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Ruxolitinib synergizes with regulatory T cells to improve inflammation but has no added benefits in decreasing albuminuria in SLE

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Background: Umbilical cord blood (UCB)-derived CD4⁺CD25⁺CD127^{low} regulatory T cells (Tregs) can decrease albuminuria and anti-dsDNA IgG in systemic lupus erythematosus (SLE). Ruxolitinib, a JAK/STAT inhibitor, has been shown to improve cutaneous manifestations of SLE. We hypothesize that the addition of ruxolitinib to UCB-Tregs may improve SLE outcomes.

Methods: *In vitro* cell suppression, phenotype change, IL-10 secretion, and cytokine levels in coculture supernatants were determined to quantify the impact of adding ruxolitinib to UCB-Tregs. A xenogeneic SLE model was utilized to study their *in vivo* combination.

Results: In a dose-dependent manner, ruxolitinib addition synergizes with UCB-Tregs to suppress SLE-PBMC proliferation, inhibit CD8⁺ T cells, and reduce phosphorylation of STAT3/STAT5/AKT in CD8⁺ T cells. UCB-Treg and ruxolitinib combination also downregulates the soluble form of inflammatory cytokines including IFN- γ , IP-10, TNF- α , IL-6, sCD40L, IL-17A, IL-17F, IL-1 α , and LIF in cocultures. The addition of ruxolitinib increases UCB-Treg cell persistence in peripheral blood *in vivo* and decreases the soluble form of human inflammatory cytokines including IFN- γ , TNF- α , and sCD40L in plasma along with improvement of skin lesions in SLE xenografts. Compared to control, significantly lesser CD3⁺, CD4⁺, CD8⁺, and Ki-67⁺ infiltrates are observed in the lung and kidney of UCB-Tregs and/or ruxolitinib recipients. No added benefit of addition of ruxolitinib is observed on the significant improvement in the urine albumin/creatinine ratio and the anti-dsDNA IgG levels induced by UCB-Tregs.

Conclusions: Our results demonstrate that the addition of ruxolitinib to UCB-Tregs increases UCB-Tregs suppressor function and their persistence *in vivo*, downregulates systemic inflammation, and controls cutaneous SLE but does not add to UCB-Treg-mediated improvement in renal manifestations.

KEYWORDS

adoptive cell therapy, regulatory T cells (Tregs), allogeneic, umbilical cord blood (UCB), ruxolitinib, JAK/STAT pathway, inflammation, systemic lupus erythematosus (SLE)

Highlights

- In vitro, the addition of ruxolitinib to UCB-Tregs augments the inhibition of pathogenic CD8⁺ T cells, and the reduction of soluble IFN-γ, IP-10, TNF-α, IL-6, sCD40L, IL-17A, IL-17F, IL-1α, and LIF production.
- In vivo, ruxolitinib improves UCB-Treg cell persistence and decreases CD3⁺, CD4⁺ T, CD8⁺ T, and Ki-67⁺ cell tissue infiltration but has no added benefits to albuminuria and anti-dsDNA IgG Ab.

Introduction

Although traditionally thought to be a B-cell disorder, recent data support a central role of T cells in the pathogenesis of systemic lupus erythematosus (SLE) (1). T cell dysregulation and autoreactive T effector (Teff) cells affect peripheral tolerance and induce inappropriate activation of B cells (2). T lymphocytes are increasingly being recognized as key contributors to disease pathogenesis, where CD4 T follicular helper cells enable autoantibody production, inflammatory Th17 subsets promote inflammation, while defects in regulatory T cells (Tregs) lead to unchecked immune responses (3).

Th1 cytokines including IFN- γ promote B-cell class switching and stimulate pathogenic autoantibody production in SLE (4). Levels of IFN- γ are elevated in patients with SLE compared to controls and positively correlate with SLE disease activity index (SLEDAI) scores (5). Th2 cytokines including IL-4 promote B-cell differentiation into plasma cells and induce antibody class switching to IgG1 and IgE (6). In lupus-prone mice, blocking IL-4 decreases anti-double-stranded DNA (anti-dsDNA) antibodies, whereas the administration of IL-4 increases the levels of this autoantibody (7). Th17 cytokines have been shown to correlate with the SLE severity where IL-17 levels correlate with SLEDAI scores, and both IL-17 and IL-23 are associated with treatment-resistant active nephritis (8, 9). Therefore, autoreactive T cells drive the severity of disease in SLE (2).

