



OPEN ACCESS

EDITED AND REVIEWED BY

Ana Cuenda,
Spanish National Research Council (CSIC),
Spain

*CORRESPONDENCE

Payel Roy,
✉ proyl@lji.org
Thomas Riffelmacher,
✉ triffelmacher@lji.org

RECEIVED 07 February 2024

ACCEPTED 08 February 2024

PUBLISHED 22 February 2024

CITATION

Roy P, Winkels H, Orecchioni M,
Quesada-Masachs E and Riffelmacher T (2024),
Editorial: Role of innate and adaptive immune
cells in the metabolic syndrome.
Front. Cell Dev. Biol. 12:1383642.
doi: 10.3389/fcell.2024.1383642

COPYRIGHT

© 2024 Roy, Winkels, Orecchioni, Quesada-
Masachs and Riffelmacher. This is an open-
access article distributed under the terms of the
[Creative Commons Attribution License \(CC BY\)](https://creativecommons.org/licenses/by/4.0/).
The use, distribution or reproduction in other
forums is permitted, provided the original
author(s) and the copyright owner(s) are
credited and that the original publication in this
journal is cited, in accordance with accepted
academic practice. No use, distribution or
reproduction is permitted which does not
comply with these terms.

Editorial: Role of innate and adaptive immune cells in the metabolic syndrome

Payel Roy^{1,2*}, Holger Winkels^{3,4}, Marco Orecchioni^{1,2},
Estefania Quesada-Masachs¹ and Thomas Riffelmacher^{1,5*}

¹Center for Autoimmunity and Inflammation, La Jolla Institute for Immunology, La Jolla, CA, United States, ²Immunology Center of Georgia, Augusta University, Augusta, GA, United States, ³University of Cologne, Faculty of Medicine and University Hospital Cologne, Clinic III for Internal Medicine, Cologne, Germany, ⁴Center for Molecular Medicine Cologne (CMMC), University of Cologne, Cologne, Germany, ⁵Immunometabolism Core Facility, La Jolla Institute for Immunology, La Jolla, CA, United States

KEYWORDS

inflammation, metabolism, immune cells, signaling, chronic disease, therapies

Editorial on the Research Topic

Role of innate and adaptive immune cells in the metabolic syndrome

Dyslipidemia, hypercholesterolemia, and hyperglycemia are well-recognized metabolic disturbances that control the pathogenesis of a multitude of diseases including obesity-associated metabolic syndrome, diabetes, cardiovascular diseases (CVD), and non-alcoholic fatty liver disease (Hotamisligil, 2017). The presence of lipid-rich and/or glucose-rich microenvironments prompts metabolic adaptations within immune and non-immune cells (Hotamisligil and Erbay, 2008). An intricate crosstalk between the diverse cell types results in the release of a myriad of inflammatory mediators that disrupt crucial metabolic processes including lipolysis, lipogenesis, insulin sensitivity, and glucose utilization (Chawla et al., 2011; Hotamisligil, 2017). This maladaptive inflammation exacerbates progression of several chronic diseases.

In this Research Topic, we curated articles that highlight latest discoveries related to immune cell heterogeneity and crosstalk, activation of signaling pathways, receptor-ligand interactions and release of inflammatory mediators that modulate disease-associated inflammation and metabolic reprogramming.

Mazitova et al. discuss how the complex interplay between fat cells (adipocytes) and myeloid immune cells (macrophages, dendritic cells, neutrophils) regulates the progression of atherosclerosis, a pathological condition that is the most common cause of many CVDs. They describe the various types of adipocytes in the perivascular adipose tissue (PVAT), their unique adipokine profiles, and how these cells in turn alter the immune cell composition and cytokine production in atherosclerotic blood vessels. In this regard, the authors highlight the contributions of Interleukin (IL)-17, Type I and Type II interferons (IFN), IL-6 and IL-27 signaling pathways as important mediators. The authors suggest a dominant role of different functional subtypes of macrophages that are abundantly present both in adipose tissue and in atherosclerotic plaques. Targeting inflammatory responses in a cell-type dependent manner could be a novel therapeutic avenue to address obesity induced meta-inflammation which drives CVD.

Ray and Odum et al. delve deeper into the role of macrophages as regulators of inflammation. They emphasize that diverse macrophage subsets have unique functions in both promoting and resolving inflammation. In this regard, the authors highlight that embryonically-derived resident macrophages and infiltrated monocyte-derived macrophages have distinct and contrasting roles in

regulating chronic inflammation. They elaborately review a large body of literature that clarifies our mechanistic understanding of how macrophages exert pro- and anti-inflammatory effects in the context of four chronic diseases - insulin resistance and adipose tissue inflammation, atherosclerosis, non-alcoholic fatty liver disease and neurodegeneration. These factors can be broadly categorized into shared mechanisms related to lipid handling, production of cytokines and lipid mediators, efferocytosis or phagocytosis, and microRNA-mediated gene regulation. These pathways are promising targets for therapy. The authors further cautioned against the blocking of pathways that are active in both pro- and anti-inflammatory macrophage subsets, such as CCR2-CCL2 driven cell recruitment, as this will also impair resolution and tissue repair.

