = 21) | Previous Immunosuppression (n = 16) | |
| Patient Demographics | | | | | |
| Sex | Female Male | 182 (79.1%) 48 (20.7%) | 16 (76.2%) 5 (23.8%) | 14 (87.5%) 2 (12.5%) | 0.78 |
| Age at first MDT-TED clinic | | 45 (34 - 54) | 48 (43.5-61.5) | 52 (44.8-57.8) | 0.04* |
| Smoking status | Current smoker Previous smoker Vapes Never Smoked | 54 (23.5%) 41 (17.8%) 4 (1.7%) 131 (57.0%) | 0 (0%) 3 (14.3%) 1 (4.8%) 17 (81.0%) | 2 (11.1%) 5 (27.8%) 1 (5.5%) 8 (50.0%) | 0.07 |
| Ethnicity | Caucasian BAME Not Stated | 62 (27.0%) 127 (55.2%) 41 (17.8%) | 6 (28.6%) 12 (57.1%) 3 (14.3%) | 6 (37.5%) 9 (56.3%) 1 (6.2%) | 0.75 |
| Thyroid Dysfunction | | | | | |
| Family history of thyroid dysfunction | | 61 (26.5%) | 3 (14.3%) | 5 (27.8%) | _ |
| Diagnosed Thyroid Pathology | Graves Hashimoto's None Other | 200 (87.0%) 7 (3.0%) 4 (1.7%) 2 (0.9%) | 17 (80.1%) 2 (9.5%) 2 (9.5%) 0 (0%) | 15 (93.8%) O (0%) 1 (6.2%) O (0%) | 0.23 |
| Thyroid status at TED onset | Hypothyroid Euthyroid Hyperthyroid | 18 (7.8%) 98 (42.6%) 114 (49.6%) | 4 (19.0%) 8(38.1%) 9 (42.9%) | 0 (0%) 9 (56.2%) 7 (43.8%) | 0.25 |
| Previous Radioiodine Therapy | ,, , | 32 (13.9%) | 4 (19.0%) | 1 (6.3%) | _ |
| Thyroid Eye Disease | | | | | |
| CAS at presentation | | 1 (1-2) | 2 (1-4) | 1 (1-2) | 0.04* |
| Selenium recommended | | 183 (79.6%) | 20 (95.2%) | 14 (87.5%) | _ |
| Proportion of patients requiring further therapy | IVMP only IVMP + MMF IVMP + Radio IVMP + MMF + Radio | 35 (15.2%) 25 (10.9%) 18 (7.8%) 9 (3.9%) | 2 (9.5%) 2 (9.5%) 1 (4.8%) 3 (14.3%) | 2 (12.5%) 0 (0%) 1 (6.3%) 0 (0%) | - |

Categoric data are presented as "count (percentage)"; all continuous variables are presented as "median (interquartile range)."

OAI, other autoimmune; TED, thyroid eye disease; BAME, black and minority ethnic; CAS, clinical activity score; IVMP, intravenous methylprednisolone; MMF, mycophenolate mofetil; Radio, orbital radiotherapy.

Thyroid Receptor Antibody (TRAb) Titer

61.4% (164/267) of patients were tested for TRAb antibodies and 88.4% of these (145/164) were TRAb positive. There were no significant differences in the TRAb titers of patients with TED only compared to those with OAI, regardless of prior immunosuppression (p=0.66; **Figure 2**). However, we note there was a tendency for OAI (no prior immunosuppression) patients conditions to have higher titers, 13.35 U/L (0.90–30.0), compared to OAI (prior immunosuppression) patients, 4.90 U/L (1.55–5.90), and those with TED only, 4.00 U/L (1.37–15.10; p>0.05).

Further sub-analysis found no correlation between TRAb titers and CAS of all included patients (r = 0.079, p = 0.32). There were also no correlations between TRAb and CAS within the three patient sub-groups.

We did not distinguish between stimulating and blocking TSH receptor antibodies as it was not possible to establish which assays had been used in the various laboratories across the three sites from 2011 to 2019.

DISCUSSION

This was the first study to evaluate the prevalence of non-thyroid OAI conditions in an ethnically diverse, metropolitan TED cohort and to assess potential associations between the presence of polyautoimmunity and TED severity or clinical activity.

The findings of this study, taken together with the results of the Cruz and Ferrari studies, suggest TED patients may have a higher prevalence of certain AI conditions compared to the general population. For example, RA and psoriasis was found to have a prevalence of 3.7% and 3% in our cohort, compared to a prevalence of 0.4%–1.2% and 1.3% in the general population (19). Our study also corroborates findings from the aforementioned studies that vitiligo and RA were among the most prevalent OAI conditions among TED patients. However, we also found a higher prevalence of psoriasis and MS in our cohort (16, 17).

A number of immune-related genes have been found in TED and OAI, presumably underpinning the inherited susceptibility to autoimmunity (20). There is evidence to suggest that

^{*}represents statistical significance, where p < 0.05.

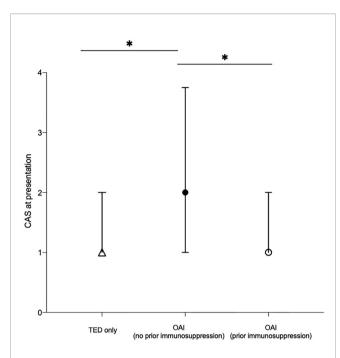


FIGURE 1 | Median clinical activity score (CAS) at presentation of patients categorized by the presence of non-thyroid autoimmune conditions and whether or not patients with non-thyroid autoimmune conditions had commenced immunosuppressive therapy prior to TED symptom onset. Median value of each cohort plotted. Error bars represent the interquartile range. *represents statistical significance, where p < 0.0167 by Bonferroni's correction. CAS, clinical activity score; OAI, other autoimmune; TED, thyroid eye disease.

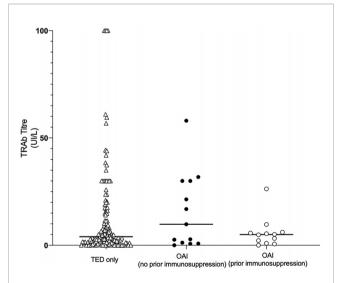


FIGURE 2 | Initial thyroid receptor antibody (TRAb) titer of patients categorized by the presence of non-thyroid autoimmune conditions and whether or not patients with non-thyroid autoimmune conditions had commenced immunosuppressive therapy prior to TED symptom onset. Data were collected from clinic notes and patient hospital records. Points represent the initial TRAb titer of each patient in the cohort; solid lines represent the median TRAb titer of each cohort. No significant differences were found in the TRAb titers of the three groups. OAI, other autoimmune; TED, thyroid eye disease.

mutations in the interleukin-23 receptor (IL23R) gene could confer increased susceptibility to AI conditions, such as RA, psoriasis and TED (21–23), which may explain the association of RA and psoriasis in our cohort. The link between IL23R gene mutations and TED pathophysiology is uncertain with only a few conflicting studies in the literature. For example, a study of Caucasian patients demonstrated the RA-associated single nucleotide polymorphisms of the IL23R gene to be associated with TED (24) while a similar study on Japanese patients did not identify such a trend (25). Further studies are needed to fully understand the role, if any, of the IL23R gene.

Vitiligo has previously been demonstrated to be associated with autoimmune thyroid diseases, such as Graves and Hashimoto's thyroiditis (26–28). Similarly, autoimmune thyroid diseases have been demonstrated to be the most common co-existing AI conditions in MS and RA patients (29). It would be interesting to compare the prevalence of each AI conditions in patients with AITD and no orbitopathy to patients with TED to establish whether there is an association between TED and these AI conditions, or rather if the presence of AITD in the studied population was a confounder. Further studies are required on the potential value of screening with thyroid antibodies in patients with OAI conditions that appear to be most prevalent in those eventually presenting with TED.

We found 13.9% of patients to have at least one non-thyroid OAI condition, of whom 48.6% had prior immunosuppression before their TED diagnosis. Patients in the TED only and OAI (no prior immunosuppression) subgroups presented with more active and severe disease compare to the subgroup with OAI (prior immunosuppression) and required more immunosuppression for their TED. It is possible this finding is a chance association and further studies are required to confirm this.

This study found a higher proportion of euthyroid patients at presentation to TED-MDT compared to figures previously published in the literature (30). The nature of the referral pathway to these TED-MDTs is such that the majority of patients are referred from an endocrine clinic where therapy such as radioiodine, which is known to have an adverse effect on TED (30), may have been used to establish a euthyroid state.

A limitation of this study is that the term "immunosuppression" included patients who had taken monoclonal antibodies, disease modifying drugs or steroids prior to TED symptom onset. It is therefore unclear if the different mechanisms of action of various immunosuppressive agents could have different effects on TED severity or activity. Although only one patient had taken alemtuzumab, a recent study has demonstrated this drug to trigger the onset of TED in "at risk" patients (31). Whether other monoclonal therapies can have a similar effect is unknown. Beyond this, the effects of alemtuzumab on TED severity remain unclear. It would therefore be interesting to further study the effects of steroids, monoclonal antibodies and other disease modifying drugs on TED severity and activity. A further limitation of this study reflects this being a single-center, retrospective study. While our findings are comparable to the ethnic diversity of London, UK (BAME population of 40.2% as per the Office of National Statistics 2011 Census), the findings may not be representative of other parts in the

UK (32). Furthermore, the study does not account for the effects of other medications, such as statins and metformin, among others, that could have an immunomodulatory properties (33).

Although the patients in this study were treated in the pre COVID-19 era, during the pandemic all first line intravenous steroid immunosuppression for non-sight threatening TED were stopped. We monitored and treated patients with severe relapse with oral steroids in line with best practice recommendations (34–36). All our second line immunosuppression patients continued treatment abiding with governmental shielding protocols. Careful consideration and constant evaluation has to be deployed regarding the choice and timing of immunosuppression in the post-COVID landscape.

In conclusion, the findings of this study were overall concordant with similar studies of monoethnic populations. Certain AI conditions, such as rheumatoid arthritis and vitiligo, appear to have an increased prevalence in TED patients. Whether or not this is due to a common pathway in their pathogeneses remains unclear. This study was also the first to demonstrate that patients with polyautoimmunity receiving immunosuppression prior to TED diagnosis may have less severe and less clinically active TED than those who have received no prior immunosuppressive therapy. Larger, prospective studies are warranted to evaluate how polyautoimmunity affects TED severity and activity and whether the presence of polyautoimmunity could be used as an early predictor of TED severity.

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DATA AVAILABILITY STATEMENT

The raw data supporting the conclusions of this article will be made available by the authors, without undue reservation.

ETHICS STATEMENT

The studies involving human participants were reviewed and approved by Imperial College Audit Office. Written informed consent for participation was not required for this study in accordance with the national legislation and the institutional requirements.

AUTHOR CONTRIBUTIONS

VL conceived and designed the study. MK acquired the data, analyzed, and interpreted the data. MK, PA, and VL drafted the manuscript. SF, RA, RJ, AA, CF, VB, and KM contributed to the manuscript. All authors contributed to the article and approved the submitted version. VL and MK are the guarantors of this work and, as such, had full access to all the data in the study and take responsibility for the integrity of the data and the accuracy of the data analysis.

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Mechanisms That Underly T Cell Immunity in Graves' Orbitopathy

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Fang S, Lu Y, Huang Y, Zhou H and Fan X (2021) Mechanisms That Underly T Cell Immunity in Graves' Orbitopathy. Front. Endocrinol. 12:648732. Graves' orbitopathy (GO), also known as thyroid-associated ophthalmopathy, is the most common ocular abnormality of Graves' disease. It is a disfiguring, invalidating, and potentially blinding orbital disease mediated by an interlocking and complicated immune network. Self-reactive T cells directly against thyroid-stimulating hormone receptor-bearing orbital fibroblasts contribute to autoimmune inflammation and tissue remodeling in GO orbital connective tissues. To date, T helper (Th) 1 (cytotoxic leaning) and Th2 (antibody leaning) cell subsets and an emerging role of Th17 (fibrotic leaning) cells have been implicated in GO pathogenesis. The potential feedback loops between orbital native residential CD34⁻ fibroblasts, CD34⁺ infiltrating fibrocytes, and effector T cells may affect the T cell subset bias and the skewed pattern of cytokine production in the orbit, thereby determining the outcomes of GO autoimmune reactions. Characterization of the T cell subsets that drive GO and the cytokines they express may significantly advance our understanding of orbital autoimmunity and the development of promising therapeutic strategies against pathological T cells.

Keywords: Graves' orbitopathy, thyroid-associated ophthalmology, T cell immunity, effector T cell, orbital fibroblast, fibrocyte

INTRODUCTION

Graves' orbitopathy (GO), also known as thyroid-associated ophthalmopathy, is the ocular abnormality of Graves' disease (GD). The prevalence of GO in Europe is about 10/10,000 people, which is above the threshold for rarity in Europe (1). However, as the most common extrathyroidal complication, GO affects 25-30% of patients with Graves' hyperthyroidism and detailed orbital imaging has revealed orbital soft tissue changes in 70% of GD patients (2, 3). Patients with GO suffer from impaired visual function, facial disfigurement, and at worst, irreversible visual loss caused by corneal ulceration or dysthyroid optic neuropathy, which result in a poor quality of life and socioeconomic status (4, 5). GO is a vexing autoimmune condition with both cellular and humoral immunities that form a sophisticated regulatory network, which leads to early orbital inflammation and late tissue remodeling (2, 4–6). Because of incomplete understanding of its precise pathogenesis, which partly results from the absence of suitable preclinical animal models, there is a lack of highly effective and well-tolerated therapies that target the most likely cause and glucocorticoids (GCs) are

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still the mainstay of treatment for active GO when inflammation is at peak (4, 5, 7, 8). Clinically, intravenous GC treatment has acceptable outcomes for most patients in the active phase. Nevertheless, a substantial number (20-30%) of active moderate-to-severe GO patients may not respond to GCs and adverse effects may occur after administration of high-dose or long-term GC use. Some patients may have disease progression despite GC treatment or relapse after steroid withdrawal (7, 8). Hence, a balance between benefits and risks of therapies for GO should be considered, which means developing more specific immunosuppressant strategies such as targeting T cells.

In the late 1980s, the role of T cell immunity was investigated in the orbital connective tissues of GO patients (9). Although thyroid-stimulating hormone receptor (TSHR) and its autoantibody play a major role in the pathological cascade of GO (2, 5), activation of humoral immunity, namely B cell immune responses, depends on defects in T cell immune modulation (10). The orbit is likely to have similar initial autoimmune reactions as those in the thyroid (5). It can be safely speculated that, among the various immune components that infiltrate the orbital connective tissues of GO patients, autoreactive T cells may act to establish and perpetuate the orbital inflammatory process. Recent studies have revealed that such disease-associated T cells include both T helper (Th) 1 (cytotoxic leaning) and Th2 (antibody leaning) subpopulations, and an emerging role of Th17 (fibrotic leaning) cells has also been implicated (6). The use of traditional non-specific immunosuppressants, such as cyclosporine that prevents interleukin (IL)-2 secretion by CD4+ T cells and mycophenolate that inhibits T cell proliferation by depleting guanosine-triphosphate, appear to be effective as a step-down from GCs to achieve stable efficacy in the long term (11). In view of the abovementioned facts, phenotypic and functional analyses of orbitinfiltrating T cells may provide better insights into the pathogenesis of GO.

In this review, we provide a detailed overview of the dysregulated T cell immunity in GO pathology. We include the early data as well as the latest research to reflect the developing course of understanding GO orbital autoimmunity. A selected listing of recommended studies on T cell pathogenesis in GO is summarized in **Table 1**. We highlight the integral role of pathological T cells that have deleterious effects on fibrocytes and orbital fibroblasts (OFs), and describe the development of targeted therapies for GO in an effective and safe manner.

CD4⁺ AND CD8⁺ T CELL IMMUNITIES IN GO

The first issue is whether cellular immunity is involved in GO inflammation. In an early study, Heufelder et al. reported the presence of CD3⁺ cells that represent total T cells in orbital and pretibial connective tissues from two GD patients with both orbitopathy and dermopathy (12). The results provide evidence of T cells infiltrating the inflamed orbit. Phenotypic analysis of four peripheral blood mononuclear cell (PBMC) samples from

four severe GO patients revealed the main subtype as CD4+ T cells (CD4/CD8 ratios 1.9-2.5), which was similar to the phenotypes of four control PBMC samples, whereas their corresponding orbital connective tissue-derived T cell lines had equal amounts of CD4⁺ and CD8⁺ T cells (CD4/CD8 ratios 0.9-1.2) (38). The ratios of CD4/CD8 were unchanged in 153 GO T cell clones cultivated from the four orbital T cell lines and 166 and 236 T cell clones cultivated from the four PBMC samples of GO patients and control subjects, respectively (38). The relatively low ratios of CD4/CD8 in orbital connective tissue-derived T cell lines and clones indicate that there is a disorder of cellular immune function in GO orbits. Grubeck-Loebenstein et al. established and characterized six T cell lines from orbital connective tissues of two severe GO patients and found they were predominantly CD8+CD45RO+ T cells (77%-96%) (39). The above two studies imply that a cytolytic T cell immunity triggered by CD8⁺ T cells may contribute to orbital inflammation in GO in a major histocompatibility complex (MHC) class I dependent manner. But the results cannot tell whether there exists a more efficient and unique antigen-presenting process to activate orbit-specific T cells. Stover et al. screened 64 orbital connective tissue-derived T cell clones expanded from two GO patients and reported an obvious predominance of the CD4⁺ T cell population (CD4/CD8 ratio 8.2) that contrasted with six PBMC samples (CD4/CD8 ratio 2.1) (39). In another study, the same research group analyzed 10 of 17 T cell lines derived from orbital connective tissues of six severe GO patients and found mainly CD4⁺ T cells (six of 10 strains) with a similar CD4⁺/CD8⁺ T cell distribution (40). The studies supporting the role of CD4⁺ T cells suggest an MHC class II pathway primed by a specialized antigenic determinant within the thyroid and at the involved orbital connective tissues. Pappa et al. investigated the extraocular muscles (EOMs) of 10 GO patients who underwent corrective strabismus surgery and examined six EOM-derived T cell lines from four patients. Five were CD4+CD45RO+ T cells (85%-97%) and CD8⁺ T cells (68%) were dominant in only one strain. The same status was found in the four corresponding PBMC samples (three were mostly CD4+ cells (89%-98%)) of each patient. They further reported detectable T cell receptor (TCR) gene expression in 10 out of 12 EOMs collected from the other five patients and in all five EOMs collected from three control subjects (41). The discrepancy of CD4⁺ and CD8⁺ T cell subsets in the above findings may lie in the small number of patients, the heterogeneity of patients involved in the different studies, and the different research methods. Notably, the T cell lines or clones in the above studies were cultured tissue- or peripheral blood-derived T cells expanded for several days to weeks, which may affect the initial status of these T cells to a certain extend. For example, CD8+ T cells may have more rapid expansion and CD4⁺ T cells gradually die during culture. Förster et al. established 18 T cell lines from orbital connective tissues of six severe GO patients and reported that 10 were predominantly the CD4⁺ phenotype, whereas three were mostly CD8⁺ cells (42). Intriguingly, in their study, even two independent T cell lines derived from the same patient had distinct T cell phenotypes (CD4 or CD8). This indicates that both CD4⁺ and CD8⁺ T cells

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TABLE 1 | Recommended Studies on T cell Pathogenesis in GO.

| Reference | Study subjects | Main findings |
|-----------------------------|---|--|
| T cell immunity and T | CR repertoires | |
| Heufelder et al. (12) | Biopsies of thyroid glands, orbital connective tissues, pretibial skins, and PBMCs from two GD patients | Both orbital connective tissues and pretibial connective tissues were infiltrated by CD3 ⁺ T cells; Marked similarities of intrathyroidal, orbital, and pretibial TCR gene |
| | with both orbitopathy and dermopathy and two non- GO controls | repertoires were found, which indicate apparent TCR restriction and T cell oligoclonality. |
| Pappa et al. (13) | Biopsies of EOMs from five early active GO patients, nine late stable GO patients, and 14 non-GO patients | CD4 ⁺ and CD8 ⁺ T cells and macrophages were significantly present in EOMs of active GO compared with both stable GO and controls; Increased HLA-DR expression on OFs, but not EOM fibres, was observed in both active and stable GO. |
| Rotondo Dottore et al. (14) | Biopsies of orbital connective tissues from 20 consecutive GO patients | A positive correlation was found between CD3 ⁺ T and CD20 ⁺ B cells infiltrating orbital connective tissues with GO clinical activity. |
| Wang et al. (15) | Biopsies of thyroid glands and PBMCs from six GD patients; PBMCs from 43 GO patients and 57 stable GD patients | A model for prediction of GO progression in GD cohort with high sensitivity and specificity. |
| Aniszewski et al. (16) | 117 CD4 ⁺ T cell clones expanded from orbital connective tissues of 6 GO patients | Th1 immune response predominated in early active GO and Th2 immune response predominated in late stable GO. |
| | d fibrocyte interaction | |
| Feldon et al. (17) | GO and control OFs; autologous T cells from PBMCs | Autologous T cells promoted the proliferation of GO OFs dependent on MHC class II and CD40-CD40L pathways. |
| Hwang et al. (18) | GO and control OFs | GO OFs expressed elevated levels of CD40 that could be further up-regulated by IFN- γ ; CD40-CD40L combination led to IL-6, IL-8, and MCP-1 production in GO OFs; CD90+ GO OFs expressed more CD40 than CD90- GO OFs. |
| van Steensel et al. (19) | Biopsies of orbital connective tissues from GO patients and controls; GO OFs | Mast cells, monocytes, and macrophages expressed increased levels of PDGF-A and PDGF-B in GO orbital connective tissues; PDGF-AB and PDGF-BB promoted proliferation and hyaluronan and IL-6 production by GO OFs. |
| Tsui et al. (20) | Biopsies of thyroid glands and orbital connective tissues; GO and control OFs; thyrocytes | TSHR levels were higher on thyrocytes than GO and control OFs; Differentiation of GO OFs, but not control OFs, into adipocytes led to increased TSHR expression; IGF-1R levels were higher on GO OFs than control OFs; TSHR and IGF-1R colocalized to the perinuclear and cytoplasmic areas of both GO OFs and thyrocytes. |
| Cao et al. (21) | GO and control OFs | CD40-CD40L combination led to the synthesis of hyaluronan and PGE ₂ in GO OFs; PGE ₂ production in GO OFs was caused by increased expression of PGSH-2 at both transcriptional and translational levels regulated by IL-1α expression |
| Koumas et al. (22) | GO OFs; myometrial fibroblasts | CD90 $^{+}$ myometrial fibroblasts and GO OFs were capable of myofibroblast differentiation by TGF- β or platelet concentrate supernatant treatment; CD90 $^{-}$ myometrial fibroblasts and GO OFs were capable of lipofibroblast differentiation by 15 deoxy- $\Delta^{12,14}$ -PGJ $_2$ or ciglitazone treatment. |
| Antonelli et al. (23) | Sera from consecutive subjects including 60 GD patients, 60 GO patients, and 60 controls; GO thyrocytes, OFs, and induced preadipocytes; Control fibroblasts and induced preadipocytes from dermal tissues of the same patients | CXCL10 was higher in GD and GO patients than controls; CXCL10 was significantly higher in active GO patients than inactive GO patients; IFN- γ and TNF- α synergistically induced CXCL10 production in GO thyrocytes, OFs, and preadipocytes, which was suppressed by PPAR- γ agonist. |
| Antonelli et al. (24) | GO thyrocytes, OFs, and induced preadipocytes; Control fibroblasts and induced preadipocytes from | IFN- γ and TNF- α synergistically induced CXCL9 and CXCL11 production in GO thyrocytes, OFs, and preadipocytes, which was suppressed by PPAR- γ agonist. |
| Han et al. (25) | dermal tissues of the same patients GO and control OFs | IFN- γ and IL-4 attenuated IL-1 β -provoked PGE $_2$ production by suppressing PGHS-2 gene promoter activity but enhanced IL-1 β -initiated hyaluronan production by up- |
| Han et al. (26) | GO and control OFs | regulating hyaluronan synthase-2 gene expression in GO OFs. IFN- γ and IL-4 attenuated IL-1 β -induced TIMP-1 production by suppressing TIMP-1 |
| Huber et al. (27) | Whole blood from 216 GD patients and 368 healthy | gene promoter activity in GO OFs. rs2201841 was strongly associated with GO development, especially AA and CC |
| Douglas et al. (28) | controls Biopsies of orbital connective tissues; PBMCs from 70 GD patients (including 51 GO patients) and 25 healthy controls; GO and control OFs; thyrocytes; fibrocytes | genotypes of II23r. CD34+CXCR4+Collagen I+TSHR+ fibrocytes were increased in PBMCs of GD patients. TSH induced fibrocytes to produce IL-6 and TNF- α ; Increased fibrocytes were found in orbital connective tissues of GO patients. |
| Gillespie et al. (29) | PBMCs from 31 GO patients and 19 healthy controls; GO OFs; GO and control fibrocytes | Fibrocytes expressed higher levels of TSHR than GO OFs; GO fibrocytes expressed higher levels of TSHR than control fibrocytes; TSH or M22 greatly stimulated the production of various cytokines and chemokines such as IL-8, RANTES, and MCP-1 i both GO and control fibrocytes. |
| Fang et al. (30) | Biopsies of orbital connective tissues; PBMCs from 34 GO patients and 36 healthy controls; GO and control OFs; <i>in vitro</i> -differentiated Th17 cells | GO peripheral Th17 cells produced IFN- γ and IL-22 and were related to clinical activity score; IL-17A enhanced TGF- β -induced fibrosis in CD90 ⁺ OFs but inhibited 15-deoxy $\Delta^{12,14}$ -PGJ ₂ -induced adipogenesis in CD90 ⁻ OFs; Th17 cells stimulated proinflammatory cytokine expression of GO OFs and GO OFs promoted Th17 cell differentiation by PGE ₂ production. |

(Continued)

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TABLE 1 | Continued

| Reference | Study subjects | Main findings |
|--------------------------|--|--|
| Fang et al. (31) | 21 GO orbital connective tissues and 38 control orbital connective tissues; CD34 ⁺ GO OFs; <i>in vitro</i> -differentiated Th17 cells | GO orbital microenvironment was composed of T cells, B cells, natural killer cells, dendritic cells, macrophages, plasma cells, and CD34 ⁺ OFs; Orbit-infiltrating Th17 cells displayed a Th1-like phenotype and expressed high levels of IL-1R and IL-23R; CD34 ⁺ OFs enhanced IL-1R and IL-23R expression on Th17 cells by PGE ₂ -EP2/EP4-cAMP signaling. |
| Fang et al. (32) | PBMCs from 16 active and 14 stable GO patients and 20 healthy controls; GO and control fibrocytes; <i>in vitro</i> -differentiated Th17 cells | IL-17A stimulated cytokine production in both GO and control fibrocytes; Autologous Th17 cells promoted inflammatory and antigen-presenting functions of GO fibrocytes; GO fibrocytes enhanced Th17 cell phenotype and recruited Th17 cells by MIP-3 and CCR6 combination. |
| Fang et al. (33) | Biopsies of orbital connective tissues; Sera and PBMCs from consecutive subjects including 37 GO patients, 38 GD patients, and 32 healthy controls | Increased CXCR3 ⁺ IFN-γ-producing Th17.1 cells were positively correlated with GO activity and associated with the development of very severe GO; In GC-resistant, very severe GO patients, CXCR3 ⁺ IFN-γ-producing Th17.1 cells remained at high levels in blood and orbital connective tissues, which were positively correlated with elevated triglycerides. |
| Fernando et al. (34) | GO OFs; GO and control fibrocytes | TSH and M22 induced IL-23, but not IL-12, expression in fibrocytes, while they induced IL-12 production in GO OFs; The shift from IL-23 expression in fibrocytes to that of IL-12 in CD34+ GO OFs was regulated by Slit2. |
| GO animal model | | |
| Moshkelgosha et al. (35) | hTSHR-A subunit plasmid-immunized BALB/c mice | TSHR was the pathogenic antigen in GO; Interstitial inflammation of extraocular muscles with CD3 ⁺ T cells, F4/80 ⁺ macrophages, and mast cells, accompanied by glycosaminoglycan deposition was observed in murine orbits. |
| Zhang et al. (36) | hTSHR-A subunit-expressing adenovirus-immunized BALB/c mice | Fibrosis and adipogenesis accompanied by CD4 ⁺ T cell infiltration were seen in murine periorbital fat tissues; Increased frequencies of Th1 cells and decreased frequencies of Th2 cells and regulatory T cells were shown in the splenocytes of GO mice. |
| Masetti et al. (37) | hTSHR-A subunit plasmid-immunized BALB/c mice | Bacteroides and Bfidobacterium counts were more abundant in mice in Center 1, while Lactobacillus counts were more abundant in mice in Center 2; Significantly higher yeast counts were found in Center 1 TSHR-immunized mice; A significant positive correlation was found between the presence of Firmicutes and orbital adipogenesis in Center 2 TSHR-immunized mice. |

are involved in GO pathogenesis. However, the phenotypic analysis was also based on T cell lines cultured *in vitro*. Therefore, direct *in vivo* T cell examination is needed to avoid biases and better reflect the real orbital immunity in GO inflammation.

Subsequently, an *in situ* study by immunohistochemistry demonstrated that both CD4⁺ and CD8⁺ T cells had infiltrated the EOMs in early active GO, which were much less evident in late inactive GO and control subjects (13). A recent study examined 26 GO patients and seven control subjects by immunohistochemistry, which showed that TCR expression was strong and diffuse in severe patients, although the orbital TCR detectable rate was similar in both active severe and inactive mild GO. Active severe GO patients had a higher CD3 detectable rate compared with inactive mild GO patients. Additionally, no expression of TCR or CD3 was found in control orbits (43). These data support the idea that GO orbital connective tissues are variably infiltrated by lymphocytes during active disease when medications are more effective than in the inactive disease.