Tregs develop in the thymus through strong T cell receptor (TCR) signaling just below the threshold for negative selection and recognize self-antigens for their differentiation (10). Tregs express the CD4⁺CD25⁺CD127^{low} phenotype with high intracellular FOXP3 (11). Autoreactive T cells that escape negative selection in the thymus can persist in the periphery where Tregs prevent their aberrant activation and expansion, thus acting as important gatekeepers (12). By neutralizing these autoreactive T cells as well as other harmful immune cells, Tregs play a central role in the resolution of unwanted inflammation (11, 13). Furthermore, Tregs homing to the areas of inflammation are based on their ability to sense the source of survival cytokine, IL-2, and migration to the zones of immune activation where they "steal" IL-2 from effector T cells promoting their apoptosis (14). In all, Tregs perform several functions including (i) preventing autoimmunity, (ii) suppressing inflammation, (iii) maintaining tissue integrity, (iv) controlling allergies including asthma, (v) inducing tolerance to dietary antigens and the fetus, and (vi) protecting commensal bacteria from being eliminated by the immune system (15). Defective and decreased Tregs can lead to autoimmune diseases including SLE (10). In fact, compared to a healthy population, SLE patients have been shown to have a lower percentage of Tregs and have defects in their suppressor function (16–19). Furthermore, the pathogenic Teff cells in SLE patients develop resistance to Treg-induced suppression that supports their unopposed proliferation, leading to excessive secretion of inflammatory cytokines that contribute to SLE pathogenesis (20).

Recently, we showed that umbilical cord blood (UCB)-derived Treg cells (UCB-Tregs) have unique properties including (i) lack of plasticity when exposed to inflammatory micro-environments; (ii) no requirement for HLA matching with the recipients; (iii) long shelf life of the cryopreserved cells; and (iv) immediate product availability for on-demand treatment (21). In addition, UCB-Tregs are able to decrease the production levels of anti-dsDNA IgG antibody and inflammatory cytokines and improve albuminuria in the xenogeneic model of SLE (22). In the same study, it was shown that UCB-Tregs are able to suppress pathogenic SLE PBMCs to a similar extent as the healthy PBMCs (22). Therefore, adoptive therapy with UCB-Tregs for SLE might be a promising approach. A concern remains for the possibility of resistance since in response to the unusually high levels of plasma cell-niche cytokines in SLE patients, the auto-antibody production is dependent on JAK/ STAT3 activation such that this process can be abrogated by inhibition of this pathway by the JAK inhibitor, ruxolitinib (23). Although ruxolitinib as a single agent has shown significant activity in steroid refractory acute graft vs. host disease (GVHD), a T cellmediated fatal complication of allogeneic stem cell transplantation (24), when used as a therapeutic strategy for SLE, despite showing activity in cutaneous lupus by decreasing inflammation, no improvement in the systemic organ involvement was observed in ruxolitinib recipients (25-27). The combination of ruxolitinib with Tregs has been examined in a GVHD model with an observed decrease in the incidence and severity and improved survival without dampening of the graft versus leukemia effect (28). We have shown a synergistic effect of UCB-Tregs and ruxolitinib in xenogeneic GVHD as well (29).

Here, we hypothesize that the combination of ruxolitinib with UCB-Tregs may lead to a synergistic activity in treating SLE manifestations.

Materials and methods

Cell source

UCB-Treg cells were generated as described previously (21). Human peripheral blood mononuclear cells from SLE patients (SLE-PBMCs, BioIVT, Westbury, NY, USA) or healthy donors (HD-PBMCs, Gulf Coast Blood Bank, Houston, TX, USA) and UCB-Tregs were cultured as described previously (22). Additionally, cryopreserved UCB-derived *ex vivo* expanded CD4⁺CD25⁺127^{low} Treg cells were obtained from Cellenkos Inc. (Houston, TX, USA).