George et al. explore the mechanisms that allow the pleiotropic cytokine, IFN- γ , to both promote and restrain inflammation in the context of type 1 diabetes. As an inflammatory mediator, IFN- γ induces apoptosis and cell death, promotes expression of chemokines, receptors and adhesion molecules that facilitate immune cell recruitment and infiltration, augments antigen presentation and autoreactive T cell activation. On the other hand, IFN- γ has anti-inflammatory effects by limiting the proliferation of pathogenic T cells and by upregulating immune checkpoint molecules in the target tissue. The authors discuss that the pro-inflammatory role of IFN- γ is mediated by the activation of the Janus kinase/signal transducer and activator of the transcription (JAK/STAT) signaling pathway, while the regulatory effect is due to inhibition of JAK/STAT signaling by SOCS1 (suppressors of cytokine signaling 1). Based on this, the authors advocate the use of JAK inhibitors as strategies to limit inflammation in type 1 diabetes.

Liu et al. further extend this possibility and describe the importance of the JAK/STAT signaling pathway in the context of diabetic kidney disease (DKD), the most important microvascular complication of diabetes. The authors extensively reviewed available literature and proposed that JAK/STAT activation can affect the pathogenesis of DKD by influencing multiple factors, including the renin-angiotensin system (RAS), fibrosis, immune cell-mediated inflammation, cellular senescence, autophagy, and epithelial-to-mesenchymal transition (EMT). As these mechanisms are interconnected, inhibition of JAK/STAT signaling can limit disease progression by blocking multiple pathways. Finally, the authors discuss the potential of specific inhibitor molecules, natural compounds and other drugs that target the JAK/STAT pathway, as prospective treatment options in DKD. The safety and efficacy of one of these drugs, Baricitinib, in treating DKD patients, has been explored in a phase II, multicenter, double-blind, randomized, controlled clinical trial.

Guo et al. focus on upstream activators of signaling pathways and shed light on the role of pattern-recognition receptors (PRRs) in orchestrating maladaptive inflammation in endometriosis, a gynecological disorder that is characterized by hormonal imbalance and abnormal growth outside the uterine cavity. Recent studies have shown a pivotal role of immune cells in the etiology of endometriosis. The authors suggest that an inflammatory endometrial microenvironment is maintained by interactions between PRRs on immune cells with pathogen- and damage-associated molecular

patterns (PAMPs and DAMPs) in the surrounding milieu. They discuss the potential involvement of all five major groups of PRRs, namely, the toll-like receptors (TLRs), c-type lectin receptors (CLRs), nod-like receptors (NLRs), retinoic acid-inducible gene I-like receptors (RLRs), and absent in melanoma 2 (AIM2)-like receptors (ALRs). The authors propose that PRRs, particularly the TLR molecules, can be targeted to block inflammation.

These studies collectively highlight several distinct and overlapping mechanisms through which immune cells regulate the pathogenesis of inflammatory and metabolic diseases. They emphasize that a complete understanding of diverse pathways needs to be combined with targeted therapies for future clinical success.

Author contributions

PR: Writing—original draft, Writing—review and editing. HW: Writing—review and editing. MO: Writing—review and editing. EQ-M: Writing—review and editing. TR: Writing—original draft, Writing—review and editing.

Funding

The author(s) declare financial support was received for the research, authorship, and/or publication of this article. HW is supported by Deutsche Forschungsgemeinschaft (GRK 2407: 360043781); SFB TRR259 (397484323 project A04 and A09 and project 535107899); the Center for Molecular Medicine Cologne; and the Neven-DuMont Foundation. MO is supported by American Heart Association's Career Development Award (941152). EQ-M is supported by NIH R01AI092453.

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.

The author(s) declared that they were an editorial board member of Frontiers, at the time of submission. This had no impact on the peer review process and the final decision.

Publisher's note

All claims expressed in this article are solely those of the authors and do not necessarily represent those of their affiliated organizations, or those of the publisher, the editors and the reviewers. Any product that may be evaluated in this article, or claim that may be made by its manufacturer, is not guaranteed or endorsed by the publisher.

References

- Chawla, A., Nguyen, K. D., and Goh, Y. P. S. (2011). Macrophage-mediated inflammation in metabolic disease. *Nat. Rev. Immunol.* 11, 738–749. doi:10.1038/nri3071
- Hotamisligil, G. S. (2017). Inflammation, metaflammation and immunometabolic disorders. *Nature* 542, 177–185. doi:10.1038/nature21363
- Hotamisligil, G. S., and Erbay, E. (2008). Nutrient sensing and inflammation in metabolic diseases. *Nat. Rev. Immunol.* 8, 923–934. doi:10.1038/nri2449