We used flow cytometric analysis and found no differences in the frequency of circulating CD4⁺ and CD8⁺ T cells or the ratios of CD4/CD8 between GO patients and control subjects (44). In agreement with the above immunohistochemistry studies, infiltrated CD4⁺ and CD8⁺ T cells extended throughout the orbital connective tissues of GO patients, especially in the active phase, compared with control subjects (44, 45). Rotondo Dottore et al. confirmed that the total number of orbit-infiltrating T cells was correlated positively with the GO clinical activity score in

simple and multiple linear regression models (14). Studies in GO murine models also supported T cell-mediated inflammation in the orbit *in vivo*. CD3⁺ total T cells were found to infiltrate into the orbital muscles and periorbital tissues of human (h) TSHR-A subunit plasmid-immunized BALB/c mice (35, 46). The same phenomenon was found in mouse TSHR-A subunit plasmid-immunized BALB/c mice (47). Intriguingly, increased CD4⁺ T cell subsets were reported in periorbital fat of SKG mice after intraperitoneal administration of zymosan A compared with wild type mice (48). A recent study used an adenovirus that expressed the hTSHR-A subunit to induce GO in BALB/c mice and also observed CD4⁺ T cell infiltration in periorbital fat tissues (36). Collectively, these data shed light on the presence and type of T cells in GO, which suggest a complex inflammatory microenvironment in the orbit.

SELF-REACTIVE T CELLS DIRECTED AGAINST OFS

The second issue is whether T cells in GO recognize autoantigens, i.e., a primary GO immune response leads to the development of antigen-specific T cell responsiveness and clonal proliferation in the orbit. This will determine whether T cell immunity is specifically directed against orbital antigens. Heufelder et al. reported that in the two GD patients with both orbitopathy and dermopathy the vast majority of TCRs in the orbital and pretibial connective tissues were $\alpha\beta$ chains and not $\gamma\delta$

chians (12). Although expression of a broad spectrum of both TCR $V\alpha$ and $V\beta$ genes was observed in the PBMCs of patients, marked restriction of TCR Vα and Vβ gene expression was found in thyroid glands and orbital and pretibial connective tissues compared with PBMCs. Furthermore, thyroid, orbital, and pretibial tissues from two control subjects did not express restricted TCR transcripts (12). These data imply the potential GO-specific oligoclonal expression of the TCR gene repertoire. To further characterize the limited variability of antigen receptors on extrathyroidal T cells in GO, Heufelder et al. examined the TCR V gene repertoire in situ in orbital connective tissues and EOMs from eight early severe GO patients and observed apparent TCR $V\alpha$ and $V\beta$ gene restriction compared with matched PBMCs. Loss of TCR gene restriction was observed in four late GO patients and no TCR gene restriction was found in samples from three non-GO control subjects (49, 50). These findings suggest that oligoclonality of T cell immunity may be lost during GO, which indicates that antigen specificity of orbit-infiltrating T cells occurs in the early active phase of GO. This is important because an early adaptive immune response implies organ-specific autoimmunity in orbital connective tissues independent of the thyroid. Development of diversity or polyclonality of the TCR gene repertoire indicates that orbital inflammation is at the burnout stage. Heufelder summarized data from three severe active GO patients with GD and dermopathy and reported not only marked TCR restriction, but also several conserved junctional motifs shared by T cells in the orbit, thyroid, and pretibial tissue despite obvious heterogeneity of the TCR genes in each patient (12, 51). This highlights the presence of certain oligoclonal T cell populations stimulated by the analogous antigenic determinants shared by the thyroid and the involved extrathyroidal compartments. A recent interesting study proposed a novel TCR clonal expansion and chaos score to predict GO development in GD by characterizing complementarity determining region 3 of the TCR VB gene repertoire in PBMCs, which indicates specific GO TCR signatures distinctive from GD (15). These selected TCR-bearing T cells are self-reactive and recruited to the orbit at GO attack, which lead to orbital inflammation.

The next issue is unraveling the specific cell type or protein that triggers GO self-reactive T cell expansion. Genetic immunization with mouse TSHR-A subunit breaks selftolerance and induces GO-like pathology in BALB/c mice (47). Splenic T cells from BALB/c mice that have received hTSHR-A subunit prepared as a maltose-binding protein fusion induce orbital pathology in naïve recipient BALB/c mice marked by the presence of CD3⁺ total T cells (52). Furthermore, splenic T cells from hTSHR-A subunit plasmid-primed GO BALB/c mice show proliferative responses to purified TSHR antigen (53). These data from animal models provide a clue to potential TSHR-specific T cell responses that may also occur in the GO patient orbit. Arnold et al. reported occasional proliferation responses to EOM antigens in 10 circulating T cell lines from 10 severe GO patients. Additionally, these T cells hardly produced interferon (IFN)-γ under EOM antigen stimulation (54). Similarly, in the in vitro

model presented by Grubeck-Loebenstein et al., six T cell lines from orbital connective tissues did not proliferate in response to EOM antigen stimulation, but all had apparent proliferation after autologous OF treatment (39). In the in vitro model of Otto et al., the established 17 orbital T cell lines responded significantly to autologous orbital connective tissue proteins (6-10 and 19-26 kDa). A similar phenomenon was seen in most GO PBMCs that were more sensitive to autologous proteins from OFs than myoblasts. Moreover, orbital T cell lines hardly responded to allogeneic orbital proteins (40). Conversely, the authors demonstrated that 18 established T cell lines were barely able to respond to TSHR (2/18), thyroidal peroxidase (2/18) or thyroglobulin (none) (42). The results suggest the primary antigen role of TSHR and antigen-specific T cell clones in GO patients. However, the relatively low proliferation rate is confusing. It is important to note that although irradiated autologous PBMCs were added as feeders to help T cell to clone in these two studies, the antigen-induced T cell-specific proliferative response is acted in an antigen-presenting cell (APC)-dependent manner. The same research group used PBMCs from 16 GO patients and 12 controls and confirmed that incubation of GO PBMCs with OFs from the same patients led to marked T cell proliferation compared with control OFs. Similarly, compared with control OFs, GO OFs also had increased proliferation responses to stimulation by autologous PBMCs (55). This implies that OFs express GO autoantigens, and we hypothesize that GO OFs may function as facultative APCs to stimulate the proliferation of antigen-specific T cells, which has been confirmed by the fact that autologous T cells also stimulate the proliferation of GO OFs, but not eyelid-derived fibroblasts, via MHC class II and CD40-CD40 ligand (CD40L) signaling (17). We and other groups have shown that GO orbital connective tissues express higher gene and protein levels of MHC II and CD40 than control subjects (18, 30, 43, 56). Moreover, MHC II⁺ cells and CD40⁺ cells are local fibroblast-shaped cells and invading mononuclear cells such as macrophages in orbital connective tissues (18, 56). Even in stable GO, orbital connective tissues are activated to persistently express MHC II (56). Similarly, murine OFs derived from hTSHR-A subunit plasmid-primed BALB/c mice showed strong expression of CD40, TSHR, and insulin-like grow factor 1 receptor (IGF-1R) (57). Taken together, these findings have revealed sensitized and orbital connective tissue-specific T cells in circulation and the orbit of GO patients and provided evidence for self-reactive T cell populations directed against OFs.

PRIMARY IMMUNE REACTION IN THE ORBIT

The third issue is how T cells and OFs communicate with each other, which causes a series of pathophysiological changes in the GO process. Evidence for both T and B cells infiltrating GO orbital connective tissues was shown as early as 2012 (58). Many other *in situ* immunohistochemistry studies have demonstrated the presence of CD4⁺ T cells (44, 45, 56, 59), CD20⁺ B cells

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(14, 60, 61), CD14⁺ monocytes (19, 56), CD68⁺ macrophages (19, 59, 62), and CD117⁺ mast cells (19, 63) as the main invading immune components in the GO orbit. Using single cell sequencing analysis, we showed that various genes are expressed in GO orbital connective tissues, which can be classified into six independent cell types: APCs, lymphocytes including T and B cells, OFs, adipocytes, endothelial cells, and myocytes (31). This indicates an extremely complicated local orbital microenvironment, in which infiltrating immune cells and non-immune stromal cells interact with each other. The presence of APCs in the orbit further supports the idea that a primary GO autoimmune reaction occurs within the extrathyroidal compartment, although it resembles the process in the thyroid. OFs that express TSHR and IGF-1R (20, 31) detect danger signals to guide the property and intensity of the GO-adaptive immune response (64, 65). TSHR has been recognized as an autoantigen, but might not be the only one that activates GO self-reactive T cells. More work is needed to explain why GO occurs in patients with Hashimoto's thyroiditis with no evident TSHR autoimmune reactivity (2, 8). Although IGF-1R might be another autoantigen, GO pathology does not involve directly stimulating IGF-1R antibodies (5, 8). Serum antibodies against the IGF-1R are greater in GD patients, regardless of the presence of GO, suggesting a less pathogenic role of IGF-1R antibodies in GO (66, 67). In fact, a functional and physical crosstalk between TSHR and IGF-1R seems to be more important (5, 8). Defects in immune modulation lead to the breakdown of self-tolerance to thyroid or orbital connective tissues. Then, antigen-presenting cells (OFs themselves, fibrocytes, B cells, macrophages, or dendritic cells in the vicinity) recognize the exposed TSHR epitopes on the OF surface and then process and present the TSHR peptides to T cells, which leads to T cell clonal expansion and migration into orbital connective tissues (5, 6, 64). It should be noted that selfreactive B cells are likely to participate in GO antigen-presenting process. In recent onset GD patients, autoreactive B cells in PBMCs expressed CD86 and no longer appeared anergic, which indicates the activation and differentiation of B cells into plasma cells, leading to autoantibody production (68). Hence, it is reasonable to postulate that the same immune response occurs in orbital connective tissues in GO, where high levels of TSHR autoantibodies are detected in the sera of patients and may precede the onset of eye disease. A vital role of rituximab in the treatment of GO may lie in the blockade of antigen-presenting function of these activated self-reactive B cells (69). In addition, the over-reactive immune process has many other complicated mechanisms including thymic and peripheral T cell deletions and T cell anergy (5, 6, 70). Activated T cells provide the second signal for self-reactive B cell activation via the interaction of CD40L on the T cell surface with CD40 on the B cell surface. Moreover, the combination of B7 on the B cell surface and CD28 on the T cell surface provides the second signal for further activation of self-reactive T cells (5, 64, 71). Autoantibodies against TSHR are produced by plasma cells differentiated from activated B cells and autoantibody class switching (IgM to IgG and IgE) is aided by IL-4 secreted by activated T cells (mainly

Th2 cells) (5, 64, 71). Autoantibodies, including stimulating, neutralizing, and blocking IgG (72), target the TSHR on OFs, which may promote cytokine and chemoattractant production, deposition of extracellular matrix (ECM) such as hyaluronan, and pathological OF differentiation into adipocytes and myofibroblasts (73). Potential cross-talk of TSHR with IGF-1R on OFs helps to augment the expression of inflammatory molecules and hyaluronan synthesis (74, 75). The above pathological processes are largely due to the cell contact between OFs and T cells and cytokines produced by various T cell types (**Figure 1**).

An important intercellular communication in GO is CD40-CD40L signaling (Figure 2). CD40 is a mitogenic receptor that belongs to the tumor necrosis factor (TNF)-α receptor superfamily (76). CD40 is constitutively expressed by human fibroblasts derived from different tissue sources including OFs (18, 76), which facilitates fibroblast proliferation (76). GO OFs express elevated CD40 at gene and protein levels compared with control OFs (18, 77). When delineated by the cell surface marker CD90, CD90+ GO OFs had considerably greater CD40 expression than that on CD90 subsets as well as both control OF subsets (18). The combination of CD40 on OFs with CD40L on T cells leads to the three following pathological effects: (1) The release of inflammatory cytokines that induce acute and chronic orbital inflammation. Activation of GO OFs by CD40 engagement elevates IL-6 and IL-8 protein levels comparable with those produced by CD40-activated control OFs (77). Additionally, GO OFs primed with IFN-γ appear to be more responsive to CD40 activation than control OFs with regard to macrophage chemoattractant protein-1 (MCP-1) expression (18). Intriguingly, overproduction of IL-6 and IL-8 has been observed in CD90⁺ GO OFs compared with CD90⁻ GO OFs after priming with IFN-y (18). Conversely, CD40-CD40L signaling stimulates relatively low IL-6 and IL-8 production in both control OF subsets even when pre-incubated with IFN- γ (18). Hence, the higher expression of CD40 on CD90⁺ GO OFs may be critical to generate IL-6 and IL-8 in response to CD40L. Moreover, time-dependent secretion of prostaglandin (PG) E2 from GO OFs induced by CD40 engagement has been attributed to the up-regulation of IL-1 α production, which enhances the expression of prostaglandin endoperoxide H synthase-2 (PGSH-2 or COX-2) at both transcriptional and translational levels (21). (2) Up-regulation of adhesion molecules promotes immune cell recruitment to orbital connective tissues. GO orbital connective tissues expressed higher levels of intercellular adhesion molecule-1 (ICAM-1) and vascular cell adhesion molecule-1 (VCAM-1) compared with control subjects (30). The gene and protein expressions of ICAM-1 (78), VCAM-1, and E-selectin (79) were increased by CD40 activation in both GO and control OFs in dose- and time-dependent manners, with a more obvious effect on the former. Moreover, sphingolipids were induced in GO OFs, but not control OFs, which attracted T cells to migrate (80). (3) Synthesis of hyaluronan leads to the edema of orbital connective tissues and late-stage tissue remodeling. These phenomena are independent of PGE2 synthesis in GO OFs (21). Signal transductions for the pathways involved in the

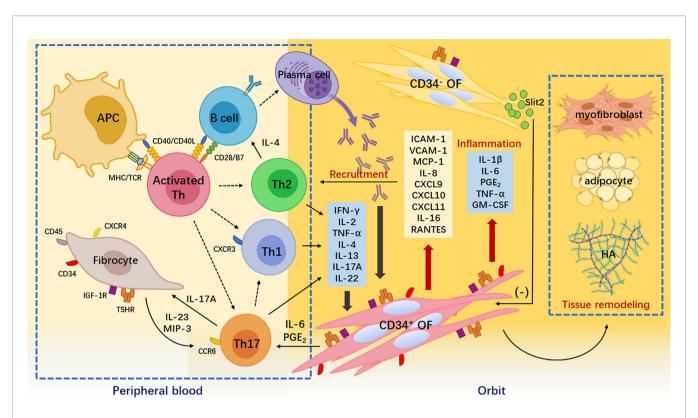


FIGURE 1 | Pathogenesis of Graves' orbitopathy. Breakdown of self-tolerance to thyroid-stimulating hormone receptor (TSHR) leads to the recognition of TSHR epitopes by antigen-presenting cells and B cells. They process the TSHR peptides to activate naïve T helper (Th) cells. Activated and expanded naïve Th cells differentiate into different subsets including interferon (IFN)-γ-producing Th1 cell, interleukin (IL)-4-producing Th2 cells, and IL-17A-producing Th17 cells. These cytokines together with autoantibodies produced by plasms cells derived from self-reactive B cells stimulate orbital fibroblasts (OFs), thereby initiating inflammatory responses in the orbit. IFN-γ is cytotoxic, IL-4 helps B cell expansion and autoantibody class switching, and IL-17A is proinflammatory and profibrotic. Meanwhile, peripheral CD34⁺ fibrocytes infiltrate orbital connective tissues and transition into CD34⁺ OFs. Upon IFN-γ and IL-17A stimulation, these CD34⁺ cells not only robustly produce chemokines such as C-X-C motif ligand 9/10/11 attracting C-X-C chemokine receptor 3⁺ Th1 cell and macrophage inflammatory protein-3 attracting C-C chemokine receptor 6⁺ Th17 cells but also secrete a large amount of cytokines such as IL-1β and prostaglandin E₂ that exacerbate orbital inflammation. CD34⁺ OFs ultimately synthesize hyaluronic acid and differentiate into adipocytes or myofibroblasts that cause orbital tissue remodeling. The orbital native residential CD34⁻ OFs express immunomodulatory molecules such as Slit2 to restrain the activities of the infiltrating CD34⁺ OFs.

CD40-CD40L combination mentioned above mainly include nuclear translocation of nuclear factor-κB, mitogen-activated protein kinases, and phosphatidylinositol-3-kinase. Further studies are needed to examine in more depth the cognate interactions between GO OFs and infiltrating T cells *via* CD40-CD40L communication.

Furthermore, T cell-OF interaction triggers the differentiation of OFs into adipocytes (8). Feldon et al. reported that GO orbital connective tissues and GO OFs highly expressed peroxisome proliferator-activated receptor (PPAR)- γ mRNA and proteins (81). In an autologous periphery T cell-OF coculture system, activated T cells drove the differentiation of GO OFs into orbital adipocytes. These activated T cells not only expressed upregulated levels of PGSH-2 but also synthesized PGD₂ and related PGJ₂ that are PPAR- γ ligands (81). This PPAR- γ -dependent adipogenic process of GO OFs provides evidence for how inflammation-provoking T cells regulate adipogenesis of orbital connective tissues and an interesting clue to the many faces of T cell immunity in GO.

PARADIGM OF TH1/TH2 IMMUNE RESPONSES

Previous studies have shown that both Th1 and Th2 cell subsets are involved in GO autoimmunity. In a study by De carli et al., 153 orbital T cell clones expanded from four severe GO patients exhibited remarkably high proportions of both CD4⁺ and CD8⁺ T cell phenotypes with a Th1-like cytokine profile including IFN-γ (82% in CD4; 88% in CD8), IL-2 (79% in CD4; 81% in CD8), and TNF-α (90% in CD4; 88% in CD8), but not IL-4 (4% in CD4; 0% in CD8) or IL-5 (1% in CD4; 0% in CD8), compared with T cell clones expanded from PBMCs of both GO patients and control subjects (38). Förster et al. examined cytokine gene expression in 18 orbital T cell lines from six severe GO patients and detected expression of Th1 cytokine genes Ifng (10/18), Tnfa (12/18), and Il2 (17/18) as well as Th2 cytokine genes Il4 (12/18) and Il5 (17/ 18). Other expressed cytokine genes were Il6 (13/18), Il10 (4/18), and Tnfb (15/18) (42). Using orbital T cell clones expanded from three severe GO patients, Yang et al. observed expression of Ifng

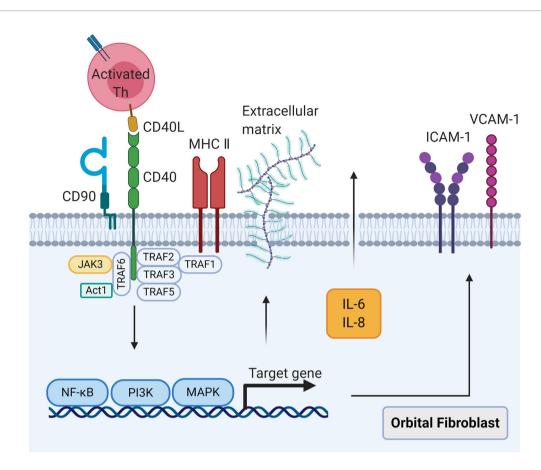


FIGURE 2 | The CD40-CD40 ligand signaling in orbital fibroblasts. When CD40 ligand on self-reactive T cells combines with CD40 on orbital fibroblasts, the nuclear translocation of nuclear factor-κB, mitogen activated protein kinases, and phosphatidylinositol-3-kinase signaling pathways will be activated, leading to the synthesis of cytokines interleukin-6 and interleukin-8, costimulatory molecules intercellular adhesion molecule-1 and vascular cell adhesion molecule-1, and extracellular matrix.

and Il2 in eight out of 14 CD4+ T cell clones and four out of six CD8⁺ T cell clones. The authors also assessed cytokine secretion in 38 CD4⁺ and 10 CD8⁺ strains, including the T cell clones for gene expression examination, and reported detectable levels of IFN-γ in most T cell clones (36/38 in CD4; 9/10 in CD8), of which some secreted IL-2 (8/36 in CD4; 5/10 in CD8). No Th2 cytokine gene profile and only three IL-4-secreting and five IL-10-secreting T cell clones were found (82). These results indicate that the great majority of orbit-infiltrating T cells express a Th1-like cytokine profile at both transcriptional and translational levels. In a study by Pappa et al., nine EOM-derived T cell lines from four GO patients were all positive for Th1 cytokine IFN-γ and IL-2. Other tested cytokines included TNF-β (5/9) and transforming growth factor (TGF)- β (9/9). They also found that Th2 cytokine IL-4 was positive in three out of five examined T cell lines (among the nine T cell lines) and IL-10 was positive in four out of five. However, the detectable rates of cytokines genes Il1a, Il2, Il4, Il6, Il8, Il10, Il5, and Tnfa varied among another 12 different EOMs of a further five patients. Expression of Ifng, Il13, Il1b, and Il12p40 was not detected in these EOMs. Il6 and Il8 were the only cytokine genes expressed in two out of five EOMs from three control subjects (41). It should be considered that gene and protein expressions are not complete coincident. Furthermore, apart from the technical problems related to the lymphocyte number and sample size, the various pre-surgery treatments that each patient had received and whether T cell clones were consecutively included or selected from independent patient cohorts will introduce biases and affect the results.

An important study by Aniszewski et al. examined cytokine production of 57 CD4⁺ T cell clones expanded from six GO patients and explicitly showed that T cell clones from recent onset GO (less than 2 years) mainly produced IFN- γ (47%) compared with IL-4 (23%), whereas those from remote onset GO (more than 2 years) dominantly produced IL-4 (75%) compared with IFN- γ (0%) (16). These findings suggest that the Th1 immune response may predominate in early active GO and the Th2 immune response is likely to play a role in late inactive GO. Unfortunately, all T cell strains examined were cultured and expanded in vitro for many days and only four T cell clones were matched for late GO, which drew criticism of the study design in which conclusions were made on the basis of data that may result in inaccurate estimates of T cell populations that invaded in situ within the orbit. Additionally, short duration GO is not exactly the same as active GO and longer duration GO does not mean that orbital inflammation is apparently absorbed.

By directly investigating the cytokine gene expression of EOMs in vivo from 14 GO patients and orbital connective tissues from 29 GO patients, expression of Ifng (13/14), $Tnf\alpha$ (5/14), Il1b (8/14), and Il6 (9/14) was mainly detected in EOMs, while they were less often expressed in orbital connective tissues in which Il4 (7/29) and Il0 (11/29) were more frequently expressed (83). Notably, the mean GO duration of the patients involved in the study was 2 years, which may account for the lower expression of *Ifng* and relatively higher expression of *Il4* in orbital connective tissues. Additionally, we cannot neglect the fact that all patients had undergone orbital irradiation before tissues were obtained and half had received high-dose GCs. Wakelkamp et al. investigated cytokine gene expression in orbital connective tissues from 17 GO patients, of whom six had untreated active disease and underwent emergency decompression surgeries. The other 11 patients were in the inactive phase and underwent rehabilitative surgeries. Compared with inactive GO patients, active GO patients had increased expression levels of Il1b, Il6, Il8, and Il10. Expression of Th1 cytokine genes *Ifng*, *Il2*, and *Il12p40* was also higher in active orbital connective tissues. However, expression of Th2 cytokine genes Il5 and Il13 was comparable in both patient groups and Il4 expression was not found in the study (84). These data imply the importance of the Th1, but not Th2, immune response and many other proinflammatory cytokines in the autoimmune

environment in the active GO orbit, although the gene expression results need to be supported at the protein level. Furthermore, the source of the examined cytokines cannot be identified in the bulk sequencing data.

Many recent studies have confirmed that orbital connective tissues from GO exhibit strong immunostaining for IFN-γ in the infiltrating cells, especially in the active phase (33, 45, 61). Using *in vivo* flow cytometric analysis, we demonstrated an increased frequency of CD3⁺CD8⁻ IFN-γ-producing T cells in both GO peripheral blood and orbital connective tissues compared with control subjects (31, 45). In hTSHR-A subunit plasmid-primed GO BALB/c mice, splenic T cells secreted IFN-γ (53). An increased frequency of CD4⁺ IFN-γ-producing T cells derived from splenocytes has also been observed in hTSHR-A subunit-expressing adenovirus-immunized GO BALB/c mice (36). In an SKG murine model of GO, expression of *Ifng*, *Tnfa*, and *Il2* was augmented in periorbital tissues and their protein levels were elevated in sera compared with wild type mice (48).

However, the pathological effects of IFN- γ on OFs are not fully understood (**Figure 3**). An essential function of IFN- γ in GO is acting as the "sinister partner" of CD40-CD40L signaling to exacerbate orbital inflammation. IFN- γ not only up-regulates CD40 on human fibroblasts derived from lung, skin, and gingiva, but also shifts fibroblasts to the G0/G1 phase of the cell cycle (76). Expression of CD40 was further augmented by IFN- γ in

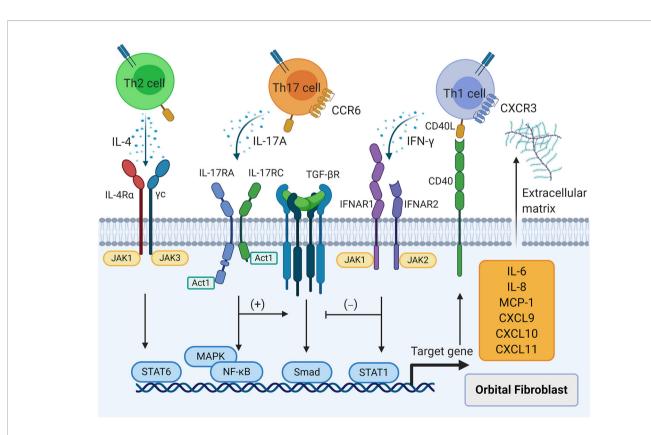


FIGURE 3 | The effects of T cell cytokines on orbital fibroblasts (OFs). T helper (Th) 1 cytokine interferon-γ, Th2 cytokine interleukin (IL)-4, and Th17 cytokine IL-17A result in the production of IL-6, IL-8, macrophage chemoattractant protein-1, C-X-C motif ligand 9/10/11, and extracellular matrix in OFs. Interferon-γ interferes with transforming growth factor-β induced fibrosis in OFs while IL-17A strengthens this process.

both GO and control OFs regardless of CD90 expression (18, 77, 85). Additionally, MHC II expression was increased in both GO and control OFs after IFN- γ stimulation (17, 77, 85) with a more potent effect on CD90° GO OFs (85). High-dose IFN- γ (1000 U/ml) alone was a potent stimulant of MCP-1 in GO and non-GO OFs as well as GO OF-differentiated adipocytes (86, 87) and induced IL-8 secretion with long term incubation (24 hours) (86). However, low- and medium-dose IFN- γ (100-500 U/ml) alone did not up-regulate IL-6 or IL-8 expression in GO and control OFs as well as their corresponding subsets (77, 85, 88), but greatly promoted IL-6 and IL-8 production provoked by CD40-CD40L signaling in mixed GO and control OF populations (77) as well as pure CD90⁺ and CD90° GO OF subsets (18, 85).

Another major pathological role of IFN-γ is establishment of a positive inflammatory feedback loop that maintains Th1 immune response in GO. The serum levels of chemokine C-X-C motif ligand (CXCL) 10 were higher in GO patients than control subjects, especially in active disease (23). Dose-dependent secretion of CXCL9, CXCL10, and CXCL11 after IFN-γ treatment has been observed in GO OFs as well as GO OFdifferentiated adipocytes (23, 24). Although TNF-α alone did not induce secretion of the chemokines in GO OFs and adipocytes, IFN-γ further promoted MCP-1, CXCL9, CXCL10, and CXCL11 release stimulated by TNF- α in these cells (23, 24, 87). The proinflammatory effect that IFN- γ with TNF- α synergistically exerts on GO OFs and adipocytes is suppressed in a dosedependent manner by PPAR-α agonists fenofibrate, gemfibrozil, or ciprofibrate, and PPAR-γ agonists rosiglitazone or pioglitazone (23, 24, 87, 89). A study regarding the role of circulating CXCL9 and CXCL10 as potential markers for GO activity revealed that GC treatment and teleradiotherapy significantly decreased CXCL9 and CXCL10 serum concentrations compared with basal values in GO patients. A positive correlation between CXCL9 and CXCL10 was also found in this study (90). Because C-X-C chemokine receptor (CXCR) 3 is particularly expressed on Th1 cells, which binds CXCL9, CXCL10, and CXCL11 (91), the above studies reflect an accurately self-modulated Th1 immunitymediated inflammatory network in GO.

Furthermore, IFN- γ results in the accumulation of ECM in GO. IFN-γ enhances hyaluronan synthesis activated by CD40-CD40L signaling in GO OFs and strengthens IL-1β-induced hyaluronan synthesis in GO OFs by promoting expression of the hyaluronan synthase-2 gene (21, 25). It does not directly induce PGE₂ secretion in GO OFs or contribute to PGE2 levels initiated by CD40-CD40L signaling (21). However, IFN-γ acts synergistically with CD40-CD40L signaling to elicit a dramatic increase in PGE₂ production in CD90⁺ GO OFs and CD90⁻ GO OFs via up-regulation of PGSH-2 proteins (85). Conversely, IFN-γ attenuates IL-1β-provoked PGE₂ production in GO OFs via down-regulation of PGHS-2 mediated by decreased Pghs-2 promoter activity and weakened PGHS-2 mRNA stability. This process is regulated by Janus kinase 2 signaling (25). The different modulation of PGE2 production by IFN-γ in combination with other molecular signals indicates a potential role of Th1 cell immunity and its related cytokines in regulating tissue reactivity and remodeling in the orbit. It is recognized that CD90+ OFs tend to differentiate into myofibroblasts, a hallmark of late GO fibrosis, whereas CD90 OFs tend to differentiate into adipocytes (2, 6, 22). IFN- γ blocks TGF- β -induced α -smooth muscle actin (SMA) expression in CD90⁺ GO OFs, which inhibits myofibroblast differentiation (22). Similarly, high levels of tissue inhibitor of metalloproteinase (TIMP)-1 gene and protein expression associated with fibrosis have been observed in IL-1 β -treated GO OFs in a dose- and time-dependent manner, which was attenuated by IFN- γ via down-regulation of *Timp1* promoter activity (26). This suggests that IFN- γ is more of a kind of proinflammatory factor that causes tissue damage and degeneration, and proves that the Th1 immune reaction is predominantly involved in early active GO.