Phenotype analysis

UCB-Treg cells, SLE-PBMCs, or UCB-Treg: SLE-PBMC (1:1) cocultures were stained using the following antibodies: APC-eFluor 780-CD45 (HI30), Alexa Fluor 532-CD3 (UCHT1), FITC-CD3 (UCHT1), PerCP-Cyanine5.5-CD8a (RTA-T8), Super Bright 600-CD19 (SJ25C1), PE-CD25 (BC96), PE-Cy5-CD127 (eBioRDR5), APC-CD56 (CMSSB), FITC-CD16 (eBioCB16) (CB16), PerCPeFluor 710 CD14 (61D3), PE-Cy7-HLA-DR (LN3), PE-Cyanine7phosphor-STAT3 (Tyr705) (LUVNKLA), and LIVE/DEADTM fixable Blue (Thermo Fisher Scientific); BV650-CD4 (L200), BV510-CD8 (RPA-T8), PE-CF594-CD27 (M-T271), Alexa Fluor 700-IgD (IA6-2), BV421-CD62L (SK11), Alexa Fluor 647-Helios (22F6), Alexa Fluor 647-FoxP3 (259D/C7), PerCP-Cy5.5-FoxP3 (236A/E7), Pacific Blue-Stat5 (pY694) (47/Stat5) (pY694), BV421-Akt (pS473) (M89-61), and BUV-395-Ki-67 (B56) (BD Biosciences); and Pacific Blue-CD45.1 (A20) (Southern Biotech, Birmingham, AL, USA). T, Treg, B, NK cells, and monocytes were gated and displayed on the t-distributed stochastic neighbor embedding (t-SNE). The following CD8⁺ T cells were further analyzed: p-Stat3⁺, p-Stat5⁺, Ki-67⁺, or p-Akt⁺ CD8⁺ T cells. Cytek Aurora (Cytek Biosciences, Fremont, CA, USA), BD LSRFortessa (BD Biosciences), and FlowJo software (FlowJo, LLC, Ashland, OR, USA) were used for phenotype analysis.

Ruxolitinib IC₅₀

Ruxolitinib (INCB018424; S1378) was purchased from Selleck Chemicals (Houston, TX). The comparative half-maximal inhibitory concentration (IC₅₀) values of ruxolitinib against HD-PBMCs (Gulf Coast Blood Bank), SLE-PBMCs (BioIVT), or UCB-Tregs after treatment with ruxolitinib for 6 days were assessed as described previously (30).

Functional ability of UCB-Treg cells

In the presence of 0, 25, or 100 nmol/L ruxolitinib and CD3/ CD28 beads, the suppressive function of UCB-Tregs on the proliferation of CD4⁺CD25⁻ conventional T cells (Tcons) or SLE-PBMCs were assessed as described previously (21, 22).

Production levels of soluble human cytokines

UCB-Tregs, SLE-PBMCs, or UCB-Treg: SLE-PBMC (1:1) cocultures were cultured for 3 or 6 days in the absence or presence of ruxolitinib. The soluble form of human cytokines was measured using Cytokine ELISA kits from Thermo Fisher Scientific or Eve Technologies (Calgary, AB, Canada).

SLE xenograft model

Animal procedures were performed according to an approved IACUC protocol by The University of Texas MD Anderson Cancer Center. In vivo efficacy assessment was done using a SLE xenograft model as described previously (21, 22). Mice were divided into four groups (No Rx, Rux, UCB-Treg, and UCB-Treg+Rux, n = 5 mice/ group) after they displayed human immune cells. A total of 5×10^6 UCB-Treg cells were administered by tail vein injection on days 28, 32, 42, and 46. Ruxolitinib was administered orally at a dose of 90 µg on days 35, 36, 37, 38, 39, 49, 50, 51, 52, and 53. The reconstitution level of human immune cells and weight loss was monitored weekly or twice per week, respectively. Mice were euthanized using 30%-70% displacement of the chamber volume per minute with compressed CO₂, and death was confirmed by cervical dislocation. At the time of euthanasia, single-cell suspensions from each organ were aseptically isolated and phenotypic analysis of human immune cells was determined based on the expression levels of human cell surface and intracellular markers.

Urinary albumin in SLE xenografts

For the kidney function assessment, the production levels of albumin and creatinine in mouse urine samples were assessed as described previously (22).

Anti-human dsDNA IgG antibody and human cytokine/chemokine in SLE xenografts

The production levels of anti-human dsDNA IgG Ab and human inflammatory cytokines in mouse plasma samples were measured as described previously (22).

Histopathology and immunohistochemistry

Mouse organs were harvested, fixed, processed, and embedded in paraffin. For histopathology analysis, tissue sections were stained with hematoxylin and eosin. The presence of human immune cells in the mouse tissues was assessed by immunohistochemical staining with the following antibodies: human CD3 (Clone F7.2.38) (DAKO, Santa Clara, CA, USA), CD4 (Clone 4B12) (Leica Biosystems Inc., Buffalo, Grove, IL, USA), CD8 (Clone C8/144B) (Thermo Fisher Scientific), and Ki-67 (Clone MIB-1) (DAKO). The stained tissue slides were scanned using Aperio AT2 (Leica Biosystems Inc., Buffalo Grove, IL, USA). Total positive cells and *H*-scores for human CD3, CD4, CD8, and Ki-67 were calculated using the HALO 3.3 software (India Labs, Albuquerque, NM, USA).