The pathological effects of Th2 cytokines on OFs have yet to be examined carefully (Figure 3). Studies in GO murine models have not been able to duplicate Th2-dominated immune responses. A decreased frequency of CD4⁺ IL-4-producing splenic T cells has been observed in hTSHR-A subunitexpressing adenovirus-immunized GO BALB/c mice (36). However, compared with wild type mice, expression of Il4, Il5, and Il13 was increased in periorbital tissues of GO SKG mice (48). In another study, serum IL-4 remained at a higher level in hTSHR-A subunit plasmid-immunized GO BALB/c mice than in normal mice with extension of the immune time when IL-6, TNF-α, and granulocyte-macrophage colony stimulating factor were gradually declining (92). These results imply a possible role of Th2 cell-triggered immune responses in orbital connective tissues of stable GO. We used flow cytometry to confirm that the frequencies of CD3+CD8- IL-13-producing T cells and CD3+CD8-GATA3+ T cells were augmented in orbital connective tissues from GO patients. Both IL-13 and GATA3 were significantly related to GO development in a multivariate logistic regression model (31). These results suggest an indispensable and major role of Th2 immunity in GO inflammation. Although IL-4 cannot up-regulate CD40 expression in fibroblasts (76), it has many similar effects in regulating the biological behaviors of GO OFs. IL-4 suppresses *Timp1* promoter activation by IL-1β, which reduces the levels of TIMP-1 gene and protein expression in GO OFs (26). IL-4 also suppresses *Pghs-2* promoter activation by IL-1β, thereby inhibiting secretion of PGE2 from GO OFs (25). However, IL-4 promotes IL-1β-initiated hyaluronan synthesis in GO OFs by up-regulating hyaluronan synthase-2 gene expression (25). The identical functions of IFN-γ and IL-4 suggest transition from Th1 to Th2 cells to maintain the delicate balance between ECM production and degradation in orbital connective tissues as GO progresses from the early to late stage. In view of the major involvement of Th2 cell immunity in tissue fibrosis (93), more research on the relationship between Th2 cytokines IL-4, IL-5, and IL-13 and GO tissue remodeling is needed.

EMERGING ROLE OF THE TH17 IMMUNE RESPONSE

The first evidence regarding the possible role of Th17 cells in GO pathogenesis was published in 2008. A total of 216 GD patients

and 368 control subjects were genotyped for single nucleotide polymorphisms of Il23r. rs2201841 was strongly associated with GO, especially AA ($P=1.0\times10^{-4}$; OR=2.4) and CC ($P=1.4\times10^{-4}$; OR=2.36) genotypes (27). This indicates that *Il23r* variants may increase susceptibility to GO by regulating the expression or function of IL-23R on Th17 cells. Soon after, Kim et al. reported significantly higher detectable rates and serum levels of IL-17A in GO patients than those in control subjects, especially in the active phase (94). This was confirmed by another study in which serum IL-17A was higher in both active and inactive GO patients than in control subjects, despite its relative reduction compared with GD patients without eve disease (95). Additionally, Wei et al. observed the highest levels of serum IL-17A in active GO patients compared with those in both inactive GO and GD patients (96). Other studies that focused on lacrimal glands and the ocular surface have revealed elevated IL-17A levels in the tears of active and inactive GO patients (97-99). An orbital magnetic resonance scan showed that the axial lacrimal gland area was positively correlated with IL-17A concentrations in GO patient tears (99). Both serum and tear IL-17A levels have been positively correlated with the GO clinical activity score (94, 96, 99). We also observed up-regulated serum levels of IL-17A, but not IL-17F, in GO patients (44). More importantly, IL-23 (44, 94), IL-6 (44, 95, 97–99), and IL-1β (44, 97-99) concentrations were elevated in both sera and tears from active and inactive GO patients and more enriched in active phase, which are crucial factors for the differentiation of Th17 cells (100, 101). Analogously, the expression of IL-17A, IL-23, IL-6, and IL-1B increases diffusely around small vessels or fibroblasts and adipocytes within GO orbital connective tissues (44). These cytokines may construct a suitable microenvironment for the survival and activation of Th17 cells both systemically and locally in GO. We found that CD3⁺ IL-17A-producing T cells were increased among GO PBMCs compared with controls. Furthermore, both CD3⁺CD8⁻ (Th17) and CD3⁺CD8⁺ (Tc17) IL-17A-producing subsets are augmented in GO peripheral blood (44, 45). The CD3⁺CD8⁻ T cells in GO also express a higher proportion of retinoic acid receptor related orphan receptor (ROR)- γ t, the key transcription factor for Th17 cells (44). Intriguingly, most Th17 and Tc17 cells are CD45RO⁺ memory T cells (30, 44, 45), which indicates that these IL-17A-producing T cells might have been exposed to autoantigens such as TSHR and activated in the very early phase of GO or even in the GD stage. This is supported by the fact that the frequency of peripheral Th17 cells is higher in new-onset and intractable GD patients (102–104). More importantly, IL-17A-producing and RORyt-bearing Th17 cells were recruited at a higher fraction in GO orbital connective tissues, which were significantly associated with GO occurrence in a multivariate logistic regression model (31).

Th17 cells facilitate the inflammatory state of OFs in GO autoimmunity (**Figure 3**). In our *in vitro* model, IL-17A promoted transcriptional and translational expression of IL-6, IL-8, and MCP-1 in GO OFs in a dose- and time-dependent manner compared with fibroblasts derived from eyelid tissues. However, IL-17A did not affect the production of IL-23, IL-1 β , or TGF- β in GO OFs (44). IL-17A alone did not stimulate RANTES (regulated upon activation, normal T-cell expressed and

secreted) production, but strongly induced its mRNA and protein expression in the presence of CD40-CD40L signaling in both GO and control OFs in a dose- and time-dependent manner (45). In a Th17 cell-OF coculture system, Th17 cells promoted the secretion of IL-6, IL-8, MCP-1, macrophage inflammatory protein (MIP)-3, TNF-α, and granulocytemacrophage colony stimulating factor from both CD90+ and CD90⁻ OFs (30). In recent years, circulating fibrocytes have been recognized to participate in GO inflammation and tissue remodeling (105, 106). These cells express CD45, CD34, CXCR4, collagen I, thyroglobulin, TSHR, and IGF-1R, and were far more frequent in the circulation of GD and GO patients than in control subjects and were highly detected in GO orbital connective tissues, but were absent in control orbits (28, 29, 107). Both GO and control fibrocytes secreted TNF-α, IL-6, IL-8, IL-12, MCP-1, RANTES, MIP-1a, MIP-1b, CXCL10, and granulocyte colony-stimulating factor when stimulated by TSH or M22, a monoclonal TSHR-activating antibody (28, 29, 108). We found that GO and control fibrocytes synthesized IL-6, IL-8, and MCP-1 robustly in response to IL-17A, while GO fibrocytes had higher levels of basal and induced secretion of these cytokines than control fibrocytes (32). In a Th17 cellfibrocyte coculture system, we found that expression of Il6, Il8, Mcp1, Mip3a, Tnfa, Cxcl9, and Cxcl10 was augmented in GO fibrocytes and their proteins had accumulated in the culture supernatants (32). Both fibrocytes and OFs as well as OF subsets delineated by CD90 express IL-17RA (30, 32, 44), which suggests consecutive stimulation by Th17 cells from peripheral circulation to local orbital connective tissues in GO.

Th17 cells also modulate the fibrosis and adipogenesis balance in GO OFs. IL-17A directly leads to various ECM depositions in orbital connective tissues. Compared with control OFs, the gene and protein synthesis of fibronectin, collagen I, collagen III, TIMP-1, TIMP-2, matrix metalloproteinase (MMP)-1, and MMP-2 was greatly induced by IL-17A treatment of GO OFs in a dose- and time-dependent manner (44). Up-regulation of α-SMA gene and protein expression has been observed in IL-17A-treated GO OFs, which demonstrates differentiation of OFs into myofibroblasts (44). Unexpectedly, when we used pure CD90⁺ and CD90⁻ GO OF subsets, IL-17A exerted distinct effects on the two cell types. Lowdose IL-17A (10 ng/ml) was sufficient to enhance the fibrotic process marked by increased protein levels of α-SMA, fibronectin, collagen I, TIMP-1, and MMP-2 in GO OFs provoked by TGF-β. However, both low- and high (100 ng/ml)-dose IL-17A interfered with adipogenic differentiation of CD90 OFs induced by 15deoxy- $\Delta^{\hat{12},14}$ -PGJ₂. The protein levels of perilipin A, adipocyte differentiation-related protein, adiponectin, and PPAR-y were down-regulated in CD90 OFs in the presence of IL-17A (30). IL-17A promoted phosphorylation of JNK/c-Jun in CD90⁺ OFs, but impeded phosphorylation of CEBP/ α in CD90 OFs. Additionally, in CD90⁺ OFs, proteomics analysis has revealed that IL-17A enhances the production of ECM and proteins that are positive regulators for TGF-β and JNK cascade, but prevents adipocyte differentiation of CD90 OFs by up-regulating proteins involved in fatty acid oxidation, degradation, and efflux processes (30). Owing to the considerably high proportion of the CD90⁺

phenotype among GO OFs (30, 109), these findings suggest that GO OFs have a repertoire of differentiation that is more skewed towards myofibroblasts under IL-17A stimulation.

However, GO OFs regulate the phenotype and function of Th17 cells. In a Th17 cell-OF coculture system, both CD90⁺ and CD90 GO OFs enhanced the secretion of IL-17A from Th17 cells. Other supernatant-enriched cytokines included IL-22 and IL-21. An increased frequency of IL-17A+RORyt+ Th17 cells was shown by flow cytometry in the coculture system, which was repressed by down-regulating PGE2 released from CD90+ and CD90 GO OFs (30). The molecular mechanisms were possibly mediated by up-regulating IL-23R and IL-1R expression on Th17 cells, which was caused by PGE2-EP2/EP4 signaling that led to intracellular cAMP formation and subsequent phosphorylation of cAMP-responsive element-binding protein (31). These in vitro findings are consistent with the observation that GO orbital connective tissues contain a level of PGE2 and orbit-infiltrating Th17 cells express more IL-23R and IL-1R (31). Moreover, the Th17 cell-OF interaction results in a dramatic elevation of the expression of CD40, MHC II, ICAM-1, and VCAM-1 on CD90⁺ and CD90⁻ GO OFs, particularly on those that are also CD34⁺ (30). Such CD34⁺ OFs may originate putatively from CD34⁺ fibrocyte progenitors (106). Flow cytometric analysis has shown that CD34⁺ GO OFs have higher levels of IL-17RA than native residential CD34 subsets, which might account for the overexpressed CD40 and MHC II on CD34+ cells (31). Moreover, Th17 cell-fibrocyte interplay not only enhances IL-17A production in Th17 cells, but also significantly promotes CD40 and MHC II expression on GO fibrocytes (32).

How are Th17 cells recruited into orbital connective tissues in GO? Both peripheral and orbit-infiltrating Th17 cells express C-C chemokine receptor (CCR) 6, a MIP-3 receptor (30–32). Therefore, the MIP-3 released by GO fibrocytes might be a strong attractant that directs Th17 cells to sites of inflamed orbital connective tissues. Guo et al. demonstrated that orbit-infiltrating T cells in GO express CD44 (110), a specific cell surface receptor for hyaluronan (111). CD44 is highly elevated on activated T cells (112, 113) and particularly on CCR6⁺ IL-17A-producing Th17 cells in our study (30). However, T cell subsets with low expression of CD44 hardly secrete IL-17A in GO patients (30). Thus, with increased pericellular hyaluronan deposition, CD44 may facilitate Th17 cell attachment to GO OFs.

In recent years, the concept of Th17 cell plasticity has become prominent. Th17 cells acquire much more complex functional phenotypes than previously thought. Although they can shift phenotype within their lineage, Th17 cells have a dynamic ability to trans-differentiate into other CD4⁺ T cell subsets such as Th1 and Th2 cells (100, 114, 115). IFN-γ- and IL-22-producing Th17 cells are detected at significantly higher levels among GO PBMCs, especially in active patients (30, 45). These so called pathogenic Th17 cells express both RORγt and Tbet. They infiltrate into GO orbital connective tissues and more likely produce IFN-γ instead of IL-17A (31). TSH and M22 robustly induced gene and protein expression of IL-23 in GO fibrocytes, but not IL-12, which was significantly produced by GO OFs under the same conditions (34). However, pure CD34⁺ OFs

preferentially expressed Il23p19, while their homologous CD34⁻ OFs greatly expressed Il12p35 (34). The distinct roles of CD34⁺ and CD34 OFs reflect the potential shift from a non-pathogenic to pathogenic state of circulating Th17 cells into orbit-infiltrating Th17 cells, which is consistent with the TSHR signaling that drives the specific cytokine milieu by CD34⁺ fibrocytes that masquerade as CD34⁺ OFs within orbital connective tissues. The expression of IL-23 by CD34⁺ fibrocyte/OF lineages might play a prominent part in reinforcing the highly IL-23R-bearing Th17 phenotype in GO orbits (31) by endowing Th17 cells with "pathogenic" effector functions. We recently reported an increase in peripheral classic CD3⁺CD8⁻CXCR3⁺CCR6⁻ Th1 cells in active moderate-to-severe GO patients and GD patients, which were decreased in active very severe GO patients. Conversely, we found that classic CD3+CD8-CXCR3-CCR6+ Th17 and nonclassic CD3+CD8-CXCR3+CCR6+ Th17.1 cells were elevated among PBMCs from active very severe GO patients compared with both active moderate-to-severe GO and GD patients. Intriguingly, the non-classic Th17.1 cells favored IFN-γ production in active very severe GO patients, but preferentially secreted IL-17A in active moderate-to-severe GO patients. Moreover, the peripheral Th17.1 cells expressed higher levels of RORyt in active moderate-to-severe GO patients, whereas they had augmented levels of Tbet in active very severe GO patients, which was in concert with the different cytokine production phenotypes of these two patient cohorts. Very severe GO patients who did not respond to intravenous GC treatment had a sustained higher frequency of circulating and orbit-infiltrating Th17.1 cells (33). Therefore, we speculate an immunological transition process from Th1 cell immunity to Th17 cell immunity may indicate the development of very severe eye disease in GD. The overactivity of Th17.1 cells may serve as a hallmark for the not yet subsided inflammatory storm in orbital connective tissues. Evidence from animal models is indicating that IL-17A and IFN-y double-producing Th17 cells are pathogenic drivers of various human autoimmune diseases such as multiple sclerosis, diabetes type 1, uveitis, dry eye, rheumatoid arthritis, and inflammatory bowel disease (100, 114). Unfortunately, no convincing evidence of detectable Th17 cells has been observed in current GO murine models (36, 53), which makes it difficult to prove our hypothesis. The distinctive genetic backgrounds of BALB/c and C57BL/6J mice may partially be responsible for their susceptibility to GD and GO as well as the different T cell responses under autoimmune disease conditions (116). In this regard, a role of the gut microbiota that influence the immunological responses of induced GO murine models cannot be neglected (37, 117). For example, the YCH46 strain of Bacteroides fragilis reduces Th17 cell numbers by releasing propionic acid in GD patients (118). An interesting study reported correlations between murine GO manifestation and gut microbial taxonomies. Significant differences in the diversity and spatial organization of the gut microbiota of hTSHR-A subunit plasmid-immunized BALB/c mice were shown in two centers from different countries (37). Thus, the impact of different regions is also a source of potentially conflicting results, since the microbiome changes

across different countries. Disease-associated gut microbiota may contribute to the induced immune responses in GO murine models. Despite the confounding deviation from real human GO, future animal models will certainly be developed from existing experience and provide researchers with novel points of study to investigate the immunopathogenesis of GO.

FUTURE PERSPECTIVES

To date, immunomodulation therapy has been widely used for treatment of GO. Traditional non-specific immunosuppressants are effective in combination with GC treatment as alternative options for active moderate-to-severe GO (8, 11). Azathioprine and methotrexate interfere with purine synthesis that is necessary for lymphocyte proliferation. Mycophenolate, which inhibits inosine monophosphate dehydrogenase, and cyclosporine, which prevents IL-2 secretion, also exert antiproliferative effect on lymphocytes (8, 11). However, none of these therapeutic approaches appear to alter the natural course of GO, which makes development of more specific drugs critical to address an important unmet medical need. Considering the complexity of GO pathogenesis, there remain many ambiguous aspects of the pathological T cell activities within orbital connective tissues. For example, T cell migration and activation induced by autoantigens, autoantibodies, and immunomodulatory proteins. Activating TSHR on thymocytes enhances thymic output and therefore the functional T cell repertoire in the periphery (119). A larger proportion of peripheral CD3⁺CD45RO⁺IGF-1R⁺ T cells is seen in GO patients compared with control subjects. IGF-1R, which increases upon TCR stimulation, not only inhibits Fasmediated apoptosis, but also supports the expansion of memory T cells in GO (120). Furthermore, the proportion of peripheral IGF-1R⁺ T cells declines with clinical improvement in GO patients after rituximab treatment (121). Autoantibodies from GO patients up-regulate T cell chemoattractant IL-16 and RANTES from GO OFs (122). Moreover, T cell immunoglobulin domain and mucin domain 3, which restrains cytokine production in effector T cells except Th2 cells, is downregulated in peripheral Th1 and Th17 cells in GO patients (123, 124). Slit2 from residential CD34 OFs might inhibit production of IL-6 from GO CD34⁺ OFs, thereby ameliorating orbital inflammation and repressing Th17 cell differentiation (125). These findings offer new insights to explore novel approaches for therapy of GO. Existing evidence for the efficacy and relative safety of rituximab against CD20⁺ B cells, tocilizumab against IL-6, etanercept, infliximab, and adalimumab against TNF- α is encouraging (7, 71, 126). The impressive results of teprotumumab have provided the unprecedented possibility for monoclonal antibodies in combination with GCs for GO therapy, although more evidence must be provided. Trials of utilizing belimumab against BAFF (EUDRACT 2015-002127-26), K1-70 against TSHR (NCT02904330), and iscalimab against CD40 (NCT02713256) are currently underway. Blocking the IL-23/

IL-17A axis as a therapeutic strategy for GO is also promising considering its effectiveness in other autoimmune diseases such as psoriasis and mandatory spondylitis (71). Notably, a recent interesting study had the first attempt of antigen-specific immunotherapy with ATX-GD-59 that contains two TSHR peptides 9B-N and 5D-K1 in GD, which suggests that ATX-GD-59 is a safe and well-tolerated treatment (127). This antigen-specific method blocks the activation of APCs by binding with HLA-DR molecules, thereby inhibiting the subsequent cascade reactions of self-reactive T and B cells, which truly represents the needed breakthrough for targeted and effective therapy with less prone to general side effects. Novel biological agent identification on the basis of advances in GO pathogenesis is time-consuming but rewarding, which ultimately benefits patients with this debilitating disease.

AUTHOR CONTRIBUTIONS

SF and YH wrote the paper. YL constructed all the figures. HZ revised the paper. XF was responsible for the writing idea and framework of the paper. All authors contributed to the article and approved the submitted version.

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Cytokines as Targets of Novel Therapies for Graves' Ophthalmopathy

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Fallahi P, Ferrari SM, Elia G, Ragusa F, Paparo SR, Patrizio A, Camastra S, Miccoli M, Cavallini G, Benvenga S and Antonelli A (2021) Cytokines as Targets of Novel Therapies for Graves' Ophthalmopathy. Front. Endocrinol. 12:654473. Graves' disease (GD) is an organ-specific autoimmune disorder of the thyroid, which is characterized by circulating TSH-receptor (TSH-R) stimulating antibodies (TSAb), leading to hyperthyroidism. Graves' ophthalmopathy (GO) is one of GD extra-thyroidal manifestations associated with the presence of TSAb, and insulin-like growth factor-1 receptor (IGF-1R) autoantibodies, that interact with orbital fibroblasts. Cytokines are elevated in autoimmune (i.e., IL-18, IL-6) and non-autoimmune hyperthyroidism (i.e., TNFα, IL-8, IL-6), and this could be associated with the chronic effects of thyroid hormone increase. A prevalent Th1-immune response (not related to the hyperthyroidism per se, but to the autoimmune process) is reported in the immune-pathogenesis of GD and GO; Th1-chemokines (CXCL9, CXCL10, CXCL11) and the (C-X-C)R3 receptor are crucial in this process. In patients with active GO, corticosteroids, or intravenous immunoglobulins, decrease inflammation and orbital congestion, and are considered first-line therapies. The more deepened understanding of GO pathophysiology has led to different immunemodulant treatments. Cytokines, TSH-R, and IGF-1R (on the surface of B and T lymphocytes, and fibroblasts), and chemokines implicated in the autoimmune process, are possible targets of novel therapies. Drugs that target cytokines (etanercept, tocilizumab, infliximab, adalimumab) have been tested in GO, with encouraging results. The chimeric monoclonal antibody directed against CD20, RTX, reduces B lymphocytes, cytokines and the released autoantibodies. A multicenter, randomized, placebocontrolled, double-masked trial has investigated the human monoclonal blocking antibody directed against IGF-1R, teprotumumab, reporting its effectiveness in GO. In conclusion, large, controlled and randomized studies are needed to evaluate new possible targeted therapies for GO.

Keywords: corticosteroids, cytokines, Graves' ophthalmopathy, rituximab, teprotumumab, tocilizumab

INTRODUCTION

Graves' disease (GD) is an organ-specific autoimmune disorder and it is the most frequent cause of hyperthyroidism in West Countries (1–3). It affects overall women, typically in their third to fifth decade, with an overall prevalence of 0.5% (4).

Clinical manifestations are linked to hyperthyroidism and to the autoimmune process. GD-associated signs and symptoms can vary markedly, influencing the overall well-being (5, 6).

GD is one of the principal autoimmune thyroid disorders (AITD), which are associated with the failure of immune tolerance against thyroid antigens [thyroid-stimulating hormone (TSH) receptor (TSH-R), thyroid peroxidase (TPO), and thyroglobulin (Tg)] (7, 8). At the basis of GD, an autoimmune multifactorial mechanism is present, which acts through environmental and endogenous factors in genetically predisposed individuals (9). GD is characterized by thyrotoxicosis, circulating anti-thyroid antibodies (ATA) and autoreactive lymphocytes into the thyroid (10).

The autoimmune reaction in GD induces the release of anti-TSH-R autoantibodies (TRAb) by B-cell clones, which infiltrate the thyroid, and are implicated in GD pathogenesis and its extrathyroidal manifestations, such as pretibial myxedema (PTM)/Graves' dermopathy and Graves' ophthalmopathy (GO). TRAb can be distinguished in: neutral antibodies; thyroid blocking antibodies (TBAb); thyroid stimulating antibodies (TSAb) (11). TSAb have similar downstream effects such as the binding of TSH to TSH-R, leading to thyrocytes proliferation, and secretion of thyroid hormones (TH; T4 and T3) (11, 12).

GD is associated to another autoimmune disease in \sim 20% of patients (13). A study evaluated prospectively the prevalence of other autoimmune diseases in GD patients (including some GO patients) vs. healthy controls, or patients with autoimmune thyroiditis (AT), or with multinodular goiter (MNG) (genderand age-matched, and with a similar iodine intake). In 1.5% of GD, three associated autoimmune disorders were present. Patients with GO had a higher prevalence (18.9%) of another autoimmune disorder than GD patients without GO (15.6%) (13).

GRAVES' OPHTHALMOPATHY

According to the European Group on Graves' Orbitopathy (EUGOGO), GO has a prevalence of 10/10,000 persons and 16/10,000 persons in Europe and Japan, respectively (14, 15).

Approximately 30% to 50% of patients with GD develop GO. Tearing, proptosis, periorbital edema, and diplopia are the characteristic signs of Graves' orbitopathy (16).

Orbital fibroblasts (OF) are the target of a variety of autoimmune responses, that taking together induce proliferation, adipogenesis and overproduction of the extracellular matrix, that includes glycosaminoglycans [GAG; i.e., hyaluronan (HA) and chondroitin sulfate] (4).

GD is a phasic disorder characterized during the active phase by hyperthyroidism, which can go in remission both after therapy or sometimes spontaneously. The recurrence of hyperthyroidism can occur after weeks or decades of euthyroidism, overall owing to physical or psychological stressful events (2). Also, GO is characterized by the presence of moderate-severe inflammatory signs during the active phase that can remit after therapy or spontaneously. Mild GO disappears spontaneously in the majority of cases (17). The most frequent factors associated with the recurrence of GO are stressful life events (2), and smoking (18). Smoking cessation can improve the outcome for GO (3, 4). Components of cigarette smoke might induce adipogenesis and synthesis of GAG, through reactive oxygen species (ROS) production, and whether vaping could have some negative effects on GO is still to be clarified, as nicotine induces the release of proinflammatory cytokines (4, 19). As well as in GD, an autoimmune response is at the basis of GO. T cells, and autoantibodies recognizing a common autoantigen both for the thyroid and retro-ocular tissues (such as TSH-R, that is expressed also on fibroblasts and orbital preadipocytes), have a crucial role in this process (20). Moreover, another autoantigen, the insulin-like growth factor-1 (IGF-1) receptor (IGF-1R), has been suggested to be linked to GO (21). Autoantibodies against IGF-1R take part in the activation of GO OF, and its elevated expression has been shown in the thyroid and in orbital tissues in GD patients (20).

THERAPIES FOR GO

An anti-inflammatory treatment is suggested in the active phase of GO. A euthyroid state is mandatory for successful treatment and radioiodine therapy should be avoided in active progressive GO (15).

In patients with active GO, corticosteroids (CS), or high dose intravenous (iv) immunoglobulins, decrease inflammation and orbital congestion (22). A prospective, randomized trial compared the efficacy and safety of two protocols of iv 4.5 g methylprednisolone (MP) in 80 patients, randomized to receive iv MP weekly or daily. The weekly protocol of iv MP therapy was more efficient and safer than the daily protocol for patients with active moderate-to-severe GO (23).

According to randomized controlled trials, a meta-analysis reported in patients treated with iv glucocorticoids that a reduction of 1.14 mm of proptosis and of 33% of diplopia is present, while non-randomized studies reported a reduction of 1.58 mm of proptosis and of 25% of diplopia (24). After 6 weeks, in case of worsening of the disease the therapy should be supplemented or substituted with second-line treatments (15), such as orbital radiotherapy (25, 26). Orbital decompression should be done in severe GO (27).

Patients with severe GO can have a reduced quality of life (QoL) during standard therapies and, for this reason, novel treatments, targeting directly the pathogenic disease

mechanisms in GO, are necessary to improve the clinical outcome in these patients.

At present, the more deepened understanding of GO pathophysiology has led to alternative immune-modulant therapies that target various antigens (28, 29).

The aim of new treatments for thyroidal and extrathyroidal GD is firstly to target the principal autoantigens of the disease and/or molecules that have a key role in the immunological response. Future treatments of GD, and GO, will involve monoclonal antibodies (mAb) or small molecules (15).

A mechanism for targeting the pathophysiology of GD, and GO, is to "mask" the TSH-R from the action of thyroid-stimulating immunoglobulin (TSI), through the TSH-R antagonist, K1-70 (15, 30, 31). An open-label clinical trial (clinical trial number NCT02904330) is currently in progress to assess its safety and tolerability in patients with hyperthyroidism in GD (32).

A small TSH-R antagonist (clinical trial number NCGC00229600) has been shown to reduce the synthesis of HA in retro-ocular fibroblasts/adipocytes in GO, with good results (33).

Furthermore, an encouraging treatment in GO patients is teprotumumab (RV 001, R1507), a recombinant, human mAb of the immunoglobulin G1 subclass. It binds to the cysteine-rich domain of human IGF-1R with high affinity, preventing the binding with endogenous ligands (IGF-1 and IGF-2), and leading to the internalization of the receptor, in this way stopping the IGF-1R signaling cascade (34–37).

As written above, active GO is linked to the autoimmune activation of OF. This causes the production of cytokines that promote T-cell infiltration into orbital tissues, triggering a local inflammatory process, and resulting in growth and differentiation of fibroblasts, and remodeling of orbital tissues. Teprotumumab blocks these pathophysiological responses (36).

Teprotumumab decreased *in vitro* TSH-R and IGF-1R, the TSH- and IGF-1-dependent phosphorylated Akt levels in fibrocytes, such as the TSH induction of IL-6 and IL-8 mRNA and protein (38).

A randomized, double-masked study was performed, to evaluate the effectiveness and safety of this drug in 88 patients with active, moderate-to-severe GO, administered with placebo, or the drug (eight infusions) (39). The primary end point was the response in the study eye. This response was defined as a reduction of two points or more in the Clinical Activity Score (CAS) and a reduction of 2 mm or more in proptosis at week 24. About 69% of patients taking teprotumumab, in comparison to 20% of those receiving placebo, had a response at week 24 (P<0.001). At week 6, 43% of patients administered with teprotumumab and 4% with placebo had a response (P<0.001). The reported findings in active GO demonstrated that teprotumumab, compared to placebo, is more effective in decreasing proptosis and CAS (39).

Another randomized, double-masked, phase III multicenter trial was done in patients with active GO in a 1:1 ratio to receive iv teprotumumab (10 mg/kg of body weight for the 1st infusion and 20 mg/kg for the following ones) or placebo once every 3 weeks for 21 weeks, for a total of 24 weeks (40). At week 24, the

response in proptosis was higher in patients treated with teprotumumab than with placebo (83% vs. 10%, P<0.001). Secondary findings were significantly better with teprotumumab than with placebo (P \leq 0.001). The data demonstrated that teprotumumab led to better outcomes (than placebo) with regard to proptosis, CAS, diplopia, and QoL, with uncommon severe adverse events (40).

Teprotumumab attained the approval by the U.S. Food and Drug Administration (FDA) as first drug for thyroid eye disease in March 2020 (41, 42).

Another emerging therapy is rituximab (RTX), a chimeric mAb against CD20. It was approved by FDA in Wegener's granulomatosis, chronic lymphocytic leukemia, non-Hodgkin's lymphoma, and rheumatoid arthritis (RA), and it is an off-label drug in different autoimmune diseases (43). Since RTX reduces B lymphocytes, the cytokine burden and the produced autoantibodies, its use has been proposed also in GO.

Two randomized trials showed contrasting results regarding the use of RTX. In the 1st prospective, randomized, placebo-controlled trial, RTX was not effective in 25 GO patients, showing no improvement of CAS (44). In the other randomized, double-blinded study, 32 patients took iv MP or RTX, and CAS decreased in both cases, overall with RTX. After 24 weeks, all patients receiving RTX ameliorated with respect to 69% after ivMP, showing RTX was more effective than ivMP in GO patients (45).

More recently, a meta-analysis and systematic review was done in 293 GO patients, treated with RTX, or glucocorticoids, or saline, to evaluate the effectiveness of RTX. It was shown a significant decrease of CAS (vs. controls) in patients receiving RTX at 24 weeks, and a not significant one of proptosis (46).

In another study (47), 219 GO patients received pulse MP, and then oral steroids and/or orbital radiotherapy. At last, 15 (6.8%) were administered with 100 mg RTX (100–400 mg as cumulative dose) for the presence of active GO. A low dose of RTX had an anti-inflammatory effect in patients with active GO resistant to standard treatments (47). Another paper agreed with those results in 12 patients with active GO treated with a 100 mg RTX infusion; it reported that a low dose of RTX is effective in these patients, thus leading to a reduced administration of systemic steroid (48).

Furthermore, 14 patients with active and moderate-to-severe GO, of whom 11 CS-refractory, were administered with ivRTX (1000 mg twice with a 2-week interval), reporting that RTX was well-tolerated (49). A modest amelioration of CAS, and disease inactivation in half of the patients, were reported. At week 12, CAS ameliorated in 14.3% of patients and inactivation of GO in 28.6%. At week 24, total eye score and proptosis improved in 28.6% and 33% of patients, respectively.

CYTOKINES AND DRUGS TARGETING CYTOKINES, IN GD AND GO

Cytokines in GD and GO

Cytokines [including interleukins (IL), tumor necrosis factors (TNFs), interferons (IFNs), lymphokines, and chemokines] are

small proteins, important in normal physiology, and in host responses to infection, trauma, reproduction, inflammation, sepsis, and tumors (10).

They are produced by different cell types, including immune cells (B and T lymphocytes, mast cells, macrophages), fibroblasts, endothelial cells, and different stromal cells, and act through receptors (**Table 1**).

Chemokines are "chemotactic cytokines", or signaling proteins, which can induce directed chemotaxis in the responsive cells. Chemokines exert their biological effects through the interaction with G-protein-linked transmembrane receptors, which are present on their target cells (10). The chemokine receptor (C-X-C)R3 binds Th1-chemokines [IFN-yinducible protein 10 (IP-10)/chemokine ligand 10 (C-X-C motif) (CXCL)10, IFN-inducible T-cell α chemoattractant (I-TAC)/ CXCL11 and monokine induced by IFN-γ (MIG)/CXCL9] (60). Th1 lymphocytes are attracted in inflamed tissues by Th1 chemokines, that are released there (61), and increase cytokines production, leading to Th1 chemokines secretion by different cells, establishing an amplification feedback loop (62). Raised serum and tissue Th1 chemokines levels have been demonstrated in specific autoimmune disorders [autoimmune thyroiditis (AT) (63-65), GD and GO (52), etc.], or systemic rheumatological diseases [systemic sclerosis, psoriasis or psoriatic arthritis (66, 67), etc.], hepatitis C virus infection related autoimmune disorders (68), and cancer (69, 70).

Certain serum cytokines are elevated in non-autoimmune (IL-8, TNF- α , and IL-6) (71) and autoimmune (IL-18 and IL-6) (72) hyperthyroidism, and this could be due to the chronic effects of TH excess, and not to the coexisting autoimmune condition at the basis of GD (73).

A Th1 immune response is more prevalent in the active phase of GD, and GO; CXCR3 and Th1 chemokines (CXCL9, CXCL10, CXCL11) are crucial in this process, while a switch in immune prevalence from a Th1 to a Th2 response is present in the inactive or later phases (1, 74).

Systemic hyperthyroidism and T cell infiltration into orbital tissue leads to ROS production, which can aggravate GO severity by increasing T cell proliferation, adipogenesis, and GAG production in OF (75). Recently, the role of the protein tyrosine phosphatase 1B (PTP1B), encoded by the PTPN1 gene, has been characterized in GO (76). PTP1B is known to be involved in immune cell signaling by regulating cytokines via dephosphorylation of janus kinase (JAK)2, signal transducer and activator of transcription (STAT)5, and tyrosine kinase (TYK)2. After 24 h of transfection with PTPN1 siRNA, the fibroblasts were exposed to IL-1β, cigarette smoke extract (CSE), H2O2, and transforming growth factor (TGF)-β stimulations. PTPN1 silencing ameliorated ROS generation in both CSE- and H₂O₂-stimulated cells. The changes in the phosphorylation level of multiple transcription factors after PTP1B inhibition in GO OF indicate a more complex network of signaling pathways. PTP1B inhibition suppressed IL-1βinduced Akt, and JNK phosphorylation, but p38 MAPK phosphorylation was reduced only in GO OF. As the p38 and JNK pathways, and the MAPK pathway, can mediate the transcription and translation of inflammatory cytokines, this study demonstrated that PTP1B mediates inflammatory reactions in GO OF (76).

Inflammation and cytokine production, adipogenesis, and HA synthesis are the prevailing processes implicated in the pathogenesis of GO (20). In the initial phases of GO, the prevalent Th1 immune response facilitates cell-mediated immunity, leading to the production of IFN- γ , TNF- α , IL-1 β , and IL-2, that increase fibroblast proliferation and GAG synthesis. IFN- γ induces the secretion of Th1 chemokines by fibroblasts, and the migration of lymphocytes is promoted (54). IL-1 β further stimulates the synthesis of GAG (77) and the production of CCL2, CCL5, IL-6, IL-16, IL-8, and neutrophils, by OF (78). Then, inflammation leads to Th2 lymphocytes activation, that release cytokines (i.e., IL-13, IL-10, IL-5, and IL-4), and to humoral reactions and the production of IgG (79). Tissue remodeling and fibrosis characterize the late phase of GO (20).

OF express also the costimulatory protein CD40, and its binding to the ligand CD154, on T cells, induces the production of different inflammatory mediators by OF [CCL2, IL-1 α , prostaglandin E2 (PGE2), IL-6, and IL-8] (80–82).

GD OF include cell subsets with specific cellular markers, such as the CD34+ CXCR4 + Collagen 1+ phenotype (CD34+ OF), while CD34- OF do not display them. On the other hand, OF obtained from normal subjects are CD34-OF (55). The axon guidance glycoprotein Slit2 is produced by CD34- OF, and it inhibits fibrocyte differentiation and modulates their characteristic gene expression profile. Slit2 upregulates IL-12 expression and it attenuates that of IL-23, which are both involved in GD and GO (83). CD34+ fibrocytes obtained from peripheral blood mononuclear cells (PBMCs) produce low basal levels of HA, very few of which were affected by bovine thyrotropin (bTSH). On the other hand, GD OF synthesize higher levels of HA, both basally and after the treatment with bTSH. The treatment of confluent cultures with rhSlit2 increases HA production in fibrocytes, while the knock-down of Slit2 expression attenuates its synthesis in GD OF. Considering HA synthase isoenzymes (HAS1-3), low HAS1 levels are present in fibrocytes, while HAS2 is the most strongly expressed in GD OF. Slit2 alters the pattern of HAS and uridine diphosphate (UDP) glucose dehydrogenase (UDPGD) expression and cytokine production, both in GD OF and fibrocytes. The exogenous rhSlit2 causes a transition from HAS1 dominating expression to that of HAS2 in fibrocytes, decreasing also the expression of TNF- α and IL-6. These data suggest that HA synthesis in GD OF depends on the CD34- OF cell subset, while TNF-α and IL-6 expression is present in CD34+ OF. For these reasons GD OF, comprising both CD34+ OF and CD34- OF, can generate inflammatory factors (84, 85).

The activation of OF by TRAb suggests the link between GD and GO. The increased OF activity contributes to the fibrosis of the orbital tissues, causing inflammatory cell infiltration, and edema. Consequently, the optic nerve can be compressed leading to optic neuropathy. The inflammation and swelling of the eye muscles are involved in the final exophthalmos (20).

TABLE 1 | Main cytokines implicated in Graves' ophthalmopathy, the cells producing them and their biological effects.

| Retrobulbar Cells | Main Cytokines | Biological effects | References |
|-------------------------------|---|---|------------|
| Fibroblasts/ preadipocytes | IFN- γ , TNF- α , IL-1 α has been observed in tissue sections and in primary OF cultures of patients with active GO | IFN- γ , TNF- α , IL-1 α in fibroblast stimulate: -the expression of a 72-kDa heat shock protein (HSP 72) -the expression of inter-cellular adhesion molecule-1 (ICAM-I). IFN- γ , TNF- α enhance the expression of HLA-DR | (50) |
| | IL-6, 8, 16, RANTES, MCP-1, IFN- γ , TNF- α , IL-1 | IL-16 acts as a ligand for CD4+ cells, and it is important for T-cell trafficking. IL-16 production is believed to follow that of RANTES, and both are responsible for T-cell trafficking in orbital and thyroid fibroblasts | (51) |
| | In retrobulbar fibroblasts and preadipocytes obtained from GO patients: 1)IFN-γ induced CXCL10 secretion in a dosedependent manner; | CXCL10 induces the migration of Th1 lymphocytes into the orbit, thereby perpetuating the autoimmune cascade | (52, 53) |
| | 2)TNF- α alone was not able to induce chemokine secretion; 3)IFN- γ +TNF- α synergistically increased CXCL10 secretion | | |
| | In retrobulbar fibroblasts and preadipocytes obtained from GO patients: 1) IFN-γ alone dose dependently induced the secretion of CXCL9 and CXCL11; 2) IFN-γ+TNF-α combination leads to a huge response of CXCL9 | C-X-C chemokines participate in the self-perpetuation of inflammation | (54) |
| | -Cytokines detected <i>in situ</i> in GO include TNF-α, IL-1α, IL-6, IL-8, IL-10, IL-12, IL-13, and IFN-γ, -Several of these are more highly expressed in active vs stable disease and include IL-1β, IL-6, IL-8, and IL-10; -predominance of T helper (Th)1 cytokines in active GO | -Both IFN- γ and TNF- α induce B cell activating factor in GD OF -IL-1 β induces both IL-16 and RANTES in GD OF, enhancing the release of T cell migration-promoting activity | (55) |
| | In cultured primary OF from GO patients: -IL-17A combined with CD40L could induce the production of RANTES in time- and dose-dependent modifications; -IL-17A alone was not enough sufficient to trigger RANTES release | Amplification of GO inflammatory process | (56) |
| | In primary cell cultures of GO fibroblasts and preadipocytes: 1)TNF-α increases the secretion of CXCL8 dosedependently; 2) IFN-γ stimulates the secretion of CXCL10, but it inhibits that of CXCL8 | This differential modulation of CXCL10 and CXCL8 chemokines could reflect a different role of the two chemokines during the course of the disease, as CXCL10 could be associated with the initial phase of the disease when a Th1 immune response (induced by IFN- γ) is preponderant, while CXCL8 could be associated with a later chronic phase of the disease, when there is a switch to a Th2 prevalent immune response (induced by TNF- α) | (57) |
| Adipocytes | High levels of MCP-1 mRNA in the orbital fat tissue of patients with GO have been reported | MCP-1 positively correlated with the degree of macrophage infiltration in patients with GO | (58) |
| Muscle cells | In primary extraocular muscle (EOM) cultures from patients with GO: 1)IFN-γ induced CXCL10 secretion in a dose-dependent manner; 2)IFN-γ+TNF-α synergistically increased CXCL10 secretion; | Self-perpetuation of inflammation | (59) |
| | 3)IFN- γ and TNF- α induce CCL2 secretion | | |

GO, Graves' ophthalmopathy; IP-10, IFN-γ-inducible protein 10;C-X-C motif (CXCL)10, chemokine ligand 10; IFN-γ, interferon-γ, IL, interleukin; MCP-1/CCL2, monocyte chemoattractant protein-1; OF, orbital fibroblasts; TNF-α, tumor necrosis factor-α.

Furthermore, TSH-R-expressing T cells, which could be activated by TRAb, could stimulate adipogenesis of OF in GO through a peroxisome proliferator-activated receptor-γ (PPAR-γ) ligand produced *via* upregulated cyclooxygenase (4). CD34+ fibrocytes, such as OF, migrate from the circulation into sites of inflammation and injury and express two of the major GO thyroid autoantigens, TSH-R and thyroglobulin (4). Both CD4+ and CD8+ T cells, and B cells, are present in the majority of GO orbits, and the level of infiltration correlates with disease activity (86).

Macrophages are found in the orbits in early disease, whereas monocytes and mast cells have been associated with secretion of platelet-derived growth factor, that stimulates OF proliferation and HA production, in particular the platelet-derived growth factor-BB isoform in OF from both patients with/without GO (4). Mast cells also produce prostaglandins, which are able to enhance adipogenesis (87). In light of the above, there are multiple overlapping factors in the development of GO, and cytokines are strongly involved in its pathogenesis.

DRUGS TARGETING CYTOKINES, IN GD AND GO

Drugs that target cytokines (etanercept, tocilizumab, infliximab, adalimumab) have been tested in GO, with encouraging results.

OF have an increased expression of TSH-R, and a strong upregulation of TNF- α and IL-6, in the pathogenesis of GO (88).

The association between elevated circulating TNF- α levels and the severity of GO has led to the use of mAb against TNF- α , such as infliximab and etanercept (89, 90). The effectiveness of infliximab in severe steroid and surgical-resistant GO has recently been demonstrated in three cases, with complete resolution after three doses of 5 mg/kg body weight given 1 month apart (91, 92). Moreover, a positive effect of infliximab administration on active GO in a 58-year-old woman with GD has been described, in whom a single dose of this drug resulted in a reduction of inflammation and improvement of visual function, determined by magnetic resonance imaging and CAS and NO SPECS scales, with no noticeable short-term side effects (93).

Etanercept is usually used to treat RA (29). The use of etanercept has been evaluated as potential treatment in 10 patients with active GO, showing remission in 6/10 (89). Moreover, a paper reported the case of a woman, initially diagnosed with primary hypothyroidism (in substitutive treatment with levothyroxine) and subsequently with RA, who had insufficient therapeutic effectiveness with a conventional medication. After 3 years, she showed symptoms and signs of GO. Then, the patient received etanercept for RA, and after 4 months, the ocular symptoms ameliorated and exophthalmos decreased, showing that RA and GO can share similar pathogenic characteristics (88).

Furthermore adalimumab, that has been approved by FDA for psoriatic arthritis, ankylosing spondylitis, inflammatory bowel disease, RA, showed a significant amelioration in the inflammatory composite score in a retrospective review (94).

In patients with active GO, IL-6 and its soluble receptor are activated. Tocilizumab is a humanized mAb recognizing the IL-6R, which attained the approval in Castleman's disease, systemic juvenile idiopathic arthritis, and RA. A prospective nonrandomized study has been conducted in 18 GO patients (refractory to CS) treated with tocilizumab (95). An amelioration of proptosis was reported in 13 patients, extraocular motility in 15, and 7/13 solved the problem of diplopia, with no relapse of GO at the end of the follow-up. These data suggested that tocilizumab might be effective in GO patients resistant to steroids (95). Furthermore, a reduction in extraocular muscle thickness and chemosis was reported after therapy with tocilizumab in patients with GO, by using an optical coherence tomography. Four women and one man with a median age of 52 years (range, 38-73 years) were enrolled. Median GO activity duration was 17 months (12-18). After tocilizumab, median muscle thicknesses and chemosis reduced. Median CAS decreased from 5 (4-8) to 1 (0-3) (96). More recently, a paper reported three GO patients refractory to CS or

with advanced diplopia, receiving tocilizumab (8 mg/kg; monthly iv), and it showed a significant amelioration in eye symptoms (97).

CONCLUSION

GO is one of the extrathyroidal manifestations of GD. In patients with GO, OF can differentiate into myofibroblasts or adipocytes, able to interact with mononuclear cells, that produce chemoattractants and cytokines, in this way reiterating orbital inflammation.

Certain cytokines are elevated in autoimmune (i.e. IL-18 and IL-6) and non-autoimmune hyperthyroidism (i.e. TNF- α , IL-8, and IL-6), supporting the idea that this may be associated with the chronic effects of TH increase, and not with the GD inflammatory, autoimmune condition.

A prevalent Th1 immune response is reported in the immune-pathogenesis of GD and GO; Th1 chemokines (CXCL9, CXCL10, CXCL11), and the CXCR3, are crucial in this process

Patients with active, mild GO usually benefit from local therapies and selenium, whereas patients with moderate-to-severe disease generally need the addition of iv glucocorticoids. In case of an inadequate response to glucocorticoid therapy, different second-line therapies have been evaluated, such as orbital radiotherapy (with additional glucocorticoids), RTX, mycophenolate mofetil, cyclosporine, and methotrexate. Novel biologic agents, in particular teprotumumab and tocilizumab, have shown strong reductions in disease activity and severity. If these data are confirmed in the future, the treatment paradigm could be changed. Moreover, new immunotherapies are now evaluated for GD, that may have treatment implications also for GO (30).

At present, the more deepened understanding of GO pathophysiology has led to different immune-modulant treatments. Cytokines, TSH-R and IGF-1R (on the surface of B and T lymphocytes, and fibroblasts), and chemokines implicated in the autoimmune process, are possible targets of novel therapies.

The remodeled tissues in GO are dominated by adipogenesis, the increase of HA into the orbit, and the local synthesis of proinflammatory cytokines (including TNF- α and IL-6). There is a high complexity in the interactions among the cells of the heterogeneous population of GD OF in the GO orbit. Slit2 seems to play an important role in the determination of the pattern of HAS and UDPGD expression and IL-6, TNF- α , and HA production in these fibroblasts, and it could be considered as an interesting therapeutic target in GO (84, 85).

In the pathogenesis of GO, OF have an increased expression of TSH-R, and a strong up-regulation of TNF- α and IL-6. Drugs that target cytokines have been tested in GO, with encouraging results. The association between elevated circulating TNF- α levels and the severity of GO has led to the use of mAb against TNF- α , such as infliximab, adalimumab, and etanercept. Also

tocilizumab (anti-IL6-R) have reached significant findings in GO.

In conclusion, large, controlled and randomized studies are needed to evaluate new possible targeted therapies for GO.

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AUTHOR CONTRIBUTIONS

PF, SMF, SB, and AA conceived the paper. All authors contributed to the article and approved the submitted version.

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Conflict of Interest: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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A 'Real Life' Service Evaluation Model for Multidisciplinary Thyroid Eye Services

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Background/Aims: There is no universal consensus on the practical implementation and evaluation of the Amsterdam Declaration on Graves Orbitopathy in a Multidisciplinary Thyroid Eye Disease (MDTED) pathway. Recent recommendations from the UK TEAMeD-5 and BOPSS initiative highlight the importance of prevention, screening, and prompt referral of patients with moderate to severe and sight-threatening thyroid eye disease to multidisciplinary (MDTED) clinics and recommends annual auditing. We propose a practical service evaluation model with Key Performance Indicators (KPI) that are achievable and could be implemented across most TED pathways.

Material and Methods: We conducted a service evaluation from an integrated TED pathway in London with three MDTED clinics. Data was collected retrospectively from consecutive TED patients included: 1) Patient demographics, 2) Referral to first appointment time, 3) Documented smoking cessation and selenium supplementation advice, 4) Presenting disease activity and severity, 5) Investigations and treatments, including radio-iodine, 6) Time from decision to treatment initiation, 7) Initial and subsequent thyroid status.

Results: The median age was 49.0 yrs, 77.5% (183/236) were female and 49.5% (101/204) Afro-Caribbean or Asian. At their first clinic attendance, 47.6% (110/231) were biochemically euthyroid and 76.7% (79/103) at discharge. All 23.1% (52/225) current smokers received smoking cessation advice and 64.8% (153/236) received selenium supplementation advice. Intravenous methylprednisolone was given to 33.9% (80/236) patients and 12.7% (30/236) received second-line immunosuppression. All 7.2% (17/236) patients with sight-threatening disease received treatment within two weeks of diagnosis.

Conclusions: This study forms a waymark for other units using TEAMeD-5 and BOPSS audit criteria. Dedicated electronic patient records with ongoing data capture, including quality of life assessments, and diagnostic coding would significantly aid future auditing,

improve patient care, and facilitate a national audit of TED management. A future survey when the TED standards have become embedded would be instructive to see whether this has improved TED care.

Keywords: thyroid-associated ophthalmopathy, Graves ophthalmopathy, optic neuropathy, Graves disease, Graves orbitopathy, thyroid eye disease

INTRODUCTION

Thyroid Eye Disease (TED) is a distressing and sometimes sight-threatening complication of Graves' Disease (GD) which affects up to 400,000 people in the UK and an estimated 9 per 10,000 population in Europe (1, 2). Women are more commonly affected than men (1, 3). There have been several European initiatives including Amsterdam Declaration of 2009 (3) and the 2016 European Group on Graves Orbitopathy (EUGOGO) recommendations for timely therapeutic interventions in order to optimise outcomes (4, 5).

In response TEAMeD-5 (Thyroid Eye Disease Amsterdam Declaration Implementation Group) was launched in 2017 (6) to raise awareness among endocrinologists and patients with GD about the importance of prevention, early detection and prompt referral of moderate to severe and sight-threatening TED to a specialist multidisciplinary thyroid eye disease (MDTED) clinic (4). TEAMeD-5 recommended mainly endocrinological Key Performance Indications (KPIs). In 2019, TEAMeD and the British Oculoplastic Surgery Society (BOPSS) published further recommendations based on a national survey of TED services and that were endorsed by all the UK TED stakeholder organizations. In our study, we aimed to carry out a service evaluation of a large multi-disciplinary TED network in a multi-ethnic metropolitan setting, as a waymark for implementation of the aforementioned recommendations.

MATERIAL AND METHODS

Patient Recruitment

We conducted a retrospective observational study of patient databases from three MDTED clinics led by consultant ophthalmic and oculoplastic surgeons (VL, RJ and AA), and endocrinology consultants (including KM VB and CF). The patients' care also included orthoptists who assessed for ocular motility disturbance and an immunosuppression specialist (RA), for patients requiring second-line immuno suppression. There is an integrated network of other specialists including head and neck radiologists, radiotherapists for orbital radiotherapy, thyroid surgeons for thyroidectomy and an elective rehabilitation pathway (including orbital decompression ocular muscle surgery for diplopia and eyelid surgery) for patients with quiescent residual disfiguring and debilitating disease.

We formulated the KPIs used in this study from the 2019 recommendations of the British Oculoplastic Surgery Society (BOPSS) and TEAMeD-5 (**Table 1**) (7).

Patients were identified from MDTED clinic databases at Imperial College Healthcare NHS Trust and Central Middlesex Hospital (CMH) (part of London North West University Healthcare NHS Trust). Data was collected at a contemporaneous post clinic 'board round' consensus of all the MDT consultants, who are also authors of this paper.

Additional clinical data was collected from the electronic record systems and included clinic letters, biochemical and radiology results. All patient data was collected from inception of the MDT clinic. The duration of the service evaluation started from the inception of the MDT clinics - CMH (Nov 2011), Imperial College sites at the Western Eye Hospital (WEH) Jan 2016) and Charing Cross Hospital (CXH) Jan 2017).

Database lock was on 31st January 2019, which was six months before the initiation of analysis.

This study was registered and approved by the Audit Department of the Trusts. This study adhered to the tenets of the Declaration of Helsinki.

Clinical Assessment

All TED patients had a Clinical Activity Score (CAS) assigned by a consultant ophthalmologist at each visit. A CAS of \geq 3 defined clinically active TED, although other factors such as proptosis, ocular motility disturbance and optic nerve compromise were also taken into account. The severity of the disease was graded according to the EUGOGO categories of disease severity (mild, moderate to severe and sight-threatening) (4). All patients with a CAS of 3 or above were classified as having at least moderate disease severity. MRI was the imaging modality of choice as it offers superior soft tissue resolution making it an appropriate assessment of TED, which is a soft tissue disease. First-line immunosuppression was once weekly intravenous methylprednisolone (IVMP) infusions given over 12 weeks, as recommended by the EUGOGO consensus (4). All patients had at least one endocrine assessment, with optimisation of thyroid status, where appropriate. In the majority of patients with hyperthyroidism, the treatment was either antithyroid medication (Carbimazole first-line and Propylthiouracil second-line), thyroxine replacement, or both (i.e., 'block and replace'). Hypothyroid patients and post thyroidectomy patients were treated with thyroxine replacement guided by their thyroid function.

The following data was retrospectively collected for each patient:

1) Demographic data: age, gender, ethnicity, family history (in first or second-degree relatives) of thyroid disease, smoking status (current, ex-smoker, never smoked), diabetes mellitus (type 1 and 2)

TABLE 1 | The British Oculoplastic Surgery Society proposed audit criteria for the review of services managing thyroid eye disease (6) and the findings of this study.

i) **BOPSS** audit criteria **Audit results** Consideration of oral selenium supplements for patients with mild, active (1) Out of the 135 new patients seen after the introduction of TEAMeD-5 in 2017. Efficacy 60.0% (81/135) were advised to take oral selenium supplements. Smoking cessation advice for patients who are smokers. The 23.1% (52/223) documented smokers in our cohort 100% received smoking cessation advise. Prompt correction of dysthyroidism and maintenance of euthyroidism. Not formally audited. 76.7% (79/103) were biochemically euthyroid at discharge from MDTED clinic Where systemic steroid treatment is indicated, use of intravenous pulses Departmental SOP is IVMP only for first line immunosuppression. No patients had of methylprednisolone in preference to oral steroids. oral prednisolone as first-line treatment. 80/236 (33.9%) of our patients received IVMP. Number receiving urgent treatment for sight-threatening orbitopathy 7.20% (17/236) received the EUGOGO protocol of 3-day high dose IVMP for including surgical decompression for patients who fail to respond to high sight-threatening orbitopathy followed by the standard 12 week course. dose intravenous steroids. 9 of these 17 patients also underwent emergency decompression surgery. Prevalence of patients treated orbital irradiation. 38/236 (16.1%) patients received orbital irradiation. Patients undergoing elective orbital decompression and rehabilitative 16/236 (6.78%) patients underwent elective orbital decompression as part of surgery for patients with inactive or minimally active disease who are rehabilitative surgery for TED. significantly impaired (socially or psychologically) as a result of TED. (2)Patient education for recognising side-effects of steroids and other Our patients receive information leaflets provided by the local trust and leading Safety immunosuppressive treatments. TED patient charities on the side effects of steroid and immunosuppressive medication. 12.7% (30/236) of our patients received second-line immunosuppression under care of the MDTED immunosuppression specialist. Appropriate selection of patients with TED who are being considered for There is a departmental SOP for RAI treatment, with recommendations for radioiodine for suitability of steroid cover. adjunctive oral steroid cover for at risk patients. 25% (2/8) patients under our MDTED clinic received steroid cover. The remainder were sent for RAI suitability screening. Safe use of immunosuppressive treatments (exclusion of those for whom 100% (80/80) of those requiring intravenous steroids were given in the day unit of there are contraindications, assessment, and monitoring of risks of a large teaching hospital with acute services. serious adverse effects 100% (30/30) of those requiring second-line immunosuppression received this under the care of an immunosuppression specialist who is part of the MDTED clinic Timely assessment of response to high dose intravenous steroids and There is a SOP that patients are seen monthly in MDT clinic and their treatment withdrawal of steroid treatment in favour of other therapies for those with response is assessed, with a personalised management plan formulated. Our inadequate response. departments compliance with SOP was not formally audited Availability of good quality information about GO, its usual course, likely 100% (236/236) given written information on TED. Patientoutcomes, and potential treatments, complemented by high quality centred written information and access to patient-led organisations. care Formulation of personalised management plans following 100% (236/236) of the patient management plans are discussed with consultant multidisciplinary discussion. Endocrinologist and Ophthalmologists at every visit. A board round is conducted at the end of each clinic to ensure a consensual and personalised management plan for each patient. Good communication between the clinical team and the patient. 100% (236/236) patients have their management plans discussed with them and are sent copies of clinic letters. Patient engagement with the decision process about management of Use of validated tools to assess the impact of GO on their quality of life. Our study found that there was no uniform QOL collection as a result of this finding the departmental SOP was changed to ensure MDTED patients have a GOQOL assessment at each clinic visit Patients with sight-threatening TED (dysthyroid optic neuropathy 100% (17/17) achieved this. (4)Timely resulting in significant reduction in visual acuity, corneal breakdown with impending or established infection, globe subluxation) should be treated urgently within 2 weeks Median (range) decision to treatment time was 1.6 (0-6) days. Patients with moderate-to-severe, active TED should be offered 96.3% (77/80) received treatment within six weeks from presentation. treatment within six weeks from presentation Multiple surgical treatments in patients requiring complex rehabilitative This sequence was not audited, however it is in the departmental SOP. surgery, should follow the sequence: orbital decompression/eye muscle surgery/lid surgery

BOPSS, British Oculoplastic Surgery Society; IVMP, Intravenous methylprednisolone; MDTED, Multidisciplinary Thyroid Eye Disease Clinic; RAI, Radio-iodine; SOP, Standard operating protocol; TED, Thyroid Eye Disease.