Statistical analysis

All statistical analyses were done with GraphPad Prism 10 software (San Diego, CA, USA). Data are presented as mean \pm SEM. *p*-values were obtained using one or two-way analysis of variance (ANOVA) with Tukey's or Šídák's multiple comparison test, *F*-test, or two-tailed unpaired *t*-test with 95% confidence interval for evaluation of statistical significance compared with the untreated controls. *p* < 0.05 was considered statistically significant.

Results

Addition of ruxolitinib improves UCB-Tregs suppressor function and decreases inflammation of SLE

We have previously demonstrated that CD4+CD25+127^{low} UCB-Tregs can inhibit various inflammatory cytokines and improve skin damage in a SLE xenograft model (22). Ruxolitinib has been shown to inhibit inflammatory cytokines in cutaneous lupus erythematosus (26). To examine whether addition of ruxolitinib to adoptive therapy with UCB-Tregs can enhance their anti-inflammatory effect in SLE, we first examined the IC₅₀ values for ruxolitinib in coculture with UCB-Tregs, HD-PBMCs, or SLE-PBMCs. As shown in Figure 1A, no differences were observed in the ruxolitinib IC₅₀ among the three different cell types after 6 days of incubation (IC₅₀ = 36.3-43.9 nmol/L, 45.5 nmol/L, and 33.4 nmol/L, respectively), suggestive of no direct impact of the pathogenicity of SLE on ruxolitinib effect. Since ruxolitinib has been shown to have differential effect on Tregs in vivo (31), we next assessed the impact of the addition of ruxolitinib on UCB-Treg function. The addition of ruxolitinib at 25 or 100 nM synergized with UCB-Tregs to suppress proliferation of HD-Tcon (two-way ANONA interaction: *p* < 0.0001; Treg: Tcon ratio: *p* < 0.0001; Rux conc: *p* < 0.0001) (Figure 1B) and SLE-PBMC (two-way ANOVA interaction: *p* < 0.0001; Treg: SLE-PBMC ratio: p < 0.0001; Rux conc: p < 0.0001) (Figure 1C), across different cell ratios, especially as low as the 1:8 ratio of UCB-Treg to SLE-PBMC. We next examined whether ruxolitinib can affect UCB-Tregs' ability to secrete the suppressor cytokine, IL10 (22). As shown in Figure 1D, on day 3 of culture, high levels of IL-10 were detected in the supernatants of UCB-Tregs (4,844 ± 78 pg/mL) whereas the production level of soluble IL-10 was 190 ± 14 pg/mL in SLE-PBMC alone and $3,505 \pm 18$ pg/mL in the coculture of UCB-Tregs with SLE-PBMC. In the presence of 25 nM ruxolitinib, the respective IL-10 secretion levels in the culture supernatants decreased to $941 \pm 53 \text{ pg/}$ mL, 46 ± 6 pg/mL, and 1,149 ± 33 pg/mL, respectively (one-way ANOVA, *p* < 0.0001).

To examine the impact of the addition of ruxolitinib to UCB-Tregs on the immune cell compartment of SLE-PBMCs, CD4⁺ T, CD4⁺CD25⁺CD127^{low} Treg⁺, CD4⁺CD8⁺ T, CD8⁺ T, CD19⁺ B, CD56⁺ NK cells, and CD14⁺ monocytes were gated (Supplementary Figure S1). As shown in Figures 1E, F and Supplementary Figure S2, coculture with UCB-Tregs decreased the inflammatory cell subpopulation of SLE-PBMCs. Three-day coculture increased the percentages of CD4⁺CD25⁺CD127^{low} Treg cells and decreased those of CD19⁺ B cells, CD56⁺ NK cells, and CD14⁺ monocytes in UCB-Tregs, SLE-PBMCs, and UCB-Treg:SLE-PBMC, respectively. Interestingly, a significant reduction of CD8⁺ T cells was observed in the UCB-Treg:SLE-PBMC coculture but not SLE-PBMC cultures in the absence and presence of ruxolitinib. The addition of ruxolitinib led to an increase in the double-positive (DP) CD4⁺CD8⁺ T cells in SLE-PBMC+UCB-Tregs coculture.