75.2% (177/236) were seen in a MDTED clinic within three months of referral.

Efficient be well-defined and seamless

Referral pathways from primary to secondary and tertiary care, should

- 2) Endocrine data:
- i) Thyroid status at first MDTED clinic visit
- ii) Management of thyroid disease treatment (drugs, radioiodine (RAI), thyroidectomy), endocrine control and relapse of thyroid dysfunction
- iii) Initial thyroid status of patients at referral and during period of treatment and at the last follow-up visit
- iv) Number of patients who received RAI treatment
- v) TSH-R antibody status
- 3) MDTED clinical metrics data:
- i) Time between referral and a specialist review at first MDTED clinic
- ii) Disease activity (determined by CAS at the first and last MDTED clinic). Patients with a CAS ≥3 were deemed to have active disease.
- iii) Disease severity (determined by a clinical assessment) in line with EUGOGO classification
- iv) Outcome of patients referred to MDTED clinic (with a confirmed diagnosis of TED), including investigations and treatment given
- 4) Management of TED:
- i) Duration of treatment for TED: this was estimated from the patient reported date of TED onset, date of referral and of first MDTED clinic
- ii) Number of patients receiving documented advice on smoking cessation and selenium supplementation
- iii) Time from decision to treat to initiation of treatment for moderate to severe and sight-threatening active disease. Time from decision to treatment was recorded for: IVMP infusions, orbital decompression (urgent and elective) and orbital radiotherapy.

Inclusion and Exclusion Criteria

Inclusion criteria:

- i) At least 18 years of age
- ii) A diagnosis of TED (any severity, active or inactive) Exclusion criteria:
- i) Seen at MDTED, where a diagnosis of TED was excluded
- ii) Where the patient's endocrinology care was not at Imperial or CMH and where it was not possible to access their blood results or advise on their endocrine care.

Statistical Analyses

Statistical analyses were performed using IBM SPSS statistics v.25 and GraphPad Prism v.8. The demographic data are presented as number and percentage. Descriptive statistics for non-normally distributed data used median and inter-quartile range.

RESULTS

Patient Demographics

There were 303 patients who attended the three MDTED clinics over the study period. After applying the exclusion criteria, 49 patients were excluded from the study as they did not have a diagnosis of TED; 15 patients had inadequate records and 3 were seen at other hospitals where we could not access their endocrine records. Consequently, 236 patients were included in this study; 106 were at CMH, 82 at the WEH, and 48 at CXH. The baseline patient demographic data are shown in **Table 2**. Out of the 13.6% (32/236) patients who were diabetic: 12.5% (4/32) were type 1, 84.4% (27/32) type 2 and one patient had diabetes secondary to thyroid hormone resistance syndrome with a G332E mutation.

TABLE 2 | Patient demographics across 3 multidisciplinary clinics.

| | All TED group n = 236 | | Central Middlesex Hospital n = 106 | | Western Eye Hospital n = 82 | | Charing Cross Hospital n = 48 | |
|---------------------------------|--------------------------|------------|---------------------------------------|-----------|--------------------------------|--------------|----------------------------------|----------------|
| | n | | п | | n | | n | |
| Age at first TED clinic(yrs) | 236 | | 106 | | 82 | | 48 | |
| Median (IQR) | | 49 (36-57) | | 43 (33-55 | | 52 (43.8-60) | | 48 (34.5-54.8) |
| Range | | 18 - 82 | | 18 - 82 | | 23 - 82 | | 23 - 77 |
| No. (%) Female | 236 | 183 (77.5) | 106 | 83 (78.3) | 82 | 63 (76.8) | 48 | 37 (77.1) |
| Ethnicity | 204 ^a | | 99 ^b | | 69° | | 36 ^d | |
| No. (%) Caucasian | | 71 (34.8) | | 25 (25.3) | | 26 (37.7) | | 20 (55.6) |
| No. (%) Afro-Caribbean | | 55 (27.0) | | 36 (36.4) | | 12 (17.4) | | 7 (19.4) |
| No. (%) Asian | | 46 (22.5) | | 33 (33.3) | | 9 (13.0) | | 4 (11.1) |
| Smoking status | 223 ^e | | 106 | | 69° | | 48 | |
| No. (%) Current smokers | | 52 (23.1) | | 26 (24.5) | | 13 (18.8) | | 13 (27.1) |
| No. (%) Ex-smokers | | 41 (17.9) | | 12 (11.3) | | 18 (26.1) | | 11 (20.8) |
| No. (%) Positive family history | 236 | 74 (31.4) | 106 | 34 (32.1) | 82 | 23 (28.0) | 48 | 17 (35.4) |
| No. (%) Diabetes | 236 | 32 (13.6) | 106 | 11 (10.4) | 82 | 14 (17.1) | 48 | 7 (14.6) |
| Thyroid status at first clinic | 231 ^f | | 102 ⁹ | | 81 ^h | | 48 | |
| No. (%) Euthyroid | | 110 (47.6) | | 57 (55.9) | | 29 (35.8) | | 24 (50.0) |
| No. (%) Hyperthyroid | | 96 (41.6) | | 36 (35.3) | | 39 (48.1) | | 21 (43.8) |
| No. (%) Hypothyroid | | 25 (10.8) | | 9 (8.8) | | 11 (13.6) | | 5 (10.4) |

^aData unrecorded n=32, ^bdata missing n=7, ^cdata missing n=13, ^ddata missing n=12, ^edata missing n=13, ^fdata missing n=5, ^gdata missing n=4, ^hdata missing n=1.

Data recorded from inception of the clinic until January 2019; July 2012 to January 2019 for Central Middlesex Hospital, January 2016 to January 2019 for Western Eye Hospital and January 2017 to January 2019.

In bold: n refers to the number of patients (participants) which had the data available in their patient record notes for each demographic/endocrinology/ophthalmology characteristic.

Timing of Referral

After referral, 75.2% (177/236) patients were seen in a MDTED clinic within three months of referral. The median time from the patients reported first onset of symptoms and date of their first MDTED clinic was 135 days (**Figure 1**).

Management of Thyroid Dysfunction

The baseline thyroid status of the patients is shown in **Tables 3** and **4**. Patients who were euthyroid at their first MDTED appointment were the result of well controlled thyroid disease or euthyroid GD. Out of 166 patients who had TSH antibody titre measurement, 75.9% (126/166) had a positive titre status. In our cohort, 23.7% (56/236) patients had a recurrence (relapse) of hyperthyroidism; 66.7% (37/56) had this before and 33.3% (19/56) after their first attendance at the MDTED clinics. One hundred and three patients were discharged from the MDTED clinics during the study period and 76.7% (79/103) were biochemically euthyroid at discharge.

Radioiodine Treatment (RAI)

Twenty-nine patients received RAI treatment; 72.4% (21/29) had the treatment before the diagnosis of TED. Two of the eight patients with an established TED diagnosis had adjunctive oral steroid cover during their RAI treatment. The others were deemed to be low risk for eye disease progression by the MDT team, therefore were not prescribed oral steroid cover.

Smoking and Selenium

Baseline smoking status, where it was documented for patients is shown in **Table 2**. All current smokers received documented smoking cessation advice. Out of the 135 new patients seen after the introduction of TEAMeD-5 in 2017, 60.0% (81/135) received documented advice regarding the potential benefits of selenium supplementation.

Activity and Severity of TED

Two hundred and nineteen patients (91.5%) had a documented CAS at their first MDTED clinic attendance; 20.1% (44/219) were had a CAS of 3 or more. In our cohort, 7.2% (17/236) had sight-threatening disease (dysthyroid optic neuropathy and/or sight-threatening corneal exposure) (**Tables 3** and **4**).

Investigations and Treatment

Seventy percent (165/236) had one or more orbital Magnetic Resonance imaging (MRI) scans; 57.6% (95/165) were reported to have radiological evidence of orbital inflammation on their baseline scan (either qualitatively with STIR sequences or quantitatively with non-echo planar diffusion weighted imaging - DWI).

As first-line immunosuppression treatment, 33.9% (80/236) received intravenous methylprednisolone (IVMP), as recommended by the EUGOGO consensus (**Table 1**). The median time (range) from decision to treat and treatment initiation was 7 (1-90) days. In our cohort, 12.7% (30/236) received second-line immunosuppressant therapy usually with

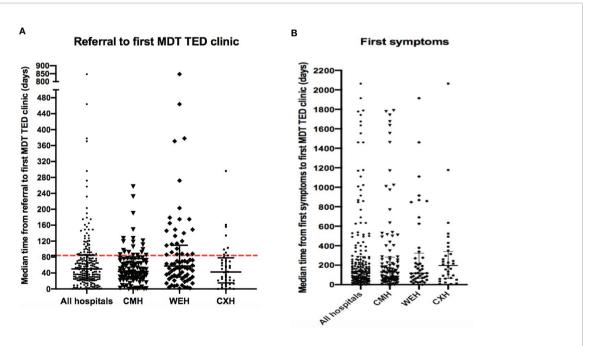


FIGURE 1 (A) Time from referral to first MDTED clinic. The median (IQR) time (days) from referral to date of first MDTED clinic. All hospitals n=215, CMH n=101, WEH n=74, CXH: n=40. Horizontal line represents the target time by TEAMeD-5 (84 days or 3 months). (B) Time from first symptoms to first MDTED clinic. The median (IQR) time (months) from patient reported onset of first symptoms of Graves' Orbitopathy and the date of their first specialist multidisciplinary thyroid eye disease clinic. All hospitals n=177, CMH n= 101, WEH n= 44, CXH n= 32. Calculation from patient data available on symptom onset and referral time. CMH, Central Middlesex Hospital (part of London North West University Healthcare NHS Trust); WEH, Western Eye Hospital; CXH, Charing Cross Hospital.

TABLE 3 | Patient baseline characteristics by disease activity at presentation.

| | Cohort 1 (CAS 0 - 1) n = 127 | | Cohort 2 (CAS 2) n = 31 | | Cohort 3 (CAS \geq 3 and non-sight threatening) $n = 44$ | | | Cohort 4 (DON) n = 17 | |
|---------------------------------|------------------------------------|------------------|-------------------------------|-----------------|--|--------------------|-----|-----------------------------|--|
| | n | | n | | n | | n | | |
| Age (yrs) | 127 | | 31 | | 44 | | 17 | | |
| Median (IQR) | | 48 (35.0 - 57.0) | | 40(31.0 - 49.0) | | 49.0 (35.5 - 53.5) | | 57.0 (53.0 - 71.5) | |
| Range | | 18 - 82 | | 21 - 73 | | 25 - 76 | | 35 - 80 | |
| No. (%) Female | 127 | 105 (82.7) | 31 | 21 (67.7) | 44 | 32 (72.7) | 17 | 12 (70.6) | |
| Ethnicity | 104 ^a | | 28 ^b | | 42° | | 15° | | |
| No. (%) Caucasian | | 36 (34.6) | | 7 (25.0) | | 16 (38.1) | | 6 (40.0) | |
| No. (%) Afro-Caribbean | | 26 (25.0) | | 9 (32.1) | | 14 (33.3) | | 3 (20.0) | |
| No. (%) Asian | | 25 (24.0) | | 10 (35.7) | | 7 (16.7) | | 2 (13.3) | |
| Smoking status | 118 ^d | | 30 ^e | | 42° | | 17 | | |
| No. (%) Current smokers | | 28 (23.7) | | 9 (30.0) | | 10 (23.8) | | 2 (11.8) | |
| No. (%) Ex-smokers | | 23 (19.5) | | 3 (10.0) | | 9 (21.4) | | 2 (11.8) | |
| No. (%) Positive family history | 127 | 39 (30.7) | 31 | 12 (38.7) | 44 | 18 (40.9) | 17 | 3 (17.6) | |

^aData missing n=23, ^bdata missing n=3, ^cdata missing n=2, ^ddata missing n=9, ^edata missing n=1.

Summary of 219 patients who had a recorded baseline clinical activity score (CAS) from July 2012 to January 2019.

DON, Dysthyroid Optic Neuropathy (sight-threatening disease).

In bold: n refers to the number of patients (participants) which had the data available in their patient record notes for each demographic/endocrinology/ophthalmology characteristic.

TABLE 4 | Endocrinological characteristics of the cohort by disease activity.

| | Cohort 1 (CAS 0 - 1) n = 127 | | Cohort 2 (CAS 2) n = 31 | | Cohort 3 (CAS ≥ 3 and non-sight threatening) n = 44 | | Cohort 4 (DON) n = 17 | |
|---|------------------------------------|----------------------|-------------------------------|---------------------|--|--------------------|-----------------------------|--------------------|
| | | | | | | | | |
| | n | | n | | n | | n | |
| Positive autoantibody | | | | | | | | |
| No. (%) TSH | 90 ^a | 59 (65.6) | 24 ^b | 20 (83.3) | 30° | 28 (93.3) | 10 ^d | 7 (70.0) |
| No. (%) TPO | 64 ^e | 31 (48.4) | 20 ^f | 8 (40.0) | 23 ^g | 13 (56.5) | 7 ^h | 3 (42.9) |
| TSH antibody titre (AU/mL) | 90 ^a | | | | | | | |
| Median (IQR) | | 4.2 (1.1 -13.7) | 24 ^b | 12.3 (3.6 - 20.9) | 30° | 15.4 (5.0 - 29.7) | 8 ⁱ | 4.0 (1.2 - 14.4) |
| Range | | 0.5 - 100.0 | | 0.7 - 59.6 | | 0.3 - 56.8 | | 0.9 - 15.0 |
| TPO antibody titre (AU/mL) | | | | | | | | |
| Median (IQR) | 46 ^j | 121.5 (12.7 - 357.2) | 20 ^f | 152.0 (5.3 - 285.0) | 23 ^g | 196.0 (20.0-491.2) | 7 ^h | 56.5 (31.1 - 83.2) |
| Range | | 2.0 - 2379.0 | | 1.0 - 617.0 | | 1.0 - 879.0 | | 25.8 - 88.9 |
| Free T3 (pmol/L) | | | | | | | | |
| Median (IQR) | 112 ^k | 5.7 (3.8 - 12.7) | 30 ¹ | 7.8 (4.0- 15.5) | 43 ¹ | 7.3 (4.4 - 22.6) | 13 ^m | 7.1 (4.8 - 14.9) |
| Range | | 0.4 - 46.2 | | 1.6 - 46.1 | | 0.9 - 47.1 | | 0.9 - 46.1 |
| Thyroid status at first clinic | 122° | | | | | | | |
| No.(%) Hyperthyroid | 36 (29.5) | 31 | 18 (58.1) | 44 | 26 (59.1) | 16 ¹ | 7 (43.8) | |
| No. (%) Hypothyroid | 15 (12.3) | | 2 (6.5) | | 6 (13.6) | | 1 (6.3) | |
| No. (%) Euthyroid | 71 (58.2) | | 11 (35.5) | | 12 (27.3) | | 8 (50.0) | |
| No. (%) Diabetes | 127 | 15 (11.8) | 31 | 3 (9.7) | 44 | 7 (15.9) | 17 | 4 (23.5) |
| HbA1c (mmol/l) | 72 ^p | | | | | | | |
| Median (IQR) | | 38.2 (36.0 - 42.0) | 20 ^f | 38.0 (35.3 - 40.8) | 31 ^q | 40.0 (36.0 - 45.0) | 14 ^r | 42.0 (36.0 - 45.3) |
| Range | | 26.0 - 48.0 | | 33.0 - 66.0 | | 30.0 - 77.0 | | 35.0 - 61.0 |
| Thyroid medication | 87 ^s | | | | | | | |
| No. (%) Carbimazole (block and replace) | | 12 (13.8) | | 5 (23.8) | | 6 (16.2) | | 1 (7.7) |
| No. (%) Carbimazole (titration) | | 58 (67.7) | 21 ^h | 12 (57.1) | 37 ^d | 23 (62.2) | 13 ^m | 9 (69.2) |
| No. (%) PTU | | 9 (10.3) | | 1 (4.8) | | 5 (13.5) | | 1 (7.7) |
| No. (%) Thyroxine | | 8 (9.2) | | 3 (14.3) | | 3 (8.1) | | 2 (15.4) |
| No. (%) Previous radioiodine | 127 | 11 (8.7) | 31 | 2 (6.5) | 44 | 5 (11.4) | 17 | 1 (5.9) |
| No. (%) Thyroidectomy | 127 | 19 (16.8) | 31 | 1 (3.2) | 44 | 7 (14.3) | 17 | 0 (0.0) |

^aData unrecorded n=37, ^bdata unrecorded n=14, ^ddata unrecorded n=14, ^ddata unrecorded n=63, ^fdata unrecorded n=11, ^gdata unrecorded n=21, ^hdata unrecorded n=10, ^fdata unrecorded n=9, ^fdata unrecorded n=55, ^gdata unrecorded n=55, ^gdata unrecorded n=13, ^fdata unrecorded n=3, ^gdata unrecorded n

Normal ranges: TSH: <0.4AU/mL, TPO: <75 AU/mL, free T3: 2.5-5.7 pmol/L.

In bold: n refers to the number of patients (participants) which had the data available in their patient record notes for each demographic/endocrinology/ophthalmology characteristic.

Summary of the endocrinological characteristics stratified by disease activities as measured by clinical activity score (CAS).

DON; Dysthyroid Optic Neuropathy (sight-threatening disease), PTU; Propylthiouracil, TPO; Thyroid Peroxidase, TSH; Thyroid stimulating Hormone.

mycophenolate mofetil (Cellcept). Further, 16.1% (38/236) patients required orbital radiotherapy (OR). Twenty-five of these patients' notes showed time of referral and date of treatment; the median time (range) from referral to OR was 30.5 (7-70) days. Twenty-five (10.6%) patients underwent orbital decompression surgery during the follow-up period, of these nine (36.0%) were emergency decompressions as adjunctive treatment for sight-threatening disease (median decision to treatment time: 1.6 days, range 0-6). All our 17 patients with sight-threatening disease started their sight preserving treatment within 2 weeks of diagnosis (**Table 1**).

DISCUSSION

The concept of combined multidisciplinary clinics has been well established in many disciplines to harness best available expertise to optimise patient management and outcomes. It has also been shown to improve the time to diagnosis and treatment in TED (5, 6, 8). Endocrinologists and ophthalmologists simultaneously seeing patients in these MDT clinics, ensures that a consistent and coherent management plan is clearly communicated and discussed with the patient and any concerns promptly addressed with readily available subspecialist expertise. The inception of two of our clinics at CMH (2011) and WEH (2015) predate the publication of TEAMeD-5 (2017) and the BOPSS audit criteria (December 2019). Data collection for this study predated the publication of the BOPSS criteria. Therefore, our multi-centre retrospective study of three MDTED clinics can be interpreted as a 'real world' snapshot of the MDT management of thyroid eye disease prior to introduction of established TED standards in a metropolitan multi-ethnic setting.

We had a similar age range and female:male ratio to other published multi-centre audits (9, 10). Our ethnically diverse cohort was comprised of 49.5% Afro-Caribbean or South Asian patients and this reflects the multi-ethnic London population. Previous audits of MDTED clinics have been less comprehensive in describing the ophthalmological and endocrinological profile of their patients (9, 10). Our prevalence of sight-threatening disease of 7.2% is higher compared to many other studies (11-13). We also had a higher number of patients requiring immunosuppression; 33.9% and 16.2% of our cohort received IVMP and orbital radiotherapy respectively compared to the PREGO study (23.0% and 6.0%), possibly reflecting a cohort with more severe disease (13). Only 75.9% of our cohort had a TSH-receptor positive titre, demonstrating that TED is primarily a clinical (and radiological) diagnosis, although serological testing is useful to identify Graves' Disease patients at risk (as per TEAMeD-5) (6). Most of our patients who received RAI did so prior to attending a MDTED clinic. There may be a very long gap between date of RAI treatment and onset of TED, so it is difficult to deduce whether RAI has contributed to the TED (14). However, out of those who received RAI after an established diagnosis of TED, only a quarter had prophylactic oral steroid cover. Oral steroids have a range of undesirable side effects for example weight gain, diabetes, and osteoporosis (15). These adverse effects must be

considered before prescribing steroids, especially in patients deemed to be low risk for eye progression. There may also be a trend that with increasing awareness of TED, pre-RAI TED screening may account for increasing referrals to eye services. It is also encouraging that some of the key initiatives of TEAMeD-5 have already been implemented with 100% smoking cessation advice given to current smokers in our cohort. There is also increasing numbers (almost two-thirds of our cohort) who have received documented advice about selenium supplementation. In our experience, many patients have started selenium supplementation by the time of their first attendance at the MDTED clinic usually on the advice of their endocrinologists.

Limitations

This is a retrospective descriptive study at a single time point prior to the introduction of established TED standards. It would be interesting and informative to repeat this service evaluation in the future to evaluate for changes in practice after the TED standards have become embedded.

Data on every variable could not be recorded due to missing information from patients' notes. TSH-R measurement was less widespread prior to 2015, therefore only 70.3% of our patients had a recorded TSH antibody titre. In our cohort, there were some patients who had clinically active TED but a low CAS. CAS, having been developed in a predominantly white Dutch TED cohort, in our experience underestimates activity in patients with a high Fitzpatrick skin type due to its reliance on soft tissue appearance, such as erythema, and this is pertinent in a cohort of high ethnic diversity. The CAS scoring system places a greater weight on symptoms resulting from acute orbital congestion, as these comprise 5 of the total 7 points which can be allocated on the first visit. Ocular muscle involvement is not reflected in the scoring system, neither is posterior orbital pathology (16). Highly relevant signs such as diplopia at the first visit are also not included in the CAS scoring system (17). Therefore, changes in visual function are not scored for in the first visit where CAS is scored out of 7, so patients can have severe ocular motility restriction and decreased acuity that is not scored for in this setting (17). Recently, non-echo planar diffusion weighted imaging (DWI) MRI has been evaluated as a valuable tool, in adjunction with CAS, for investigating posterior orbital disease in TED and in providing higher resolution orbital images (16, 17). This imaging has been an invaluable adjunct for TED patient management in our clinics for the past eight years. Nevertheless, CAS also has a strong inter-specialty understanding and is used interchangeably by endocrinologists and ophthalmologists. Our aim with our ongoing collaboration alongside our radiological colleagues is to develop a Radiological Activity Score (RAS) to complement CAS to enable a more comprehensive assessment of disease activity and severity. It is well recognised that TED can significantly impact on Quality of Life (QoL) (18-21). QoL measurement with a validated instrument such as the GO-QOL, which is the gold standard in assessing the impact of TED, is recommended as part of the ongoing clinical assessment (19-21). However during the study period, the regular use of the validated GO-QOL questionnaire remained inconsistent in the aforementioned clinics.

Thyroid Eye Care in the Post Covid-19 Landscape

This service evaluation was carried out before the Covid-19 pandemic. In the height of the COVID pandemic, all routine face to face MDT clinic appointments were suspended. We stopped all our patients undergoing intravenous steroid treatment for non-sightthreatening disease and a significant proportion relapsed meaning we had to resume immunosuppression generally in the form of oral steroids during the lockdown period. In practice TED management is complex and relies on many face to face interactions such as orthoptic measurements, blood tests and ophthalmic and systemic assessments and we have found that a significant proportion of patients had significantly deteriorated despite reporting stability over a virtual consultation. The thyroid MDT clinic was one of the first, with the support of our management, to resume face to face activity for all patients once the height of the pandemic subsided and decisions regarding starting or resuming immunosuppression were made on the basis of these face to face consultations. With the investment of National Health Service (NHS) information technology in the UK, there may be a role for virtual MDT-lite clinics, which can enable more patients to have access to this modality, especially in parts of the country where there is significant geographical separation between endocrine and ophthalmic services. MDT clinics reduce duplication of appointments, avoids patients attending hospitals twice for the same condition and offers a consistent management strategy with a wholistic approach. We recognise however in a post COVID healthcare landscape, there will be significant fiscal constraints as regards service developments.

Implications for Future Work

There is consensus that TED management should be safe, timely, efficacious, equitable and patient-centred. In the UK, TEAMeD has worked closely with BOPSS to set up sustainable targets (7). Regular auditing of key performance indicators (KPIs) is of paramount importance. One of our recommendations is the set up of a dedicated electronic patient record platform to record this data prospectively in the clinic setting to avoid an onerous audit process, with a system to enable continuous and effortless GO-QOL evaluations at every consultation. The MDT service would also benefit from having dedicated co-ordinators as presently this administrative burden is mainly borne by clinical staff. This would be highly beneficial to provide more patientcentred care. We highlight the important of signposting patients to support organisations for additional support. In the UK, these include the British Thyroid Foundation and Thyroid Eye Disease Charitable Trust (TEDct). Ideally patients should have access to psychological counselling in view of the burden of significant disfigurement from TED. We believe that our approach,

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integrating TEAMeD-5 audit standards with the BOPSS recommendations forms a more complete service evaluation model for future MDTED services to implement the aims of the Amsterdam Declaration on Graves Orbitopathy. The clinical findings of our service evaluation in a multi-ethnic metropolitan population may not be widely applicable to all TED populations. Our main aim is to show the feasibility of producing key performance indicators for benchmarking TED care in a multi-disciplinary setting. We would look forward to other centres with a different cohort demographic to conduct a similar service evaluation to establish the range of TED presentations in an MDT setting designed to enable earlier diagnosis and treatment of TED.

DATA AVAILABILITY STATEMENT

The raw data supporting the conclusions of this article will be made available by the authors, without undue reservation.

ETHICS STATEMENT

The studies involving human participants were reviewed and approved by Imperial College Audit Office. Written informed consent for participation was not required for this study in accordance with the national legislation and the institutional requirements.

AUTHOR CONTRIBUTIONS

VL conceived and designed the study. SF acquired, analysed, and interpreted the data. SF, CF, and VL drafted the manuscript. RA, RJ, AA, VB, and KM contributed to the manuscript. All authors contributed to the article and approved the submitted version.

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Conflict of Interest: VL is the British Oculoplastic Surgery Society National Lead for TEAMeD (UK Thyroid Eye Disease Amsterdam Declaration Implementation Group).

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Orbital Signaling in Graves' Orbitopathy

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Graves' orbitopathy (GO) is a complex and poorly understood disease in which extensive remodeling of orbital tissue is dominated by adipogenesis and hyaluronan production. The resulting proptosis is disfiguring and underpins the majority of GO signs and symptoms. While there is strong evidence for the thyrotropin receptor (TSHR) being a thyroid/orbit shared autoantigen, the insulin-like growth factor 1 receptor (IGF1R) is also likely to play a key role in the disease. The pathogenesis of GO has been investigated extensively in the last decade with further understanding of some aspects of the disease. This is mainly derived by using *in vitro* and *ex vivo* analysis of the orbital tissues. Here, we have summarized the features of GO pathogenesis involving target autoantigens and their signaling pathways.

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INTRODUCTION

Graves' orbitopathy (GO) or thyroid eye disease is the most common overt thyroidal manifestation of Graves' disease (GD) with substantial morbidity and socioeconomic impact (1–4). Extensive orbital tissue remodelling in GO is mainly shown as adipose tissue expansion and tissue edema *via* increased adipogenesis and hyaluronan production, respectively. These pathogenetic processes produce disfiguring proptosis and underpin all GO signs and symptoms. There is a close clinical and temporal association between GD and GO suggesting an autoimmune response to common antigen/s in the orbit and thyroid gland. The thyrotropin receptor (TSHR) is expressed in orbital adipose tissue (OAT) (5–8) and virtually all patients with hyperthyroid GO have thyroid stimulating antibodies (TSAB). Therefore, the TSHR is the most logical candidate (9), which is further supported by the existence of TSHR-induced GO in an animal model (10). The incidence of GO is estimated to be 16/100,000 in females and 2.9/100,000 in males (11). On the other hand, the prevalence estimate is about 10/10,000 (12). A recent meta-analysis reported that current GD patients have a milder phenotype than in the past; as a consequence, a smaller proportion display

Abbreviations: GO, Graves' orbitopathy; TSAB, thyroid stimulating antibodies; TSHR, thyrotropin receptor; IGF1, insulin like growth factor 1; IBMX, 3-isobutyl-1-methylxanthine; PPARγ, peroxisome proliferator-activated receptors gamma; PI3 kinase Phosphoinositide-3- kinase; MAP, mitogen-activated protein kinase; FOXO, Forkhead Box O1; C/EBP-δ, CCAAT-enhancer-binding proteins Delta; cAMP, Cyclic adenosine monophosphate; PKA, protein kinase A; GCRP, G protein coupled receptors; FRET, fluorescence resonance energy transfer; CREB, cAMP responsive element binding protein; TG, thyroglobulin; TPO, thyroid peroxidase; NIS, sodium iodide symporter (NIS), TTF, thyroid transcription factors; PKC, Protein kinase C; PLC phospholipase C; DAG, di-acyl-glycerol; NFAT, Nuclear factor of activated T-cells; HA, hyaluronic acid; GAG, glycosaminoglycan; SNP, single nucleotide polymorphism.

GO symptoms (13). As with other autoimmune conditions there is female preponderance towards the condition with 6:1 female to male ratio, although in GO the ratio is less skewed than in GD. In addition, most patients with GO have reduced quality of life (QOL) (14) and suffer long-term psychological distress due to the disfiguring appearance of the proptosis, also known as exophthalmos (15). Available treatments for GO are unsatisfactory and more research is needed to address the pathophysiology of the disease which may lead to early preclinical diagnosis promoting preventative/early interventions. This in turn will improve long-term morbidity and socioeconomic impact.