We next examined whether ruxolitinib can affect STAT3 signaling in SLE-PBMCs and/or UCB-Tregs. A significant reduction of p-STAT3⁺, Ki-67⁺, and p-Akt⁺CD8⁺ T cells was observed in 100 nM ruxolitinib-treated UCB-Tregs and SLE-PBMCs (p < 0.0001) and UCB-Treg: SLE-PBMC coculture (p < 0.01 for p-Stat3⁺ and p < 0.0001for Ki-67⁺ and p-Akt⁺) (Figure 1G). In addition, SLE-PBMCs (p < 0.0001) and UCB-Treg: SLE-PBMC coculture (p < 0.01)showed a significant reduction of p-STAT5+CD8+ T cells after treatment with 100 nM ruxolitinib. As shown in Figure 1H, the addition of ruxolitinib and/or UCB-Tregs to SLE-PBMCs significantly decreased the soluble form of inflammatory cytokines, including IFN- γ , IP-10, TNF- α , IL-6, sCD40L, IL-17A, IL-17F, IL-1 α , and LIF (p < 10.0001). Interestingly, ruxolitinib did not have any effect on TNF- α production and surprisingly seemed to increase soluble IL-17A production by SLE-PBMCs in a dose-dependent manner whereas UCB-Tregs led to a significant reduction of IL-17A.

The addition of ruxolitinib to UCB-Tregs improves their persistence *in vivo* and cutaneous lesions of SLE

Using the SLE xenograft model as described previously (22), 4 weeks were allowed for human disease to be established in immunedeficient mice followed by multiple treatments of UCB-Tregs (5 \times 10^6 cells) alone, ruxolitinib alone (90 ×g), or a combination of UCB-Tregs and ruxolitinib. UCB-Tregs were administered by tail vein on days 28, 32, 42, and 46. Ruxolitinib was administered by oral gavage on days 35, 36, 37, 38, 39, 49, 50, 51, 52, and 53. UCB-Tregs plus ruxolitinib were administered at their corresponding time points (Figure 2A). SLE disease phenotype was evident in the control arm, where mice developed malar, discoid, and erythematous skin rash, and/or hair loss, similar to that observed in human disease (22). The mice in ruxolitinib alone arm but not UCB-Treg recipients with or without ruxolitinib developed noticeable hair loss (Figure 2B). The burden of SLE disease was measured by monitoring the change of body weight and GVHD score, longitudinally. An increase in body weight measured over 68 days was observed in all groups (Figure 2C, two-way ANOVA Time and Group; p < 0.0001). The control SLE group (No Rx) had the highest GVHD score (Figure 2D, two-way ANOVA interaction, Time and Group; p <0.0001), circulating human CD45⁺ cells (Figure 2E, p < 0.05), and circulating human CD8⁺ T cells (Figure 2F, p < 0.05), when compared to either of the treatment arms (Rux, UCB-Treg,

and UCB-Treg+Rux). The percentage of circulating human $CD4^+CD25^{high}CD127^{low}$ Treg cells was significantly increased following UCB-Tregs injection and was detected up to day +70 in the UCB-Treg+ruxolitinib recipients (Figure 2G, p < 0.0001).

As shown in Figure 2H, lung tissue in the control arm (No Rx) and in the ruxolitinib recipients (Rux) showed significant alveolar disruption whereas UCB-Treg and UCB-Treg+ruxolitinib recipients preserved their alveolar air space. Compared with the control arm (No Rx), lung tissue in all treatment groups had a significant reduction in the total positive cells and the *H*-score for human CD3, CD4, CD8, and Ki-67 (Figure 2I, p < 0.0001 for Group and Human Markers). All treatment groups have well-preserved spleen tissue architecture whereas the control arm (No Rx) showed lymphocytic infiltration into red and white pulp with splenomegaly (Supplementary Figure S3A). Compared with the control group (No Rx), all treatment groups had a significant reduction in the total positive cells and the *H*-score for human CD3, CD4, CD8, and Ki-67 (Supplementary Figure S3B, p < 0.0001 for Group and Human Markers).

The addition of ruxolitinib to UCB-Tregs reduces systemic inflammation without added benefits for reduction in albuminuria and anti-dsDNA IgG Ab

We have previously shown that UCB-Tregs can treat SLE and decrease albuminuria and anti-dsDNA IgG Ab (22). Here, we examined the impact of adding ruxolitinib to UCB-Treg recipients. As shown in Figure 3A, a significant decrease in albuminuria (A/C ratio) was observed in all treatment groups but not the control arm (No Rx vs. Rux or UCB-Treg+Rux, p < 0.05; No Rx vs. UCB-Treg, p < 0.01). A corresponding decrease in the anti-dsDNA IgG Ab was observed in all treatment arms when compared to control. (Figure 3B; control vs. UCB-Treg or UCB-Treg+ruxolitinib, p < 0.01; for UCB-Treg vs. Rux, p < 0.05). Specifically, no differences were seen between UCB-Treg vs. the UCB-Tregs+Rux arm. We observed that a significant reduction of human CD20⁺ B cells was observed in the spleen of treatment arms when compared to control (Supplementary Figure S3C; control vs. Rux, p < 0.05; control vs. UCB-Treg+ruxolitinib, p < 0.01).