ADIPOGENESIS

Adipogenesis is a process in which preadipocytes differentiate into mature adipocytes to form adipose tissues. Our current understanding of adipogenesis has been largely derived by using the murine 3T3L1 cell line. This cell line can spontaneously differentiate into adipocytes when maintained in a high concentration of fetal calf serum for several weeks but the process can be accelerated by employing adipogenic cocktails including insulin, steroid and 3- isobutyl-1-methylxanthine (IBMX) (16). Further components of the differentiation cocktails may also include proliferation-activated receptor gamma (PPARy) agonists such as pioglitazone and indomethacin (17). Insulin, in common with insulin-like growth factor-1 (IGF-1) activates PI3 kinase (18) and MAP (mitogenactivated protein kinase) (19) pathways. Phosphorylation of protein kinase B (PKB/Akt) in turn phosphorylates forkhead box protein O1 (FOXO1) causing it to exit from the nucleus leading to increased transcription of adipogenic genes (20). Steroids induce the expression of the early adipogenic gene, CCAAT enhancer binding protein delta (C/EBP-δ). This transcription factor contributes to an increase in PPAR-γ expression and production of prostacyclin leading to elevated intracellular cAMP. IBMX is a nonselective phosphodiesterase inhibitor whose presence further elevates levels of intracellular cAMP and protein kinase A (PKA). IBMX is thus required for transcriptional activation of the master regulator of adipogenesis, PPARγ.

Adipogenesis contributes to OAT expansion because a fibroblast has an approximate diameter of 30 microns, whereas the diameter of a mature adipocyte is approximately 150 microns, i.e. 5 times larger. The increased adipogenesis has been demonstrated by using *in vitro* cultures of human fibroblasts and analysis of *ex vivo* samples from patients with GO (21). By using both *in vitro* lineage specific differentiation protocols and flow cytometry, studies have indicated that orbital fibroblasts (OF) possess mesenchymal stem cell (MSC) properties including positivity for Thy-1 (CD90) which is a marker of MSC (22–25). In the orbit, Thy-1 negative OF can be induced to differentiate when cultured in appropriate adipogenic medium whereas Thy-1 positive cells are more likely to undergo differentiation to myofibroblasts and cause

fibrosis (23, 24). The orbital fibroblast is also able to undergo neurogenesis, myogenesis, osteogenesis and chondrogenesis *in vitro*, indicating their pluripotency (22, 25).

EXTRA-CELLULAR MATRIX

Several extracellular matrix (ECM) components are overproduced in GO including collagens and glycosaminoglycans (GAGs). The excess ECM accumulation in OAT and extraocular muscle (EOM) lead to oedema with consequent proptosis and diplopia respectively (26). The main GAG produced in GO is hyaluronic acid, which is generated by three synthase enzymes (HAS1, HAS2 and HAS3) and broken down by hyaluronidases. Activation of cAMP-protein kinase A signaling *via* the TSHR, increases cAMP response element binding protein (CREB) at CREB binding sites in the promoters of HAS1 and HAS2 genes, thereby enhancing hyaluronan production (27).

TSHR INTRACELLULAR PATHWAYS

Several studies, including from our group, have shown that activation of the TSHR in OF leads to an increase in hyaluronan production and adipogenesis (20, 28). TSHR expression has been shown to increase during adipogenesis (5). We demonstrated that 'neutral' TSHR antibodies were capable of binding but had no effect on traditional TSHR signaling pathways (described below) (29). Indeed, TSHR signaling may be far more complex than initially thought (30). Little is known about the effects of TSHR activation at various stages during differentiation. The downstream cascade triggered by TSHR will depend on the types and abundance of guanine-nucleotide binding proteins (G proteins) available in the cell (31). G protein coupled receptors (GPCR) can exist as monomers or oligomers. Oligomerization is the term used to describe dimeric, tetrameric, or higher-order complexes between GPCR monomers. The activation of different GPCR complexes will have major influence on subsequent G protein signaling pathways. The evidence that TSHR may exist in an oligomeric state was initially provided by studies using antibodies (32) and more recently by fluorescence resonance energy transfer (FRET) technology (33). Interestingly, the presence of dimerization influences TSHR behavior. Unstimulated TSHRs have been shown to form oligomers that return to the monomer state with TSH (34). TSHR autoantibodies with stimulating properties are (TSAB) proposed to favor formation of TSHR dimers, whilst TSHR blocking antibodies, are unable to bring about this conformational change. After TSH binding, a constitutively oligomeric TSHR dissociates into active monomers (or dimers when TSAB bind). Subsequently the monomers or dimers are recruited to the lipid rafts and interact with G proteins, thereby initiating the signaling cascade. In the case of TSH, the signal is rapid and brief because of faster movement of monomers into the lipid rafts, in contrast to the slow motion of the dimers. Multivalent blocking TSHR antibodies may cross-link the

oligomers, thus preventing them from dissociating and impeding their entry into lipid rafts (35). In cells with low levels of TSHR expression, homo-heterodimer formation is less likely. This may change during adipogenesis, as TSHR expression increases, and may lead to activation of different signaling cascades from that predominating in orbital fibroblasts.

TSHR is known to activate mainly the guanine-nucleotide protein alpha stimulation (Gs)-cAMP pathway. In addition, TSHR may activate several other G protein subtypes, as detailed below (36, 37), non G protein pathways such as βarrestin-1 (38) and other signalling pathways (39, 40). When TSH binds to its receptor, GTP replaces GDP in the heterotrimeric G protein, which dissociates into Gsα and Gβγ subunits with the former activating all isoforms of adenylate cyclase (41). This enzyme increases levels of cAMP in the cell and activates PKA, also known as cAMP-dependent protein kinase. The activated PKA phosphorylates multiple downstream target proteins one of which is cAMP responsive element binding protein (CREB). CREB then binds to its receptors on the promoter region of the DNA exerting various gene transcription processes including expression of thyroglobulin (TG), thyroid peroxidase (TPO), sodium iodide symporter (NIS), the thyroid transcription factors TTF1/NKx2.1, TTF2/ FoxE1, and PAX (42, 43). Every intermediary in the pathway described above may additionally interact with different molecules belonging to other pathways.

In human thyrocytes and rat FRTL-5, guanidine binding protein alpha a/alpha 11 (Gαq/α11) coupling has been shown to stimulate Protein kinase C (PKC) pathways by generating phospholipase C (PLCβ). The PKC pathways has been associated with hyaluronan generation in GO (44). Activation of PKC pathways requires supraphysiological TSH concentrations although not all research agrees with this finding (45). PLC catalyses hydrolysis of phosphatidylinositol in cell membranes yielding di-acyl-glycerol (DAG) and inositol tri phosphate (IP3) as second messengers. DAG directly stimulates PKC. IP3 increases cytosolic Ca+2 levels which act through a number of effectors including PKC itself (46) and Nuclear Factor of Activated T-cells (NFAT) transcription factor protein. NFAT plays an important role in cytokine gene transcription regulation (47). Calcium via calmodulin -a calcium sensor protein - activates the serine/threonine phosphatase calcineurin (inhibited by cyclosporin and FK506). This in turn rapidly dephosphorylates NFAT proteins, resulting in a conformational change that exposes a nuclear localization signal leading to NFAT nuclear import (48). TSHR may also couple to guanine nucleotide binding protein alpha inhibition (Gαi1), which inhibits adenylyl cyclase and decreases cAMP levels. The accompanying Gβγ dimers may induce multitudes of other pathways, including adenylyl cyclase, PI3K/Akt (PKB)-FOXO and PLC cascades (49-51). Others have reported that TSHR activation of OF signals via p70s6 kinase (52). The finding may explain our lack of success when using gain-of-function mutants of the TSHR, which signal predominantly via Gsα, to stimulate adipogenesis (28) and concurs with the study from van Ziejl et al. who investigated TSH/TSAB induced hyaluronan

production (53). It contrasts with the studies of Neumann and colleagues, who report increased M22-mediated cAMP, even at baseline. However, these authors maintain their OF in a semi-adipogenic medium which likely increases TSHR expression (54).

Our previous work has demonstrated that adipogenesis and HA production, are linked in the orbit. HA accumulation increases in the orbit during adipogenesis but not in other fat depots (55). In this study, adipogenesis in orbital preadipocytes was accompanied by HA accumulation and significantly increased HAS2 transcripts (but not HAS 1 and 3). In contrast, adipogenic differentiation in subcutaneous preadipocytefibroblasts significantly decreased secreted HA and HAS2 transcript levels. IGF-I alone did not increase HAS2 levels, but inhibition of PKB/Akt increased orbital HAS2 transcripts but not subcutaneous preadipocytes. Furthermore, our study suggested that mTORC1 negative feedback in IGF1-PI3K-Akt signalling is absent in OF but present in subcutaneous adipose tissue (55). The difference might be explained by the fact that human OF originate from neural crest, while subcutaneous adipose tissue is of mesodermal origin. In addition, our most recent studies demonstrated a depot specific fatty acid-uptake driven adipogenesis with unique gene signatures in OAT. These result in hyperplastic-type expansion of adipocytes in GO (56, 57). Taken together, these findings suggest a very distinctive mechanism underlying the orbital adipogenesis process.

INSULIN LIKE GROWTH FACTOR -1 RECEPTOR SIGNALLING

While there is strong evidence supporting the role of TSHR in GO, IGF1R is also likely to play a key role in the disease progress. The IGF1R was first proposed by Weightman and colleagues who demonstrated high affinity IGF1 binding sites in OF (58). More recently extensive work from Terry Smith and his colleagues has confirmed this finding and further showed that TSHR and IGF1R co-localize to orbital cell membranes (59). The same group has further reported a wide range of IGF1R mediated effects in OF including increases in proliferation, GAG production and cytokine production (60, 61). Our own study demonstrated that activation of TSHR and IGR1R has additive effect on HAS2 transcripts/HA production (62). Krieger et al. found that M22 stimulation of HA secretion by OF involves cross talk between IGF-1R and TSHR. The relationship relies on TSHR activation per se rather than direct activation of IGF-1R which leads to synergistic stimulation of HA secretion (63). TSH induced ERK phosphorylation can be blocked by an IGF-1Rblocking monoclonal antibody suggesting that IGF-1R might mediate some TSH-provoked signalling. Further studies have highlighted the importance of down stream factors of IGF1-PI3K signalling and revealed that FOXOs, may mediate both TSHR and IGF1R signalling pathways in GO (64). The notion is further supported by recent successful trial of teprotumumabmonoclonal antibody which blocks IGF1R - in reducing proptosis in patients with GO (65, 66). Whilst effective medical

treatment for GO is welcome, some concerns have been raised about these trials including the lack of orbital imaging and the fact that despite QOL scores being improved in the teprotumumab group, all patients scores remained low (67). Furthermore, the activation of Fibroblast Growth factor (FGF) and its receptor has been shown to increase the expression of Insulin like growth factor-2 (IGF-2) in mesenchymal stem cells *via* IGF-2 and IGF1-R (68). The FGF signalling pathway has also been shown to play a role in OAT expansion in GO (69). Our most recent study used RNA-seq analysis to demonstrate that FGFs, FGFR2, IGF-2 and IGF1-R were highly expressed in OAT compared with white adipose tissue, supporting the aforementioned successful trial of IGF1R inhibition in GO (56).

TSH/TRAB binding proteins or even as autoantigens. Early northern blot analysis of thyroid tissue identified the expected full-length transcript plus 2 additional transcripts at 1.3 and 1.6 kb (70); the transcripts were also detected in OF (71). Of interest, the exon 1-8 variant is similar in structure to the TSHR A subunit which is generated following cleavage of the full-length receptor (72, 73). Furthermore, induced murine models of GD and GO are more effective when immunizing with the A subunit than with the complete TSHR (74, 75). We have reported that the 1.3 variant is expressed as a protein and can affect TSHR activation (76). Thus, these variants could have impact on the pathogenesis of GO by inducing further production of TSAB or protect against GO by 'neutralizing' TSAB, respectively.

TSHR VARIANTS

To add to the complexity of the molecular events associated with GO, several TSHR variants have been described which lack the transmembrane domain. If the variants are expressed as protein, they would yield soluble receptor products which could serve as

DISCUSSION

TSHR and IGF1 signaling are important in orbital tissues (summarized in **Figure 1**) but more complex than generally thought. Although these signals are mainly activated through G protein signalling pathways, other cascades may also be involved.

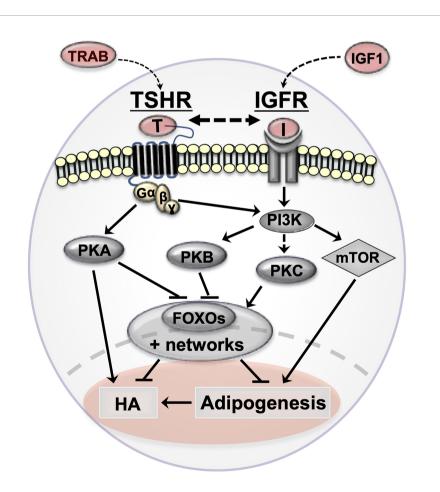


FIGURE 1 | Cartoon summarizing orbital fibroblast signaling cascades in Graves' orbitopathy (GO) and how they affect pathogenetic mechanisms (adipogenesis and hyaluronan production). TSHR/TRAB and IGFR/IGF are shown in red with arrows indicating the possible crosstalk between the pathways in GO. thyrotropin receptor (TSHR, serpentine structure); TSHR auto-antibodies (TRAB); Insulin-like growth factor 1 receptor (IGF-1R) and IGF1; protein kinase A (PKA); protein kinase B (PKB/Akt); protein kinase C (PKC); phosphoinositide 3-kinase (PI3K); forkhead box protein O (FOXO); hyaluronan production (HA).

As our understanding expands, additional extracellular or intracellular factors, which regulate signaling, may be identified. The abundance of the receptors may also dictate which pathways are activated. The recent success of TSHR extracellular domain crystallization is likely to catapult these areas of research and may lead to further alternative treatment strategies for GO (77).

As discussed above, a human monoclonal anti-IGF-1R-blocking antibody, Teprotumumab has been approved by FDA for treatment of patients with GO specifically in reducing proptosis and has recently been reported to be highly effective in active GO (65). The potential for treatments based on TSHR antagonism, which have been demonstrated to be effective *in vitro*, is keenly anticipated either with blocking antibodies or

small molecule antagonists which in theory could inhibit both TSHR and IGF-1R related and/or unrelated pathways (78). The beneficial effects on GD and GO following administration of a monoclonal TSHR blocking antibody (TBAB) in a patient with thyroid cancer has recently been described (79). Furthermore, manipulating the two pathways concomitantly may provide even more effective treatment for GO and merits investigation.

AUTHOR CONTRIBUTIONS

MD and LZ wrote the manuscript with input from ML. CD and ML reviewed the manuscript. All authors contributed to the article and approved the submitted version.

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Emerging Insights Into the Role of Epigenetics and Gut Microbiome in the Pathogenesis of Graves' Ophthalmopathy

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Wang Y, Ma X-M, Wang X, Sun X, Wang L-J, Li X-Q, Liu X-Y and Yu H-S (2022) Emerging Insights into the Role of Epigenetics and Gut Microbiome in the Pathogenesis of Graves' Ophthalmopathy. Front. Endocrinol. 12:788535. doi: 10.3389/fendo.2021.788535 Graves' Ophthalmopathy (GO) is an organ-specific autoimmune disease that is often characterized by infiltration of orbital tissues and is considered as the most common extrathyroid manifestation of Graves' disease (GD). Although genetic susceptibility has been found to be critical for the phenotype of GO, the associated risk alleles in a single gene are generally insufficient to cause the disease. Accruing evidence has shown that epigenetic disorders can act as the potentially missing link between genetic risk and clinically significant disease development. Abnormal epigenetic modifications can lead to proinflammatory cascades and activation of orbital fibroblasts (OFs) by promoting the various inflammatory response pathways and regulating the diverse signaling molecules that are involved in the fibrogenesis and adipogenesis, thereby leading to the significant expansion of orbital tissues, fibrosis and inflammation infiltration. Additionally, emerging evidence has shown that the gut microbiome can possibly drive the pathogenesis of GO by influencing the secretion of Thyrotropin receptor antibody (TRAb) and T-helper 17 (Th17)/regulatory T cells (Treg) imbalance. This paper describes the latest epigenetic research evidence and progress made in comprehending the mechanisms of GO development, such as DNA methylation, histone modification, non-coding RNAs, and the gut microbiome.

Keywords: Graves' ophthalmopathy, epigenetics, DNA methylation, histone modification, noncoding RNAs, gut microbiome

INTRODUCTION

Graves' ophthalmopathy (GO), also known as thyroid-related ophthalmopathy, is an autoimmune disease associated with Graves' disease (GD) that involves orbital tissues. It has been found that about 25% of GD patients develop GO during their disease course (1, 2). Approximately 90% of GO develop in hyperthyroidism caused by GD, whereas GO sometimes occurs in patients with euthyroid or hypothyroid chronic autoimmune thyroiditis (3, 4). Although the ratio of female to male (F/M) has been found to vary based on the different studies, the incidence of GO is significantly higher in women than in men (5). The common clinical characteristics of GO are

eyelid retraction, swelling, exophthalmos, retrobulbar pain, and even vision impairment resulting from optic neuropathy (6). Similar to GD, the autoreactive inflammatory process of orbital tissue plays a major role in the pathogenesis of GO. The extraocular muscles and connective tissue of GO patients are infiltrated by activated monocytes (such as T cells), as well as a small number of plasma cells, macrophages and mast cells (7). Activated T cells, mainly CD4⁺T cells, produce a large number of cytokines to amplify and maintain orbital inflammation and stimulate the proliferation of orbital fibroblasts (OFs) and the synthesis of glycosaminoglycans (GAG) (8). OFs are heterogeneous cells, and when activated, fibroblasts with a high level of thymocyte differentiation antigen 1(Thy1) surface expression (Thy1+) are more likely to form myofibroblasts, on the contrary, fibroblasts with little or no Thy1 (Thy1 -) are more likely to form adipocytes (9, 10). Accumulation of GAG leads to edema of extraocular muscles (11) and eventually leads to orbital swelling, tissue expansion and fibrosis development.

As mentioned above, cytokines and immune mechanisms play an important role in the pathogenesis of GO. CD4⁺T cells can be divided into auxiliary T helper 1(Th1), Th2, Th17, regulatory T (Treg) cells and other subtypes, which are considered to be related to the occurrence and development of GO (8). Studies have shown that Th1 cytokines are not only related to the progression of GO, but may play a dominant role in the orbital involvement in the early stage of GO. On the contrary, Th2 cytokines have no direct correlation with disease progression, but may play a greater role in the later stage of the disease (12-14). IL-17A secreted by Th17 cells has proinflammatory and pro-fibrosis effects, which further drives the clinical manifestation of GO (15). Treg cells are a subgroup of T cells with immune regulatory function, and impaired Th17/Treg balance may also be an important cause of GO (16). Furthermore, activated T cells activate the differentiation of B cells into plasma cells and the production of antibodies by secreting IL-4 (8). GD related hyperthyroidism is primarily caused by binding of TRAb to the thyrotropin receptor (TSHR) on the thyroid follicular endothelial cells, thereby stimulating the excessive secretion of the thyroid hormones. Furthermore, recent studies have reported that TRAb can effectively stimulate the proliferation of OFs, which is positively correlated with the clinical features of GO and can adversely affect the prognosis (17). Another crucial autoantigen involved in the pathogenesis of GO is insulin-like growth factor I receptor (IGF-IR), which forms a physical and functional complex with TSHR in OFs from GO patients (18). Antibodies of the IgG class generated by GD patients are able to bind to IGF-1R and initiate the signaling from the TSHR/IGF-IR protein complex. Furthermore, Teprotumumab, an IGF-IR blocking monoclonal antibody, has been approved in the US FDA for the treatment of GO and works by attenuating signaling from TSHR or IGF-IR (19). Currently, substantial progress has been made in treating GO, but the moderate-to-severe GO patients usually do not completely respond to available medical treatments, and complete restitutio ad integrum almost never occurs (5, 20). Therefore, more extensive studies are needed to

explore the pathogenesis of GO and develop prompt prediction, timely referral and novel therapeutic strategies for GO.

In the past decade, increasing evidence has shown that genetic susceptibility plays a crucial role in the pathogenesis of GD and GO (21-24), but the associated risk alleles in a single gene are generally insufficient to cause disease. Moreover, several studies have performed Human leukocyte antigen (HLA), cytotoxic T lymphocyte antigen-4 (CTLA4), Interleukin (IL) - 23 receptor (IL23R), and TSHR genotyping on GO and non-GO patients in GD patients. The results show no statistical differences in allele and genotype frequency between the two groups, thereby plausibly suggesting that GO might not have significant genetic susceptibility in GD patients (25). These results suggested that epigenetics and/or environmental influences could possibly also play an important role in the pathogenesis of GO. Epigenetics as a research hotspot in recent years, researchers have found that the decreased level of ICAM1 methylation is significantly associated with exophthalmos in patients with GD (26). Similarly, micro RNAs (miRNAs) have been correlated with the level of autoimmune antibodies such as TPOAb, TgAb and TRAb (27). These findings provided us with vital background to explore the potential pathogenesis of GO. A number of previous epigenetics related studies have shown that different environmental factors can also functionally regulate gene expression and phenotypes in the disease development process without altering the DNA sequences, primarily through epigenetic mechanisms such as those of DNA methylation, histone modification, and non-coding RNAs (28). Epigenetic modifications have been found to be involved in the dysregulation of the different signaling molecules and receptors in a variety of autoimmune/inflammatory conditions, including GO (29-32). Thus, epigenetic studies could serve as interesting targets for elucidating the pathogenesis of GO. Interestingly, all environmental factors involved in the pathogenesis of GD and GO can effectively alter the balance of microorganisms located in the gut and influence the immune system, particularly the proportion of Treg and Th17 cells. A recent study demonstrated that gut dysbiosis can lead to the imbalance of Th17/Treg through the propionic acid regulated pathway, and promote the occurrence of GD (33). A number of studies in GD/ GO mice model have also shown that the gut microbiome might induce the clinical manifestation of GO by influencing the levels of TRAb and Th17/Treg imbalance (34, 35). Therefore, this paper reviews the latest epigenetics landscape and progress made in deciphering the mechanisms of GO by encompassing the various aspects of DNA methylation, histone modification, noncoding RNAs and gut microbiome, with an aim to provide new effective potential therapeutic strategies for the management of GO.

ABNORMAL DNA METHYLATION AND GO

As one of the most common forms of epigenetic modification, DNA methylation is catalyzed by DNA methyltransferases (DNMTs) by using S-adenosyl-L-methionine (SAM) as a

methyl donor, and forming 5-methylcytosine (5-MC) by the transfer of the methyl group to the five carbon atoms of the CpG dinucleotide cytosine base. Accumulating evidence has suggested that DNA methylation in the promoter regions of the CpG island is primarily linked to gene transcriptional inhibition; and the methylated CpG binding domain (MBD) family can effectively recognize and bind to methylated CpG to cause significant transcriptional inhibition (36).

In recent years, several studies have shown that DNA methylation exhibits important functions in the pathogenesis of GO by altering the gene expression profile. Xin et al. (37) have identified 841 differentially methylated sites by sequencing the whole wide genomic DNA methylation in 6 patients with GO. It was found that 148 different genes were located near or at the methylation sites. These genes include various genes mediating immune responses such as Cluster of Differentiation 14 (CD14), interleukin 17 receptor E (IL17RE), beclin1(BECN1), cyclin dependent kinase 5 (CDK5), oxidative stress regulation genes including ATP-Binding cassette transporter superfamily D1 (ABCD1), homeobox B13 (HOXB13), thyroid function regulating gene such as dopamine receptor D4 (DRD4). The aforementioned genes IL17RE and CDK5 have been positively associated with Clinical Activity Score (CAS) and TRAb, respectively. The authors also performed enrichment of genes associated with the diverse biological processes, cellular composition, and molecular functions. To further understand the pathogenesis of GO, pathway analysis of these identified genes was carried out, and four potentially important pathways, including toxoplasmosis pathway, axon guidance pathway, focal adhesion pathway, and proteoglycans in cancer pathway, were found to be closely associated to the occurrence and development of GO. Moreover, other important pathway genes such as LDLR, CDK5, and phosphatidylinositol-4,5-bisphosphate 3-kinase

catalytic subunit beta (PIK3CB) that play an important role in regulating inflammatory have also been found to be significantly correlated with GO phenotypes (32). These observations further confirmed the potential relationship between the DNA methylation level and its functions in the regulation of specific genes involved in the pathogenesis of GO. In another study, Shi et al. (38) carried out genome scale screening of DNA methylation in the peripheral blood samples of patients with GO and healthy controls. They next screened the different candidate genes with a good topological performance by constructing a gene regulatory network of hypermethylated and hypomethylated genes. They further expanded the sample for verification by monomethylated sequencing, thus confirming that, the methylation level of boule homolog (BOLL) was lower, and the methylation level of myelin basic protein (MBP) and CDK5 was markedly higher. BOLL, MBP, and CDK5 genes were found to be associated with increased risk of GO. The possible mechanism can be described by the fact that MBP is related to the activation of MBP specific CD8 lymphocytes and CDK5 could promote oxidative stress. Khong et al. (39) used microarray analysis to detect the gene expression profile of orbital adipose tissue obtained from GO patients. In combination with gene set enrichment analysis (GSEA), they found that compared with inactive GO, the gene enrichment of active GO showed significantly more prominent epigenetic characteristics of acute myeloid leukemia (AML). Moreover, the abnormal hypomethylation of epigenetic markers in AML was associated with immunodeficiency signaling, cytotoxic T cell-mediated apoptosis, and dysregulation of T cell receptor signaling (40). In summary, the pathogenesis of GO is driven by aberrant DNA methylation of specific genes such as ABCD1 and PIK3CB, which may cause oxidative stress and immune dysregulation leading to inflammation (Figure 1).

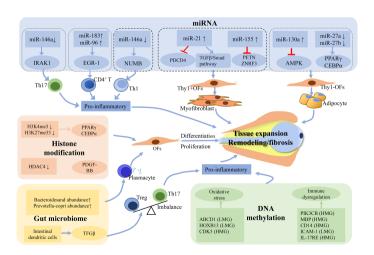


FIGURE 1 | Potential mechanism of epigenetics and gut microbiome in the pathogenesis of GO. Abnormal epigenetic modifications may promote pro-inflammatory cascades and disrupt the expression of signaling molecules that are involved in the fibrogenesis and adipogenesis of orbital fibroblasts. Gut microbiome possibly drives the pathogenesis of GO by influencing secretion of TRAb and Th17/Treg imbalance. The process of all these changes will lead to the hyaluronan production and the activation of orbital fibroblasts, which eventually develop into tissue expansion, fibrosis and inflammation. LMG, lowly methylated genes; HMG, highly methylated genes; TRAb, Thyrotropin receptor antibody; Treg, regulatory T; Th17, T-helper 17.

Furthermore, OFs play a crucial role in the initiation and maintenance of the inflammatory response as well as in orbital tissue expansion and remodeling. Hu et al. (41) performed DNA methylation sequencing and orbital computer tomography (CT) measurement on the peripheral blood samples from patients with GO and healthy controls. They found that the abnormal methylation of MBP gene was significantly correlated with the ratio of the total cross-sectional area of the orbital muscles (OM) to the total orbital area in GO patients. Virakul et al. (42) similarly carried out the detection of various OFs, and further reported that OFs in active versus inactive GO patients might have different proteomic and DNA methylation characteristics, thereby indicating that OFs can show specific pathological characteristics that might be related to the disease stage during the progression of GO. These observations highlighted the potential role of OFs as a future therapeutic target in GO. These findings also suggested that DNA methylation occurs in the fibroblasts and epigenetic understanding of the transition of OFs from "inflammatory/pre-lipogenic" to "remodeling/prefibrotic" effector cells should be a critical area of research in the future studies.

Genetic polymorphisms of a few selected DNA methylation regulatory genes can also lead to abnormal DNA methylation and dysfunction of these genes, which can further increase the susceptibility of the host (43). For example, methylenetetrahydrofolate reductase (MTHFR) reduces 5, 10-methylenetrahydrofolate to 5-methylenetrahydrofol -ate in the folic acid metabolic pathway, thus extending to the methyl transport pathway and indirectly facilitating methyl groups for the DNA methylation and protein methylation through the re-methylation of homocysteine residues. Lee et al. (44) reported that MTHFR polymorphism was closely associated with GO susceptibility and that MTHFR 677 TT can serve as an independent risk factor for GO. Moreover, SNPs 677C>T of the MTHFR gene have been demonstrated to result in pronounced alteration of its enzymatic activity (45), providing indirect evidence for the important role of DNA methylation in GO pathogenesis.

Taken together, DNA methylation plays an important role in the occurrence and development of GO, but the related research is rather limited. Therefore, more studies are needed to explore the key role of DNA methylation in the pathogenesis of GO, to develop biomarkers for the rapid detection of early DNA methylation changes and establish effective GO treatment strategies.

HISTONE MODIFICATION AND GO

The nucleosome constitutes the basic structural unit of chromosomes. It consists of an octamer composed of four different pairs of core histones (H3, H4, H2A, and H2B) and a 147bp length of the double-stranded DNA surrounding the octamer. The core histones moieties are generally spherical, and the non-structural N-terminal amino acids extended by each histone can be modified after translation. The common histone modifications events include methylation, acetylation, phosphorylation, ubiquitination, adenosine diphosphate (ADP)

ribosylation, etc. (46). Currently, histone modification studies have primarily focused on the methylation and acetylation, thereby providing some evidence for the possible role of histone modification in regulating immune tolerance and autoimmune thyroid diseases (47–49).