FIGURE 1

Addition of ruxolitinib synergizes with UCB-Tregs to suppress SLE-PBMCs and decrease inflammatory cytokine secretion. (A) Cytotoxicity of ruxolitinib against HD-PBMCs, SLE-PBMCs, or UCB-Tregs when used as a single agent. (B) Assessment of UCB-Tregs suppression function on proliferation of Tcons from healthy donor (left panel) and representative histogram (right panel) of suppressive activity of UCB-Tregs against Tcons when cocultured at 1:1, 1:2, and 1:4 ratio at 0, 25, or 100 nM ruxolitinib. (C) Assessment of UCB-Tregs suppression function on proliferation of SLE-PBMCs (top panel) in the absence and presence of ruxolitinib and representative histogram (bottom panel) of suppressive activity of UCB-Tregs against SLE-PBMCs when cocultured at 1:1, 1:2, 1:4, or 1:8 ratio at 0, 25, or 100 nM ruxolitinib. (D) Production levels of soluble IL-10 in the absence and presence of 25 nM ruxolitinib. Data are presented as mean \pm SEM (n = 3). p < 0.05 was considered statistically significant. *p < 0.05; **p < 0.01; ***p < 0.001 by one- or two-way ANOVA with Tukey's multiple comparison tests. (E) Subset analysis on the tSNE map. (F) Quantification analysis of subsets. CD4⁺ T, CD4⁺CD25⁺CD127^{low} Treg⁺, CD4⁺CD8⁺ T, CD8⁺ T, CD9⁺ B, CD56⁺ NK cells, and CD14⁺ monocytes were quantified. Data are presented as mean \pm SEM (n = 5). (G) Quantification analysis of p-Stat5⁺, Ki-67⁺, or p-Akt⁺CD8 + T cells were quantified. Data are presented as mean \pm SEM (n = 5). (G) Quantification analysis of p-Stat5⁺, Ki-67⁺, or p-Akt⁺CD8 + T cells were quantified. Data are presented as mean \pm SEM (n = 3). p < 0.05 was considered statistically significant. (H) Production levels of soluble cytokines on heatmap. UCB-Tregs, SLE-PBMCs, or UCB-Tregs:SLE-PBMC NC = 3.). p-Stat5⁺, Ki-67⁺, or p-Akt⁺CD8 + T cells were quantified. Data are presented as mean \pm SEM (n = 3). p-stat5⁺, Ki-67⁺, or p-Akt⁺CD8 + T cells were quantified. Data are presented as mean \pm SEM (n = 3).



circulating human CD8⁺ T cells in all treatment recipients. **(G)** Sustained increase in circulating human CD4⁺CD25⁺CD127^{low} Treg cells in UCB-Treg +ruxolitinib recipients. Data are presented as mean \pm SEM (n = 5). p < 0.05 was considered statistically significant. *p < 0.05; ***p < 0.001; ****p < 0.0001 by two-way ANOVA with Tukey's multiple comparison tests or Student *t*-test. **(H)** UCB-Tregs in combination with ruxolitinib improve lung tissue damage in SLE xenografts. **(I)** Quantification analysis of total positive cells and *H*-scores for human CD3, CD4, CD8, and Ki-67 in lung tissues. Data are presented as mean \pm SEM (n = 5). p < 0.05 was considered statistically significant. *p < 0.01; ***p < 0.001; ***p < 0.001 by two-way ANOVA with Tukey's multiple comparison tests.

Histopathological analysis of kidney tissue showed significant lymphatic infiltrate in the control arm (No Rx) compared with preservation of the kidney architecture in UCB-Treg and UCB-Treg +ruxolitinib recipients (Figure 3C). A significant decrease in the total number (Figure 3D, left panel, p < 0.0001 for Group and p = 0.0288 for Human Marker) of human CD3⁺, CD4⁺, and Ki-67⁺ cells and the *H*score (Figure 3D, right panel, p < 0.0001 for Group and p = 0.0131 for Human Marker) was observed in all treatment groups but not in the control arm. In addition, all treatment arms showed an improvement at the kidney tissue and functional level and a decrease in systemic inflammation and the expression levels of circulating human inflammatory cytokines including IFN- γ , TNF- α , and sCD40L, which overlapped with those impacted in *in vitro* studies (Figure 3E; *p < 0.05, ***p < 0.001, ****p < 0.0001).