Histone methylation can be catalyzed by histone methyltransferase (HMT) and occurs mainly on the lysine and arginine residues of H3 and H4 histones, but the transcriptional output from histone methylation primarily depends on the nature of the specific residues involved. Histone H3 methylation is one of the most well characterized epigenetic modifications and actively involved in the adipogenesis of OFs. The differentiation of OFs into adipocytes is primarily a peroxisome proliferator-activated receptor gamma (PPARy) and CCAAT/enhancer binding protein alpha (CEΒPα) dependent process (50). Khong et al. (39) conducted GSEA on the expression profile of the various differentiated genes in the active phase of GO and found that the unmethylated histone H3 in the genes with high CpG density promoters (HCP) found in embryonic fibroblast was significantly downregulated compared with the normal controls. Methylation at K4 (H3K4me3) and K27 (H3K27me33) on histone 3, which is in bivalent status on HCP has been directly related to cell lineage commitment. PPARγ often remains in a bivalent state during embryonic fibroblast differentiation (51). Matsumura et al. (52) reported that the H3K4me3 and H3K27me3 bivalent state might change to H3K4/H3K9me3 during lineage specification. The characteristics of H3K4/H3K9me3 bivalent chromatin signature can possibly maintain $CEBP\alpha$ and $PPAR\gamma$ gene expression at a very low level in lineage-committed mesenchymal stem cells (MSCs) and preadipocytes by silencing or by maintaining developmental genes to poise for subsequent activation, and they may constitute a distinct pseudo heterochromatin/euchromatin boundary which is important for adipocyte differentiation. The above studies suggested that histone H3 methylation might play an important role in the pathogenesis of GO by targeting some genes that are related to adipogenesis. However, at present, there are only few related studies, and there are no reports to establish that abnormal histone methylation might be related to the occurrence of GO. Therefore, additional studies are needed to decipher the specific mechanisms underlying these observations.

Histone acetylation and deacetylation are important components of the gene regulatory machinery. Histone acetylation is usually catalyzed by histone acetyltransferase (HAT), and addition of acetyl groups to the target histones can activate the various transcription factors which can specifically bind to the different DNA binding sites and promote gene transcription. Histone deacetylase (HDAC) can also catalyze the deacetylation of histones to make DNA more tightly encapsulated around the nucleosomes, thus inhibiting gene expression (46). Histone acetylation and deacetylation can effectively maintain a dynamic balance in the nucleus. For example, loss of HDAC expression in the thyroid and immune cells of patients with autoimmune thyroid diseases has been confirmed (53). Recently, Ekronarongchai et al. (31) reported HDAC4 mRNA upregulation and protein expression in OFs by Platelet-Derived Growth Factor-BB (PDGF-BB). It was

observed that the expression of HDAC4 mRNA in the OFs of GO patients was significantly higher as compared to the healthy control group, and the lysine 9 acetylation of histone H3 (H3K9ac) was markedly decreased under PDGF-BB stimulation. It was also found that the expression of hyaluronan synthase 2 (HAS2), collagen type I alpha 1 chain (COL1A1), proliferation marker Ki67 (Ki67), and α-smooth muscle actin (α-SMA) mRNA and the production of hyaluronic acid in PDGF-BB stimulated OFs were significantly decreased upon HDAC4 silencing. These data suggested that HDAC4-induced H3K9 deacetylation could potentially exacerbate GO OFs hyperproliferation and extracellular matrix production by functionally regulating PDGF-BB, thus suggesting that HDAC4 might serve as a new target for GO therapy.

Therefore, histone modification has been related to the occurrence and development of GO. Histone H3 methylation and HDAC may be potential targets for GO therapy, but the specific mechanisms are still not clear. More studies are needed to identify abnormal histone modifications associated with GO and to clarify their molecular mechanisms in GO.

NON-CODING RNAS AND GO

Non-coding RNAs are a class of RNA transcripts that lack the function of coding proteins. They can be classified into miRNAs, circular RNAs (circRNAs) and long noncoding RNAs (lncRNAs), which have been reported to play an important role in the pathogenesis of the various autoimmune thyroid diseases (54).

MiRNAs and GO

MiRNAs are endogenous small non-coding RNA molecules that contain 18 to 23 nucleotides in length, and that by binding to the untranslated 3' domain of the target gene mRNA, miRNAs can effectively mediate the degradation of mRNA or inhibit its translation, thus exerting a strong negative regulatory effect on the gene expression at the post-transcriptional level (55). MiRNAs can regulate more than 60% of protein- coding genes in the humans and have been implicated in the various biological processes such as the lineage commitment in the immune system, cell proliferation, differentiation, apoptosis, and maintenance of immune homeostasis (56, 57). An abnormal expression of miRNAs has been closely related to the pathogenesis of various autoimmune-mediated eye diseases such as GO (58). Circulating miRNAs can be detected in the peripheral blood and orbital tissues of GO patients, with different expression levels in the active disease stage of GO.

Circulating miRNAs and GO

A number of recent studies have shown that some circulating miRNAs might be directly involved in the development of GO, and have been found to be closely related to the regulation of differentiation or activation of immune cells and immune responses.

MiR-146a-5p is one of the most widely studied miRNAs and has an important role in regulating GO progression. MiR-146a is a typical multifunctional miRNA whose expression can be induced

by toll-like receptor 4 (TLR4)/nuclear factor-kappa B (NF-κB) pathway, which acts as an important regulator of immune and inflammatory response pathways (59). By targeting tumor necrosis factor receptor-associated factor 6 (TRAF6) and IL-1 receptor-associated kinase 1 (IRAK1), miR-146a can negatively inhibit the feedback of NF-кВ signaling cascade and modulate the signal transduction by the related receptors and subsequent cytokine release (60, 61). A few studies have shown that the peripheral blood level of miR-146a-5p in GO patients was significantly decreased as compared with their healthy counterparts as well as GD patients which lack GO phenotype (62-64). Additionally, reduced expression of miR-146a-5p has been negatively correlated with the CAS and IL-17 levels in GO patients. IL-17 levels have been positively correlated with CAS, and the downregulation of miR-146a can effectively lead to a paradoxical increase in IL-17 levels possibly due to their inability to inhibit IRAK1. This finding suggested that IRAK1 could play a key regulatory role in the differentiation of Th17 cell (62). The orbital tissues are infiltrated by activated mononuclear cell such as CD4⁺ T cells, which exert a key role on GO activity (7). A number of recent studies have also suggested that the down-regulated expression of miR-146a in CD4⁺T cells of GO patients might facilitate GO development by targeting NUMB genes, thus resulting in the development of inflammatory Th1 response, which can lead to orbital inflammation in GO patients (63, 64). In other words, the downregulation of circulating miR-146a may indirectly promote the development of GO by regulating the expression of various inflammatory mediators. Contrary to the downregulation of miR-146a, miR-155 levels have been reported to be increased in the peripheral venous blood in GO patients. In turn, both miR-155 and miR-146a have multiple target genes in the TLR4/NF-κB pathway, which can mutually regulate the immune response. The upregulation of miR-155 may also promote autoimmune inflammation through the targeted inhibition of cytokine signaling 1 (SOCS1) and SH2 domaincontaining inositol-5-phosphatase 1 (SHIP1) (65). Other miRNAs such as miR-Let7d-5p, miR-96-5p and miR-301a-3p, were also found to be differentially expressed in the serum of GO patients versus control group, which might contribute to the regulated expression of the different inflammatory mediators in vivo and can lead to the occurrence as well as the development of GO (27). In addition, Thiel and colleagues (66) have reported an increased expression of miR-183 and miR-96 in CD4⁺T cells of GO patients and in activated T cells of mice. It has been found that these two kinds of miRNAs promote the activation of CD4⁺T cells by effectively regulating the expression of early growth response 1 (EGR-1), which can directly be related to the activation of PTEN/ Akt signal axis, and thus participate in the occurrence as well as the development of GO. At present, based on the special characteristic of the differential expression of miRNAs, it will be of great significance to study the role of the various influencing factors involved in the pathogenesis and to decipher the mechanisms of the transition from an active stage to static stage, to facilitate early diagnosis and treatment possibilities of GO.

The clinical utility of miRNAs in GO has also been explored. For instance, Shen et al. (67) found that the low serum miR-224-

5p could be correlated with GO glucocorticoid (GC) insensitivity, and overexpression of miR-224-5p in vitro increased GC sensitivity and glucocorticoid receptor protein via targeting glycogen synthase kinase 3 beta. Moreover, the combination of baseline serum miR-224-5p and TRAb was an independent risk factor for GC insensitivity. Recently, a number of studies have analyzed the possible role of miRNAs in the assessment of GO severity and disease outcomes. One study by Martinez-Hernandez et al. (27) reported that miR-let7D-5P expression was markedly decreased in GO patients as compared with GD patients without GO and was also found to be negatively correlated with CAS. Meanwhile, Zhang et al. (68) performed miRNAs profiling of the thyroid tissues in GO patients using next-generation sequencing (NGS) and then identified their serum expression. They found that novel:19_15038 and hsa-miR-27a-3p were up-regulated in GO, while hsa-miR-22-3p was down-regulated in GO as compared to healthy controls. To sum up, the detection of the various differentially expressed miRNAs might be used as the potential biomarkers to predict the development and progression of GD into GO.

Orbital Tissue miRNAs and GO

OFs are both the target cell and effector cell in the pathogenesis process of GO. A number of recent studies have suggested that miRNA inhibitors or mimics can be used to up-regulate or down-regulate miRNAs expression in orbital tissues, thus regulating the corresponding biological behavior. Recent studies have also indicated that some miRNAs might be associated with orbital fibrosis, adipose tissue formation and inflammatory cell development.

For example, miR-146a acts as a key regulatory factor of GO orbital tissue fibrosis. To our knowledge, the level of miR-146a might be variable in the different types of tissues (63, 69). Unlike the downregulation of miR-146a in peripheral blood summarized previously, it is mostly up-regulated in orbital tissues. Wang et al. (69) reported that up-regulation of miR-146a in the orbital connective tissue of GO patients could significantly increase IL-6 level by inhibiting Notch2, thereby effectively promoting mitotic activity of fibroblasts in GO patients, inhibiting cellular apoptosis, and subsequently worsening the deterioration of GO. Woeller et al. (70) reported that TSHR signaling could induce the production of miR-146a and miR-155 on the OFs of GO patients. The expression of the different cell proliferation suppressor genes ZNRF3 and PTEN were substantially decreased, thereby indirectly promoting the proliferation response of OFs, which could possibly explain the pathological mechanisms of partial fibroplasia observed in GO. However, a few studies have also shown that miR-146a can inhibit the fibrosis process. MiR-146a also plays an active role in the antiinflammatory and anti-fibrotic processes of OFs. Jang et al. (71, 72) have reported that inflammatory stress can significantly up-regulate miR-146a expression in OFs, but inhibit the expression of IL-6, ICAM-1, and other inflammatory proteins. It can also negatively regulate levels of transforming growth factor beta (TGF-β) -induced fibrosis markers such as fibronectin (FN), collagen I α , and α -SMA. Type I collagen is considered to be a critical marker of fibrosis, and hyaluronic acid is a glycosaminoglycan. Liu et al. (73) recently

reported that miR-146a expression was markedly down-regulated in the secretion of hyaluronic acid and type I collagen in GO OFs in vitro, which can reduce the aggregation of glycosaminoglycan and collagen deposition, and thereby delay the progression of the disease. Li et al. (65) hypothesized that miR-155, in contrast to miR-146a, can enhance the development of inflammatory T cells by promoting different autoimmune responses. In particular, the levels of miR146a in circulating and orbital tissues have been found to be inconsistent, which may be because the inflammation caused by miR-146 in peripheral blood may not fully reflect the inflammation in orbit. What is more, the contradictory functions about the relationship between miR-146a and differentiation of OFs can also be difficult to interpret, mainly due to the manifestation of the different targeting effects. Furthermore, miR-146a was found to play different functions in the regulation of the diverse signaling pathways or in interactions with the different target mRNAs. However, elucidation of the detailed mechanism of miR-146 in GO requires further investigation.

In addition to miR-146a, miR-21 has also been reported to play an important role in orbital muscle fibrogenesis. Lee et al. (74) found that in human OFs, PDGF-BB can significantly inhibit the expression of Programmed cell death factor 4 (PDCD4) by up-regulating miR-21, and thus promoting the proliferation of OFs. It is consistent with the study that the translation of PDCD4 was inhibited by miR-21 by Young et al. (75). Therefore, PDGF-BB/miRNA-21/PDCD4 pathway might form the basis of novel strategies for developing therapeutic interventions against GO. Tong et al. (76) reported that expression of miR-21-5p in GO eye fibroblasts was markedly higher than that of the control group and that it could enhance TGF-β 1-induced expression of total type I collagen and mRNA level of type I collagen. In addition, anti-miR-21 not only blocked decapentaplegic3 (Smad3) phosphorylation but can also mimic activated Smad3 phosphorylation. These results suggested that miR-21 could substantially enhance Smad3 phosphorylation and activate TGF-β/Smad signaling pathway to participate in orbital muscle fibrogenesis. In summary, these studies indicated that miR-21 may provide new targets for the treatment of GO.

Adipogenesis has been found to be closely related to the degree of ocular protruding (77). Therefore, efforts are being made to find effective ways to block the adipogenesis pathway as a new therapeutic option. Hammond et al. (78) reported that when Thy1- OFs were accumulated in GO, the miR-130a level was significantly increased, resulting in the decreased AMPK activity and promotion of the fat accumulation in orbit. Moreover, the expression level of miR-27a and miR-27b in the orbital adipose tissue of GO patients was significantly lower than that of the controls, which explains the inability to negatively regulate the mRNA expression of the various adipogenic genes such as $PPAR\gamma$ and $CEBP\beta$ in GO fibroblasts, thus indirectly causing the adipogenic differentiation of OFs (79). Therefore, blocking orbital muscle fibrogenesis and adipose generating pathways through modulating miRNA expression could provide novel therapeutic options.

In summary, multiple miRNAs can play an important role in the occurrence as well as the development of GO, and especially miR-146a may serve as an important target (**Table 1**). However,

TABLE 1 | Non-coding RNAs in the pathogenesis of GO.

| | | Expression change | Function | Effects in GO | References | |
|-------------------------|-------------------|-------------------|---|---------------------------------------|------------|--|
| miR-146a | Plasma | Downregulation | Promote the differentiation of Th17 cell | Pro-inflammatory | (62) | |
| | CD4+ T | Downregulation | Increase Th1 response | Pro-inflammatory | (63, 64) | |
| | Orbital tissue | Upregulation | Increase the IL-6 level | Exacerbate GO | (69) | |
| | Orbital tissue | Upregulation | Decreases FN, collagen I α , and α -SMA | Inhibit fibrosis process | (72, 73) | |
| miR-183 and miR-96 | CD4+ T | Upregulation | Contribute to the activation of CD4+ T cells | Pro-inflammatory | (66) | |
| miR-21-5P | Orbital tissue | Upregulation | Promote collagen $I\alpha$ and total collagen production | Promote fibrogenesis | (76) | |
| miR-146a and miR-155 | Orbital tissue | Upregulation | Reducing the expression of PTEN and ZNRF3 | Promote proliferation | (70) | |
| m iR-27a and miR-27b | Orbital tissue | Downregulation | Cause the adipogenic differentiation of OFs | Promote adipogenesis | (79) | |
| m iR-130a | Orbital tissue | Upregulation | Enhance lipid accumulation | Fatty tissue accumulation | (78) | |
| circRNA_14940 | Orbital tissue | Upregulation | Regulate the Wnt signaling pathway, ECM receptor interaction and PIK3-AKT signaling pathway | Participate in the pathogenesis of GO | (80) | |

the therapeutic applications of miRNA have not been fully developed, and hence more studies are needed to clarify the possible effects of miRNAs on other target genes in GO or its pathogenesis. In the future, miRNAs are expected to aid significantly in the precise diagnosis and targeted therapy of GO.

Other Noncoding RNAs and GO

CircRNAs are endogenous entities formed by the covalent binding of 3' and 5' end reverse splicing, which can display important biological functions in acting as miRNA sponge adsorption, attaching to the different RNA-binding proteins, and participating in protein translation (81, 82). However, the role of circRNAs in GO development is still in its infancy. For example, Wu et al. identified (80) 163 differentially expressed circRNAs from the orbital fat/connective tissues of GO patients. Through circRNA-mRNA co-expression and circRNA-mRNA interaction analysis, possible crosstalk of circRNA_14940 with down-regulated mRNA tenascin XB (TNXB) and up-regulated mRNA cyclin D1(CCND1) was analyzed and it was observed that abnormal regulation of the Wnt signaling pathway, PI3K-Akt signaling cascade, and extracellular matrix (ECM) receptor might also be involved in the pathogenesis of GO. CircRNA_10135 can also interact with prostaglandin F receptor (PTGFR) through modulating the calcium signaling pathway and participating in the adipogenesis process of GO. Meanwhile, circRNA_14936 has been found to be correlated with up-regulated mRNA TNF receptor superfamily member 19 (TNFRSF19), and hsa-miR-10392-3p that may potentially affect the occurrence and development of GO by regulating the interactions between circRNA_14936 and TNFRSF19 which can influence B cell survival. Therefore, circRNAs might play an important role in the pathogenesis and progression of GO (Table 1). However, there are only few studies conducted to evaluate the role of circRNAs in the pathogenicity of GO, so the

functions of most circRNAs are not clear and their molecular mechanism remains to be further elucidated.

LncRNAs are an important class of non-coding RNAs with a length of more than 200 nucleotides. LncRNAs can regulate the gene expression through affecting various mechanisms, including those related to epigenetics, transcriptional, post-transcriptional, and miRNAs regulation. A number of previous studies have reported that lncRNAs may play an important role in the pathogenesis of GD (83). For instance, Christensen et al. (84) reported the first lncRNA Heg to be associated with GD and showed that was negatively correlated with monocyte CD14 mRNA and serum TRAb level in GD patients. Moreover, antithyroid therapy could not change lncRNA Heg level in GD patients. The intimate temporal relationship between the onset of GD and GO also clearly suggests that these two diseases may have the same underlying etiology. Xia et al. (85) identified a novel lncRNA named Lnc-Smad3, which can maintain the tight chromatin structure of Smad3 promoter through direct interaction with HDAC1, thereby inhibiting Smad3 transcription and affecting inducible Treg (iTreg) cell polarization through modulating the TGFB/Smad3 signaling pathway.

LncRNAs are also involved in the development of Treg cells, which provides a new direction to further study of the possible relationship between Lnc-Smad3 and GO. Although there are some difficulties in screening and studying lncRNAs, identifying their role and mechanism in GO may be a necessary direction for future research.

GUT MICROBIOME AND GO

The human gut microbiome consists of about 100 trillion different gut microbial cells and serves as the main entry point for bacteria. In a symbiotic manner, the human host primarily provides the gut

microbiota with crucial nutrients for promoting growth. In contrast, the gut microbes play an important role in the food digestion, detoxification, protection from pathogens, and the regulation of the host immune system. In fact, the highest concentration of T cells in the human body is located in the intestinal mucosa, where lymphocytes are primed to respond to the various microorganisms. Furthermore, the intestinal microbiome can also effectively regulate the local intestinal immune system and the systemic immune response (86, 87).

Moreover, multiple human and animal studies have shown that gut microbiome have been closely related to a variety of autoimmune diseases, such as inflammatory colitis (IBD) (88), rheumatoid arthritis (RA) (89), type I diabetes (T1D) (90), and systemic lupus erythematosus (SLE) (91). For instance, Jiang et al. (92) sequenced the fecal samples of patients with GD and healthy controls, and found that the richness of intestinal microflora in patients with GD decreased significantly. This alteration was characterized by a significant increase in Bacteroides, which damaged intestinal barrier function, resulting in the release of a large number of pro-inflammatory factors outside the intestine, thus resulting in immune dysfunction. In addition, it was also found that changes in thyroid hormone levels can lead to an increase in the number of Lactobacillus. It may play a harmful role in GD by activating NF- κB signaling pathway and autoimmune response to thyroid gland. At present, there are only few studies on intestinal flora in GO. Several reports have suggested that there is a marked decrease in bacterial community diversity in GO. Shi et al. (93) sequenced the fecal samples of patients with GO and healthy controls and found that compared with the control group, the proportion of Bacteroidetes and Prevotellaceae in patients with GO at different levels increased significantly, while the proportion of Blautia and Fusicatenibacter decreased significantly. The proportion of the Succinivibrionaceae and Subdoligranulum was positively correlated with TRAb, and there was a negative correlation between Parabacteroides_disasonis and TRAb. Further analysis showed that Prevotellaceae could clearly distinguish GO patients from the control group. The level of *s_Prevotea_copri* was reported to be positively correlated with the level of serum TRAb, which may be related to active orbital inflammation (94). Thereafter, the researchers conducted a comparative analysis of GD, GO and healthy people for the first time. They found that the proportion of Deinococcus-Thermus and Chloroflexi in patients with GO was significantly lower than that in GD (95). Interestingly, after transplanting the normal human feces into type 2 diabetic mice, researchers found that the abundance of Bacteroides and some bacteria decreased and blood glucose levels recovered. This led to the speculation that it may be possibly increasing the diversity of intestinal microflora that can affect the different types of microorganisms in the intestines of mice to achieve the effect of treatment (96). Therefore, whether fecal microbiome transplantation can also have a therapeutic effect on GO can be an important direction for future research. Additional in-depth studies about the role of intestinal microorganisms in GO will possibly provide new methods and ideas for the treatment of GO. In addition, some studies have reported that Salmonella that can cause enteritis possesses an antigenic cluster similar to the thyroid stimulating hormone receptor protein, where antibodies against

the thyrotropin receptor proteins might be produced after infection (96). These results laid a sound foundation for exploring the interaction between intestinal microorganisms and TRAb. However, the current findings may also be related to thyroid autoimmunity rather than GO, and might not be entirely specific for GO.

Masetti et al. (34) used intramuscular injection of eukaryotic expression plasmid (thyroid stimulating receptor) in two distinct experimental centers in two different countries (Center 1 and 2). The GD/GO disease model of BALB/C female mice was established by immunological method of TSHR. The results showed that only BALB/C mice from Center 2 displayed hyperthyroidism and apparent changes in orbital tissues. Interestingly, it was found that the abundance and species composition of intestinal flora were significantly different between BALB/C mice tested from the two centers by 16S rRNA gene sequencing and traditional microbial culture. In Center 2, the intestinal flora structure of different intervention groups (TSHR group, β Gal group) was analyzed by 16S rRNA gene sequencing and it was found that the intestinal flora of TSHR group changed significantly (increased relative abundance of Lactobacillus, decreased relative abundance of bacteroides, etc.). These alterations were also associated with some clinical features such as thyroxine levels (fT4), TRAb, orbital adipogenesis, and muscular atrophy. The same group constructed another GD/GO model by using BALB/c and C57BL/6J mice. The results indicated that the C57BL/6 mice had higher TRAb, but compared with BALB/c mice, the spleen T cells of C57BL/6 mice exhibited poor proliferation of TSHR response, and secreted mainly anti-inflammatory cytokines such as IL-10, without elevated thyroxine levels or any orbital lesions, while resembling immunized BALB/C mice showed orbital pathology. These observations suggested that there was a different pattern of correlation between the microbiomes and disease (endocrine and immune) characteristics between the two different mouse strains (97). In addition, Shi et al. (94) found that operational taxonomic units (OTUs) derived from the bacteroidesstercoris species were markedly associated with the thyroglobulin autoantibodies, whereas OTUs from the bacteroides genus were associated with CAS. Masetti et al. (98) deciphered that the effect of intestinal microorganisms on the immune system can be primarily revealed by maintaining a balance between the Tregs and pro-inflammatory Th17 cells and Th1 cells. When the microbiome might not be ideal, intestinal dendritic cells can secrete TGF-β, which can cause the Th17/Treg imbalance, thus leading to the loss of immune tolerance. In individuals with a particular genetic predisposition, the result can potentially lead to GD and GO. These results clearly suggested that the gut microbiome could play a critical role in shaping the immune response that might significantly influence the TRAb levels and immune tolerance, thereby influencing thyroid and eye disease phenotypes.

The above studies have provided a novel perspective research pathway to clarify the potential mechanisms of intestinal microbiome disorders observed in GO. Hence, understanding that how the composition of the gut bacterial microbiome and the abundance of specific bacterial strains might affect

the autoimmune response in GD and GO remains the next important aspect.

CONCLUSION

With the emergence of novel bioinformatics tools and high-throughput gene sequencing technologies, the important role of epigenetics and gut microbiomes in GO has been gradually elucidated (**Figure 1**). Still, several crucial mechanisms especially from GD to GO remain unclear. At present, the study of GO epigenetics focuses primarily on only one kind of epigenetic modifications, but the role of its regulatory network in controlling molecular mechanisms of GO remains unknown. Therefore, there is an unmet need to conduct a combined analysis of the varying epigenetic modifications that can further explore the possible roles of various less explored epigenetic factors that have been implicated in the pathogenesis of GO.

Epigenetic inheritance can link environmental factors and diseases. How environmental factors can lead to the progression of GO by facilitating epigenetic inheritance and intestinal microbial dysregulation in GO remains to be elucidated. To date, GO is clinically diagnosed by an ophthalmologist with an adjunct imaging such as CT or magnetic resonance imaging (MRI). A few studies have shown that some epigenetic markers and intestinal microflora might be different in GD patients with or without GO. However, it is still difficult to predict which GD patients will develop into GO or severe GO, and which GD

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patients will be responsive or unresponsive to the different types of treatment. Further studies should focus on the current challenges in the future.

AUTHOR CONTRIBUTIONS

YW and H-SY participated in the conceptualization and writing. X-MM, XW, XS, L-JW, X-QL, and X-YL participated in the review and editing. All authors contributed to the article and approved the submitted version.

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Therapy With Different Dose Regimens of Rituximab in Patients With Active Moderate-To-Severe Graves' Orbitopathy

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Campi I, Vannucchi G, Muller I, Lazzaroni E, Currò N, Dainese M, Montacchini B, Covelli D, Guastella C, Pignataro L, Fugazzola L, Arosio M and Salvi M (2022) Therapy With Different Dose Regimens of Rituximab in Patients With Active Moderate-To-Severe Graves' Orbitopathy. Front. Endocrinol. 12:790246. doi: 10.3389/fendo.2021.790246 **Background:** Immunosuppressive therapy of Graves' orbitopathy (GO) is indicated during the active phase of disease. Intravenous steroids (IVGC) are effective in about 70% of patients, although unresponsiveness or relapse are observed. In previous studies, rituximab (RTX) has been shown to be effective in inactivating moderate-to-severe GO when used early in the disease, but its optimal dosage has never been studied in randomized clinical trials. Aim of this study was to compare the efficacy and safety of different doses of RTX, based on a *post-hoc* analysis of two open label studies and one prospective trial randomized to IVGC.

Methods: of 40 patients (35 women, 5 men), with active moderate-to-severe GO treated with RTX, 14 received a single dose of 100 mg (Group 1), 15 a single dose of 500 mg (Group 2) and 11 two 1000 mg doses, administered one week apart (Group 3). Thyroid function, TSH-receptor antibodies (TRAb) and peripheral CD19+ cells were measured. Primary endpoint was disease inactivation, measured as a decrease of the Clinical Activity Score (CAS) of at least two points. Secondary endpoints were improvement of proptosis, diplopia, quality of life and safety.

Results: Baseline CAS decreased significantly in all groups (P<0.0001), independently of GO duration or whether patients had newly occurring or relapsing GO after IVGC. Proptosis did not significantly change. There was an inverse correlation between the Gorman score for diplopia and RTX dose (P<0.01). The appearance score of the GO-QoL improved in Group 1 (P=0.015), and the visual function score, in Group 2 (P=0.04). A reduction of serum TRAb was observed in Group 1 (P=0.002) and Group 2 (P<0.0002), but not in Group 3. CD19+cell decreased in all groups (P<0.01), independently of the dose.

Conclusions: We studied the optimal dosage of RTX in the treatment of active moderate-to-severe GO. In this analysis, we considered the efficacy of RTX in inactivating GO, in

changing its natural course, its effect on disease severity and on the patients' quality of life. Based on our clinical findings, and balancing the cost of therapy, a single 500 mg dose regimen is suggested in the majority of patients.

Keywords: Graves' disease, Graves' orbitopathy, Rituximab, B lymphocytes, TSH-receptor antibodies

INTRODUCTION

Graves' disease (GD) is a thyroid autoimmune disorder in which anti-thyrotropin (TSH) receptor autoantibodies (TRAb) with stimulating activity induce hyperthyroidism. Graves' orbitopathy (GO), characterized by inflammation and remodelling of orbital tissues, is the main extra-thyroidal manifestation of GD. Moderate-to-severe GO determines major facial disfigurement, psychosocial impact and, rarely progression to sight loss (1, 2). Typically, GO has a diphasic course (3) with an inflammatory and progressive phase, followed by stabilization and subsequent scarring due to tissue fibrosis. Medical treatment, based on immunosuppression, is indicated during the active phase of GO. Intravenous steroids have been shown to be effective in about 70% of patients, although unresponsiveness and disease relapse are observed in a good proportion of patients (20-30%) (4).

Rituximab (RTX) is a human/mouse chimeric monoclonal antibody that binds the CD20 antigen on the surface of B cells and induces rapid depletion of B cells in both periphery and lymphoid organs. Clinically approved indications for the use of RTX in autoimmune disease are currently rheumatoid arthritis and ANCA-related vasculitis (5). In previous studies (6), RTX has been shown to be effective in inactivating moderate-to-severe GO when used early in the disease, but may have little effect in longer duration disease (7). The clinical effect of RTX in GO is quite rapid, since disease generally inactivates within 4 – 6 weeks after the first infusion (8). RTX induces B-cell lysis and may activate anaphylatoxins and other inflammatory cytokines with subsequent recruitment of phagocytes within the orbital tissue and depletion of inflammatory cells yielding tissue fibrosis (9, 10). This may explain why, after B cells repopulation, reactivation of GO was only observed rarely.

RTX therapy is not devoid of side effects: infections may arise due to hypogammaglobulinemia, although they are also commonly associated to therapy with steroids or other immunosuppressive drugs (11). Acute infusion reactions related to the release of cytokines produced by macrophages, monocytes, lymphocytes and NK cells, can be observed in 10-30% of patients at the first infusion (12). These effects may be associated to larger doses of RTX and significant lymphocytic infiltrates in target organs (13).