Discussion

Here, we show that the addition of ruxolitinib improves UCB-Tregs' ability to suppress proliferation of healthy as well as SLEderived PBMCs. Such synergy translates into prolonged persistence

of Tregs in vivo and improvement in the skin lesions. Interestingly, we observed that the addition of ruxolitinib decreases Treg cell population within SLE-PBMCs and decreases IL-10 secreted by UCB-Tregs. Such a discrepancy might be explained by the differential effect of ruxolitinib on Tregs based on an underlying disease biology that has been previously described in bone marrow failure, where ruxolitinib increased splenic Tregs in diseased mouse but not in the healthy control (31). A decrease in the distribution of cell populations including T-cell compartment consisting of the CD4 and CD8 subsets as well as Tregs, B-cell compartment, NK cells and monocytes in response to UCB-Tregs, ruxolitinib, and their combination was also observed in our study, which is similar to the reported decrease in the frequencies of splenic CD4⁺ T, CD8⁺ T, and NK-T cells with a significant augmentation of splenic Tregs in a mouse model of autoimmune cholangitis after treatment with ruxolitinib (32). Additionally, hemophagocytic lymphohistiocytosis, a fatal complication of SLE, primarily driven by uncontrolled activation of pathogenic CD8⁺ T cells, has been shown to be responsive to ruxolitinib in a case report (33).

The negative impact of ruxolitinib on the IL-10 secretion from UCB-Tregs was an interesting observation that might be a function of

the decrease in the Treg cell number. In contrast, we show that the addition of ruxolitinib in fact increased the cell suppressor function of UCB-Tregs in vitro and increased their persistence in vivo. Such ruxolitinib-mediated synergy on UCB-Treg function and survival is suggestive of possible additional mechanisms at play, beyond IL-10 secretion. We show that ruxolitinib-mediated suppression of the STAT3/5-AKT pathway in the CD8⁺ T cells in SLE-PBMCs in the presence or absence of UCB-Tregs might suggest an independent mechanism of targeting SLE pathogenesis, since inhibition of STAT3 in T cells in lupus has been shown to delay the onset of nephritis (34). Furthermore, a decrease in the release of multiple inflammatory cytokines, in vitro and in vivo, including IFN-y, IP-10, IL-6, sCD40L, IL-17F, IL-1 α , and LIF, similar to that shown by ruxolitinib in GVHD (35) and COVID-19 (36), also supports the ruxolitinib-mediated mechanism of synergizing with UCB-Tregs independent of IL-10 secretion in SLE. Specifically, IFN-y has been shown to be complicit in SLE pathogenesis (37, 38). JAK inhibitors can suppress the IFN signaling in human dendritic cells, reduce CD80/CD86 expression and T-cell stimulation ability (39), and reduce the production of various inflammatory cytokines including IFN- γ (40) in SLE mice. On the other hand, UCB-Tregs, as a single agent, have also been shown to resolve SLE inflammation and decrease inflammatory cytokines including IFN-γ, IP-10, TNF-α, IL-6, IL-17A, sCD40L, and IL-1a (22).

Furthermore, a unilateral decrease in pathogenic CD8⁺ T cells in circulation and in tissues underscores the synergistic impact of adding ruxolitinib to enhance UCB-Tregs ability to suppress SLE-PBMC (22).

In SLE, pathogenic IL-17-producing CD4CD8 double-negative (DN) T cells are thought to originate from autoreactive CD8⁺ T cells, which also contribute to the induction and maintenance of systemic autoimmunity (41, 42). Specifically, these DN T cells promote autoantibody production; essentially, they are considered a key pathogenic cell population in SLE (43). In our study, we observed that UCB-Treg cells increased CD4/CD8 double-negative T cells after treatment with 100 nM ruxolitinib; however, such an increase was not associated with an increase in the production of IL-17A and IL-17F. In fact, the coculture of UCB-Tregs with SLE-PBMC significantly decreased the production of IL-17A and IL-17F by SLE-PBMCs (Figure 1H). On the other hand, the CD4CD8 double positive (DP) T cell population was shown to have a suppressive effect on the production of autoantibodies including antinuclear antibody and antidsDNA Ab in SLE (44) and is increased in response to UCB-Tregs in a SLE xenogeneic model (22). Although their exact significance is unclear, these DP T cells may represent a transition of T-cell population towards further differentiation (45). In vivo terminally differentiated effector CD4⁺ T cells may acquire the alpha-chain of CD8⁺ T cells, and these CD4⁺CD8⁺ T cells have been identified in autoimmune and chronic inflammatory disorders (46). Another study shows no difference in the DP T-cell population in SLE and healthy controls (47). Future studies would be required to better understand the impact of this subpopulation of T cells in SLE.