The optimal dosage of RTX in GO has never been studied in randomized, dose finding clinical trials. Standard dosing in rheumatoid arthritis and other autoimmune diseases consists of two i.v. administrations of 1000 mg with a 2-week interval. This dose was also initially used in most open studies (8) and in individual case reports (10) in patients with GO. Rapid and complete B lymphocyte depletion was shown to occur already

after very low dosages (25-100 mg) of i.v. RTX (9). In a prospective randomized clinical study it was shown that the efficacy of a single 500 mg RTX dose was comparable to two 1000 mg doses (7) in inactivating GO and more effective than i.v. methylprednisolone (6). Later observation that in GO total peripheral (and orbital) B cell depletion was occurring even after low doses of RTX (100 mg) (9), has allowed us to reduce the therapeutic dose to a single infusion of 100 mg RTX (14).

Aim of the present study was to compare the efficacy and safety of different doses of RTX, employed prospectively in two open label studies (14, 15) and in one trial in which RTX was randomized to i.v. steroid therapy (6), in active GO.

PATIENTS AND METHODS

Patients

Forty patients, 35 women and 5 men, 32-81 years of age (mean \pm SE 58.7 ± 2.7 yr), with active moderate to severe GO were grouped based on the RTX dose that they received (Table 1). Of these patients, 14 were treated with a single dose of 100 mg (Group 1), 15 with a single dose of 500 mg (Group 2) and 11 with two 1000 mg doses of RTX, administered one week apart (Group 3). Of the 40 patients included in the study, 34 had GD, 11 in group 1, 12 in group 2 and 11 in group 3, five had Hashimoto's thyroiditis, of whom three in Group 1 and two in Group 2, and one had euthyroid GO (Group 2). Thirty of 34 GD patients were euthyroid or subclinical hypothyroid because slightly overtreated with methimazole and four slightly hyperthyroid at the time of treatment. Nineteen patients were smokers, not significantly distributed in the three groups (Table 1). All patients had mean GO duration of 11 ± 3 months. Nineteen patients were previously treated with corticosteroid therapy, nine in Group 1, six in Group 2 and four in Group 3. Glucocorticoids were discontinued at least three months before RTX administration. Patients were followed-up to 76-weeks follow-up to assess the need of rehabilitative surgery.

RTX Administration Schedule

RTX dosing and treatment schedule were adapted based on the experience achieved with this drug throughout the years. The dose of RTX initially employed in the treatment of active moderate-to-severe GO was based on previous work in autoimmunity (6, 16, 17) and consisted of a single dose of 100 mg and 500 mg (Group 1 and Group 2, respectively) or 1000 mg twice at two weeks interval (Group 3).

All patients, one hour prior to RTX infusion, received oral paracetamol (1 gr), chlorphenamine (10 mg) and i.v. hydrocortisone (100 mg) to prevent possible infusion reactions.

TABLE 1 | Baseline clinical characteristics of patients with moderate-to-severe GO treated with different doses of RTX.

| Group (dose of RTX) | GROUP 1 (100 mg) | GROUP 2 (500 mg) | GROUP 3 (1000 mg x2) | P |
|---|------------------|------------------|----------------------|--------|
| Number of patients | 14 | 15 | 11 | |
| Age * (years) | 55.6 ± 3.3 | 56.9 ± 3.1 | 63.5 ± 1.6 | 0.09 |
| Gender (F/M) | 12/2 | 14/1 | 9/2 | 0.67 |
| Smoker (Yes/No) | 5/9 | 8/7 | 6/5 | 0.55 |
| FT4 (pmol/L) * | 11.9 ± 1.0 | 11.7 ± 1.0 | 9.8 ± 1.4 | 0.15 |
| TRAb (mU/mL) * | 27.6 ± 13.8 | 12.7 ± 3.0 | 11.6 ± 4.1 | 0.93 |
| GO duration (months) * | 4.9 ± 1.6 | 6.3 ± 1.5 | 10.7 ± 3.0 | 0.14 |
| New onset/relapse (%) | 4/10 (30) | 9/6 (60) | 7/4 (64) | 0.14 |
| CAS * | 4.6 ± 0.3 | 4.3 ± 0.2 | 4.4 ± 0.3 | 0.73 |
| Proptosis right eye (mm) * | 23.8 ± 0.7 | 23.0 ± 0.6 | 21.4 ± 0.9 | 0.13 |
| Proptosis left eye (mm) * | 23.6 ± 1.0 | 22.6 ± 0.7 | 21.3 ± 0.9 | 0.21 |
| Gorman score N (0/1/2/3) | 4/4/2/4 | 3/4/7/1 | 4/2/4/1 | 0.45 |
| CD19 ⁺ cells * (cells/mm ³) | 293.5 ± 24.7 | 236.5 ± 31.2 | 269.7 ± 50.4 | 0.21 |
| GO-QoL Appearance * (%) | 56.6 ± 6.3 | 61.0 ± 6.7 | 69.0 ± 13.0 | 0.69 |
| GO-QoL Function *(%) | 56.4 ± 9.7 | 42.3 ± 7.3 | 77.0 ± 14.6 | 0.18 |
| Thyroid status | 11/1/2 | 12/2/1 | 5/1/5 | 0.15 |
| (0 = Euthyroid, 1 = Hyperthyroid, | | | | |
| 2 = Hypothyroid) | | | | |
| Thyroid diagnosis N (GD/HT/EGO) | 11/3/0 | 12/2/1 | 11/0/0 | 0.66** |
| Thyroid disease duration | 33.9 ± 9.3 | 44.1 ± 14.9 | 50.8 ± 35.6 | 0.68 |
| N of GD patients treated with RAI/TX before RTX (on L-T4) | 4 (3) | 2 (2) | 2 (1) | 0.58 |

^{*}All values are expressed as mean ± SE.

CAS, clinical activity score; EGO, euthyroid Graves' orbitopathy; GD, Graves' disease; GO, Graves' orbitopathy; HT, Hashimoto' thyroiditis; L-T4, levothyroxine; QOL, quality of life; RAI, radioiodine treatment; RTX, rituximab; TX, total thyroidectomy.

Patients were defined as euthyroid (0) hyperthyroid (1) or hypothyroid (2) in case of normal, elevated or low FT3 and/or FT4, independently on TSH serum levels.

Patients with severe chronic diseases, ongoing infections or neoplastic diseases were excluded from treatment. RTX was also not administered in patents with known significant coronary artery disease, cardiac arrhythmias, congestive heart failure, active infection, primary or secondary immunodeficiency, history of hypersensitivity, known anaphylaxis to mouse-derived proteins, positive Purified Protein Derivative (PPD) test without documentation of treatment for tuberculosis (TB) infection and denied consent to HIV testing. During treatment and follow-up, complete blood count, serum glucose, aminotransferases and gamma glutamyltransferase were closely monitored. Side effects were classified as major (diabetes mellitus, depressive syndrome, increase of aminotransferases levels 5 times or above the upper normal limit) and minor (dyspepsia, laryngeal itching, rhinorrhea, insomnia).

Biochemical Analysis

Serum free-thyroxine (FT4), free-triiodothyronine (FT3) and TSH concentrations were measured using an electrochemiluminescent immunoassay (ECLIA, Roche Diagnostics) and normal ranges were 8-17 pg/ml, 2-5 pg/ml and 0.26-5.2 mU/L, respectively. Serum TRAb were measured as TSH binding inhibitory immunoglobulins (TBII), using a 2nd generation TRAK human lumitest (Thermofisher, AG, Henningsdorf/Berlin, Germany; n.v.<1.5 U/L). Lymphocyte subpopulations were measured at baseline and at each follow-up examination.

Cytofluorimetric Analysis

The pattern of peripheral blood lymphocytes was studied before RTX and subsequently during the study period and the follow-up. We tested the standard immunophenotypic panel (CD3+,

CD3+4+, CD3+8+, CD3+DR+, CD20+, CD19+5+, CD56+16+3) on aliquots of around 10⁵ lymphocytes, submitted to triple staining procedures for immunogating with CD45, and the pairs of monoclonal antibodies to subpopulations of T, B and NK cells, subsequently processed in the flow cytometer (BD Facsan, Cell-quest software).

Study Endpoints and Clinical Assessment

Primary endpoint of the study was GO inactivation, defined as a reduction of 2 points of CAS, or a CAS of 3/10 or less, at 12 and 24 weeks after treatment with any RTX dose, when compared to baseline. The analysis of efficacy was also carried out after stratifying patients for disease duration of more or less of six months at the time of intervention.

Secondary endpoints at 24 weeks were: 1) changes of the severity of GO: significant improvement was considered a 2 mm reduction of proptosis, and of at least one class of the Gorman's score for diplopia (18); 2) improvement of the quality of life (QoL), assessed with a specific and validated questionnaire for GO (GO-QoL) (19): an increase of at least 6 points of the score of the questionnaire at 24 weeks was considered significant; 3) decrease of serum levels of TRAb; 4) decrease of peripheral B-cell lymphocytes; 5) occurrence of any major adverse events.

Statistical Analysis

This was a *post-hoc* analysis of three prospective trials, one randomized to intravenous steroids and two open label. All values are expressed as mean ± standard error (SE) or ± standard deviation (SD), as specified. Analysis by Fischer exact test, Wilcoxon and Mann-Whitney test were applied, as appropriate, and performed using SPSS 8.0 for Windows.

^{**}in the Chi-Square analysis HT and EGO were added together.

Statistical significance was defined as P <0.05. An ANOVA or Friedman model was used to study the changes of the CAS values during treatment and follow-up. A *per protocol* analysis of the data was carried out and patients undergoing surgical orbital decompression during the observation period up to 24 weeks, were not included in the analysis of the disease outcomes.

RESULTS

Disease Inactivation and Relapse

There were no significant differences in baseline clinical and immunological features in the patients of the three groups of the study (**Table 1**).

In Group 1, baseline CAS value was 4.6 ± 0.3 and decreased to 2.1 ± 0.4 at 12 weeks (P<0.05) and to 1.1 ± 0.2 at 24 weeks (P<0.0001). In Group 2, baseline CAS was 4.3 ± 0.2 and decreased to 1.4 ± 0.4 (P<0.001) and to 0.5 ± 0.3 (P<0.0001) at 12 and 24 weeks, respectively. In Group 3, baseline CAS was 4.4 ± 0.3 , 1.8 ± 0.4 (P<0.05) and 0.7 ± 0.2 (P<0.0001) at 12 and 24 weeks, respectively (**Table 2**).

There was no difference in the CAS reduction at 12 weeks in the three Groups of patients (2.6 ± 0.4 in Group 1, 2.9 ± 0.4 in Group 2, 2.6 ± 0.4 points in Group 3; P=N.S). No difference was also observed at 24 weeks (3.6 ± 0.3 in Group 1, 3.8 ± 0.3 in Group 2 and 3.6 ± 0.4 in Group 3; P=N.S; **Figure 1**). GO inactivation was not significantly associated to disease duration (more or less than 6 months) independently of the dose employed (not shown, P=N.S). Response to RTX therapy was not different whether patients had newly occurring GO or relapse after previous steroid therapy (not shown, P=N.S.).

No patient showed disease reactivation at 24 weeks and during follow-up to 76 weeks. Two patients treated with 100 mg RTX, despite an initial response to treatment, eventually developed dysthyroid optic neuropathy (DON) and required prompt surgical orbital decompression.

Disease Severity and Quality of Life

Mean proptosis values did not significantly change at 24 weeks in any of the 3 groups after RTX (**Table 3**).

Patients receiving 500 mg of RTX tended to have more improved or stable diplopia (Chi-square P<0.06; **Figure 2**), and this finding was confirmed by the observation of a significant inverse correlation of the Gorman score with the dose of RTX (r -0.43, P< 0.01).

In Group 1 the appearance score of the GO-QoL questionnaire improved in 6/14 patients and the visual function score in 4/14 (**Table 3**), with significant increase of the overall appearance score (P=0.015, **Figure 3A**), but not of the function score (**Figure 3B**). In Group 2 improvement of the appearance score was observed in 7/15 patients and in one of the visual function score, although improvement was significant only for the overall visual function score (P=0.04, **Figure 3B**), and not of the appearance score (**Figure 3A**). In Group 3 analysis of the overall score was not performed due to the low number of patients who completed the questionnaire.

Thyroid Function and Serum TRAb

All but one patients at 24 weeks were euthyroid (P=0.04, not shown). Detailed data on baseline treatments for the thyroid diseases are reported in **Table 1**. A significant reduction of serum TRAb levels at 24 weeks was observed after RTX in Group 1 (P=0.002) and Group 2 (P<0.0002), but not in Group 3 patients (P=NS; **Figure 4**).

Lymphocytes Depletion

Peripheral lymphocyte subpopulations were studied at 24 weeks in order to assess how different RTX doses affect CD19+ cell depletion. A significant reduction in CD19 cell numbers was observed in all three groups (P< 0.01), independently of the RTX dose employed (**Figure 5**). At 24 weeks, the number of CD 19+ cells in patients of Group 1 were 75.1 \pm 15.3, significantly higher than in patients of Group 2 (27.1 \pm 7.1; P=0.05) Group 3 (7.5 \pm 4.3; P=0.0001), while the difference is not statistically significant between patients of Group 2 and Group 3 (**Table 3**).

Overall, rehabilitative surgery (evaluated at 76 weeks post-treatment with RTX) was performed in 20 patients. Ten patients underwent orbital decompressions, of whom eight elective and two emergency (DON), six patients strabismus surgery and four eyelid surgery. The distribution of surgical procedures among patients was not related to the RTX dose employed.

We did not observe any significant changes in the CD4+ or CD8+ T lymphocytes levels following RTX infusion, in any Groups (**Supplementary Figure 1**). Interestingly, we observed a slightly increase in the number of natural killer cells (NK) in Group 1 and 2 after RTX. This difference reached the statistical significance only at 8 weeks in Group 1 and at 32 and 40 weeks in Group 2 compared to baseline values. On the contrary, the number of NK cells remained stable in Group 3 (**Supplementary Figure 1**).

TABLE 2 | Clinical activity score (CAS) at baseline and 12 and 24 weeks after treatment with different doses of RTX.

| | Baseline 12 weeks | | P* 24 weeks | | P*0 vs 24 weeks | P *12 vs 24 weeks |
|------------|-------------------|---------------|-------------|---------------|-----------------|-------------------|
| | Daseille | 12 Weeks | | 24 WEEKS | F 0 VS 24 WEEKS | F 12 V3 24 WEERS |
| GROUP 1 | 4.6 ± 0.3 | 2.1 ± 0.4 | < 0.05 | 1.1 ± 0.2 | <0.0001 | NS |
| 100 mg | | | | | | |
| GROUP 2 | 4.3 ± 0.2 | 1.4 ± 0.4 | < 0.001 | 0.5 ± 0.3 | <0.0001 | NS |
| 500 mg | | | | | | |
| GROUP 3 | 4.4 ± 0.3 | 1.8 ± 0.4 | < 0.05 | 0.7 ± 0.2 | <0.0001 | NS |
| 1000 mg X2 | | | | | | |

*Friedman TEST.

All values are expressed as mean \pm SE. NS, not significant.

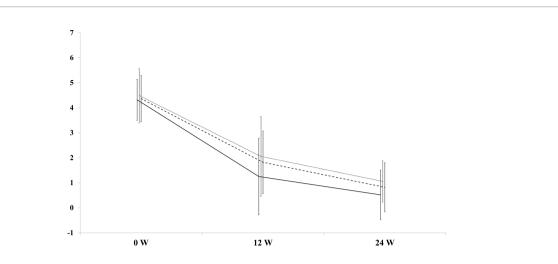


FIGURE 1 | Decrease of the Clinical Activity Score (CAS) at 12 and 24 weeks after rituximab (RTX) in patients with Graves' orbitopathy (GO). Dotted line= Group 1 (RTX 100 mg); solid line=Group 2 (RTX 500 mg); long dash line=Group 3 (RTX 1000 mg x 2).

Adverse Events

Minor adverse reactions such as itching of the throats and stuffiness of nose with rhinorrhea, were experienced by many patients. In these cases, resolution of the symptoms occurred after slowing down the rate of RTX infusion. In two patients of Group 1 a major adverse event known as the syndrome of release of cytokines occurred early during RTX infusion. This was characterized by orbital edema and progressive decline of vision rapidly occurring 30 minutes after the beginning of the infusion. This reaction was controlled by the administration of 100 mg of i.v. hydrocortisone with complete recovery of vision after 3 hours (9). Two patients treated with 100 mg RTX developed DON, possibly present at a subclinical stage at the time of treatment. This event is to be considered treatment failure, rather than an adverse event.

Cost of Treatment

The cost of RTX therapy depends the dose employed. The actual cost of one single 100 mg dose is of € 338, that of one 500 mg

dose is \in 1,698 and that of a two-dose cycle of 1000 mg is \in 2.154, as released by the manufacturer in Europe (Roche Ltd.)

DISCUSSION

The first and most important finding of this study is that doses of RTX ranging from 100 to 2000 mg invariably induce inactivation of moderate-to-severe GO with only some differences in the clinical response, especially on diplopia, quality of life assessment and CD 19+ cells repopulation. This study is based on a *post-hoc* analysis of data collected from one prospective RCT and two open label studies that have employed different protocols and dosing of RTX administration.

Our findings show that any of the three RTX dosages used (100 mg, 500 mg and 1000 mg x 2) is equally effective in inactivating the disease in >95% of patients at 12 and 24 weeks after infusion. Efficacy of RTX has been known to be associated to the duration of B cell depletion in the circulation (14). In this

TABLE 3 | Clinical outcome of patients with moderate severe GO assessed 24 weeks after therapy with different doses of RTX.

| | GROUP 1 (100 mg) | GROUP 2 (500 mg) | GROUP 3 (1000 mg x2) | P * |
|--|------------------|------------------|----------------------|---|
| Number of patient | 14 | 15 | 11 | |
| TRAb (mU/L) | 14.9 ± 7.5 | 5.98 ± 1.98 | 10.5 ± 4.0 | 0.48 |
| CAS | 1.07 ± 0.2 | 0.5 ± 0.3 | 0.7 ± 0.2 | 0.14 |
| Proptosis right eye (mm) | 22.6 ± 0.8 | 22.97 ± 0.7 | 22.05 ± 1.4 | 0.73 |
| Proptosis left eye (mm) | 23.0 ± 0.9 | 22.7 ± 0.6 | 21.5 ± 1.1 | 0.58 |
| Δ Proptosis right eye (mm)** | -1.2 ± 0.4 | -0.08 ± 0.3 | 0.64 ± 1.1 | 0.08 |
| Δ Proptosis left eye (mm)** | -0.6 ± 0.6 | 0.10 ± 0.2 | 0.2 ± 0.5 | 0.50 |
| GO-QoL Appearance (%) | 71.6 ± 4.6 | 59.8 ± 6.4 | 83.2 ± 5.97 | 0.18 |
| GO-QoL Functions (%) | 52.0 ± 9.8 | 58.9 ± 9.2 | 72.3 ± 16.8 | 0.66 |
| CD19 ⁺ cells (cells/mm ³) | 75.1 ± 15.3 | 27.1 ± 7.1 | 7.5 ± 4.3 | G1 vs G2 P<0.05; G1 vs G3 P<0.0001; G2 vs G3 P = NS |

All values are expressed as mean ± SE

^{*}Mann-Whitney test

^{**}Difference in between proptosis at 0 and 24 weeks.

GO, Graves' orbitopathy; CAS, clinical activity score; GO-QoL, quality of life questionnaire; NS, not significant.

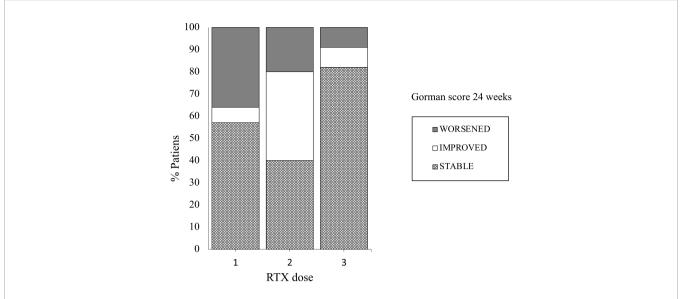


FIGURE 2 | Proportions of patients with GO with modifications of the Gorman score for diplopia after treatment with three different doses of RTX: 1= Group 1 (RTX 100 mg); 2= Group 2 (RTX 500 mg); 3 Group 3 (RTX 1000 mg x 2). Chi Square test.

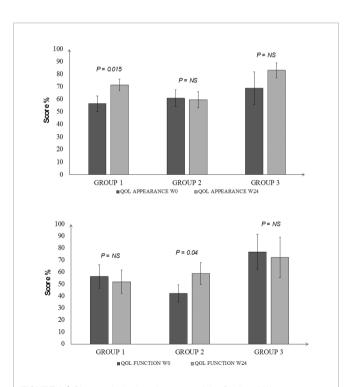


FIGURE 3 | Changes in the baseline score of the Quality of Life assessment of patients with GO at 24 weeks after treatment with different doses of RTX. Panel **(A)** (upper): appearance score, Panel **(B)** (lower): function score. Group 1 = 100 mg; Group 2 = 500 mg; group $3 = 1000 \text{ mg} \times 2$; baseline values are in dark grey and values at 24 weeks in light grey. Wilcoxon matched pair test.

study we show that peripheral CD19+ cells begin repopulating only slightly earlier than 24 weeks in patients treated with the very low dose (100 mg), but later in those administered either 500 or

2000 mg. B cell return in the circulation, though, does not modify the disease outcome or favor disease relapse during follow-up (up to 12 months) (6, 14, 20). Based on these observations, we believe that in moderate-severe GO lower doses of RTX, which are generally burdened by a lower risk of adverse events such as reactivation of infections (12) or the induction of other autoimmune processes after depletion of B cells (21), should be preferred.

We studied whether RTX induced inactivation may depend on GO duration of less or more than 6 months, as previous studies have suggested that in GO of longer duration RTX might be less effective (7, 17). We did not find differences in the therapeutic response based on the duration of disease with any of the doses employed. We suggest that efficacy of RTX treatment is likely more related to the early inflammatory phase of GO, than to disease duration. We observed no difference in the response of patients with newly diagnosed GO or of those with relapsing disease after a complete cycle of intravenous steroids, who invariably had longer disease duration (up to 66 months). This findings, besides confirming that disease activity rather than duration is important for the effect of RTX in GO, also suggest that this modality of therapy can be proposed as second line treatment when steroids fail (1, 2).

The therapeutic outcome on parameters of disease severity, especially those related to muscle involvement, appear to be of importance, as the presence of diplopia in inactive GO has an impact on the quality of life of patients. While there were no significant changes in the degree of proptosis with any of the three dosages used, we did observe a significant improvement of the Gorman score for diplopia when higher RTX doses (Group 2 and Group 3) were employed. A possible explanation for this is that higher doses might be more effective than lower doses in preventing the orbital tissue fibrosis that follows the active disease phase.

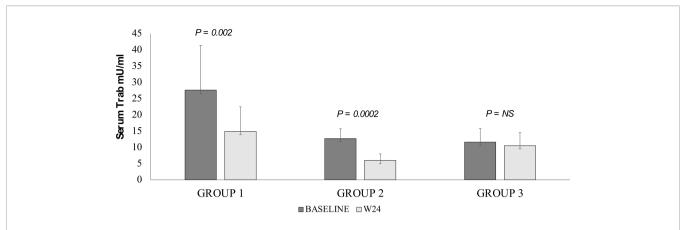


FIGURE 4 | Changes of serum TRH receptor antibody levels at 24 weeks in patients with GO treated with different doses of RTX. Group 1 = 100 mg; Group 2 = 500 mg; group 3 = 1000 mg x 2. Baseline values are in dark grey and values at 24 weeks in light grey. Paired T test.

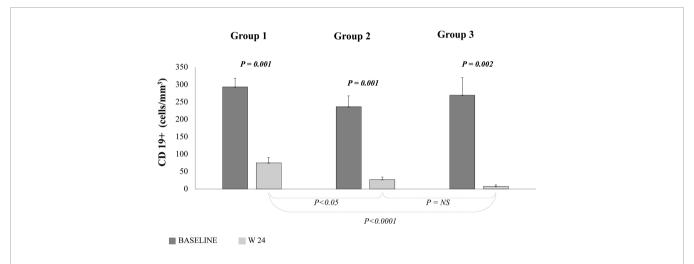


FIGURE 5 | Peripheral B cell count measured at baseline (dark grey) and 24 weeks (light grey) after treatment with different doses of RTX. Group 1 = 100 mg; Group 2 = 500 mg; Group 3 = 1000 mg x 2. Paired T test.

The QoL questionnaire used in the study has been validated and is specific for assessing quality of life changes in GO patients (18). It is divided into two sections, the first one assessing the impact of the aesthetic changes (appearance) on patients, the second one that of the eye dysfunction. The perception of self-appearance was found to improve at 24 weeks only in patients treated with 100 mg RTX in a single administration, while improvement of function was reported by those treated with a single larger dose of 500 mg. Improvement of inflammatory signs were in fact more rapidly observed after treatment with the low dose (>90% of patients inactivated at 12 weeks), whereas the impact of RTX on diplopia was found to be associated to the use of larger doses. This suggests that high doses of RTX might be preferable in patients with significant extraocular muscle involvement, as it would reduce the need of rehabilitative surgery. Unfortunately, we could not assess QoL outcome in Group 3 because of the small number of questionnaires returned, and this is a limitation of this study. Differences in the quality of life assessment were not associated to differences in the number of rehabilitative surgeries required, whichever dose of RTX was employed.

In two patients treated with the low dose, RTX therapy failed to inactivate GO and to prevent progression to DON. These patients may have had a subclinical form of DON at the time of treatment which might have been further exacerbated by the increase of intraorbital edema induced by an acute release of cytokines mediated by RTX as previously reported (9). In other studies RTX has been used in patients with DON (10) resulting in improvement of vision. In addition, the observation of the syndrome cytokine release in two patients receiving only 100 mg RTX, suggests that this adverse event is not related to the dose employed and can be prevented only by pretreating patients with fairly high intravenous methylprednisolone doses (up to 500 mg) (22). Caution is therefore suggested for the use of RTX in patients with suspected subclinical DON (1).

Restoration of euthyroidism after RTX may be due to either its effect on antibodies stimulating the TSH receptor (23) or to

remission of hyperthyroidism after prolonged antithyroid treatment (24). While some authors have hypothesized that RTX does induce a decrease of serum TRAb in patients with GD (25, 26), others have shown that the levels of antibodies stimulating the TSH receptor (TSAb) are not modified in GO patients (23). RTX causes depletion of B lymphocytes in the peripheral blood but also in the thyroid (27) and in the orbit (9) and histological analysis of orbital tissues of patients who received RTX showed absence of both B and T lymphocytes (8). This observation supports the hypothesis that RTX blocks B cell function as antigen presenting cells and interaction with helper T cells, required for the initiation of the autoimmune process (28). Interestingly, we observed that the number of NK cells increased following RTX infusion in Group 1 and 2, compared to baseline values. Previous studies on rheumatoid arthritis and idiopathic membranous nephropathy have shown a rise in the NK number, associated with an altered NK cell function. Although, the clinical significance of this observation needs to be established, it has been associated with a better response to RTX treatment (29, 30).

This may explain why after CD19+ cells repopulation, inflammation does not relapse and the CAS continues to decrease until stable disease inactivation (31). We in fact did observe that although CD19+ cells return earlier (<24 weeks) in patients treated with the 100 mg dose, when compared to Groups 2 and 3, this does not result in GO reactivation.

In analyzing the use of RTX in GO, we also looked at the cost of therapy in relation to the administered dose. When compared to first line treatment with intravenous steroids, the cost of a full cycle of treatment with RTX in Europe is about 4.8, 24 and 61 times more expensive for the 100, 500 and 1000 x 2 mg dose, respectively. Given the more favorable response/relapse rate and the clinical evidence shown by this study, we believe that a single RTX 500 mg dose is best indicated as second line treatment for the majority of GO patients.

The results of this study have to be seen in light of some limitations: in first instance the retrospective study design, the small sample size and its heterogeneity. Indeed, cases had a different duration of GO (which was longer in patients belonging to Group 3 compared to Group 1 and 2) and were enrolled among Graves'disease, Hashimoto's thyroiditis and euthyroid GO patients. Furthermore, some patients received previous corticosteroid treatment that may have had influenced the final outcome.

In this study, in conclusion, we sought to understand, from the limited available evidence, which dose of RTX is optimal as second line therapy of active moderate-to-severe GO. In this analysis we considered, both the effectiveness of the drug in

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inactivating the disease, the possibility of changing its natural course, the effect on disease severity and the impact on the patients' quality of life. Prospective randomized studies specifically addressing RTX dosing in GO are warranted.

DATA AVAILABILITY STATEMENT

The datasets presented in this article are not readily available because they contain some sensitive personal data. Requests to access the datasets should be directed to the corresponding author.

ETHICS STATEMENT

The studies involving human participants were reviewed and approved by Fondazione IRCCS Ca' Granda, Ospedale Policlinico, Milan, Italy. The patients/participants provided their written informed consent to participate in this study.

AUTHOR CONTRIBUTIONS

Data of each trial were collected by IC, DC, GV, MS, EL, LF, NC, LP and CG, BM and MD performed some serological assays and collected biochemical and cytofluorimetric data, the final database was produced by IM and EL, results were analyzed by IC and EL, the draft of the manuscript was written by IM and MS, the final manuscript was reviewed and revised by MS and MA. All authors contributed to the article and approved the submitted version.

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SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at: https://www.frontiersin.org/articles/10.3389/fendo.2021.790246/full#supplementary-material

Supplementary Figure 1 | Peripheral T cell, CD4+ (light gray) and CD8+ (white) and NK (dark grey) count measured at baseline and at different time-points after treatment with different doses of RTX. Test: Repeated measure ANOVA.

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