It is not surprising that the addition of ruxolitinib to UCB-Tregs did not add to their existing benefit of controlling systemic renal disease manifestations, as measured by urine albumin/creatinine ratio as well as anti-dsDNA IgG Ab. We have previously shown that single-agent UCB-Tregs can decrease anti-dsDNA IgG antibody and improve renal function in SLE (22). Therefore, the maximum benefit of adding ruxolitinib might be for control of dermal manifestation as seen in our data as well as reported by others (25–27).

Taken together, our results demonstrate that the addition of ruxolitinib to UCB-Tregs augments the inhibition of pathogenic CD8⁺ T cells expressing p-STAT3⁺, p-STAT5⁺, and p-Akt⁺, and the reduction of soluble IFN- γ , IP-10, TNF- α , IL-6, sCD40L, IL-17A, IL-17F, IL-1 α , and LIF production, and ruxolitinib improves UCB-Treg cell persistence and decreases CD3⁺, CD4⁺ T, CD8⁺ T, and Ki-67⁺ cell tissue infiltration but has no added benefits to albuminuria and anti-dsDNA IgG Ab. In conclusion, the addition of ruxolitinib to UCB-Tregs in SLE is a viable therapeutic strategy and should be explored in the clinical setting, especially for dermal manifestations.

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 humans were approved by The University of Texas M.D. Anderson Cancer Center Institutional Review Board. The studies were conducted in accordance with the local legislation and institutional requirements. The human samples used in this study were acquired from a by-product of routine care or industry. Written informed consent for participation was not required from the participants or the participants' legal guardians/next of kin in accordance with the national legislation and institutional requirements. The animal study was approved by The University of Texas M.D. Anderson Cancer Center Institutional Animal Care and Use Committee. The study was conducted in accordance with the local legislation and institutional requirements.

Author contributions

M-AL: Conceptualization, Data curation, Formal analysis, Investigation, Methodology, Project administration, Resources, Software, Supervision, Validation, Visualization, Writing – original draft, Writing – review & editing. XT: Formal analysis, Methodology, Writing – review & editing. MGR: Formal analysis, Writing – review & editing. MH: Formal analysis, Investigation, Writing – review & editing. KZ: Formal analysis, Investigation, Writing – review & editing. KZ: Formal analysis, Investigation, Writing – review & editing. CRF: Formal analysis, Investigation, Resources, Supervision, Validation, Writing – review & editing. SP: Conceptualization, Data curation, Formal analysis, Funding acquisition, Project administration, Resources, Supervision, Validation, Writing – original draft, Writing – review & editing.

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Conflict of interest

SP has an equity interest in, holds patents for, receives royalties and research funding from, and is a member of the board of directors/ advisory committee for Cellenkos Inc. CF received research funding from 4D, AbbVie, Acerta, Adaptimmune, Allogene, Amgen, Bayer, BostonGene, Celgene, Cellectis, EMD, Gilead, Genentech/Roche, Guardant, Iovance, Janssen Pharmaceutical, Kite, Morphosys, Nektar, Novartis, Pfizer, Pharmacyclics, Sanofi, Takeda, TG Therapeutics, Xencor, Ziopharm, Burroughs Wellcome Fund, Eastern Cooperative Oncology Group, National Cancer Institute, V Foundation, Cancer Prevention and Research Institute of Texas: CPRIT Scholar in Cancer Research and consulting fees from AbbVie, Bayer, BeiGene, Celgene, Denovo Biopharma, Foresight Diagnostics, Genentech/Roche, Genmab, Gilead, Karyopharm, N-Power Medicine, Pharmacyclics/ Janssen, SeaGen, and Spectrum. TS is an employee of Cellenkos Inc.

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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Supplementary material

The Supplementary Material for this article can be found online at: https://www.frontiersin.org/articles/10.3389/fimmu.2025.1449693/ full#supplementary-material

SUPPLEMENTARY FIGURE 1

Representative FACS analysis of subsets. The percentage of CD4⁺ T, CD4⁺CD25⁺CD127^{low} Treg⁺, CD4⁺CD8⁺ T, CD8⁺ T, CD19⁺ B, CD56⁺ NK cells, and CD14⁺ monocytes were acquired on a BD LSRFortessa X-20 flow cytometer and analyzed using FlowJo software.

SUPPLEMENTARY FIGURE 2

Quantification analysis of CD4⁺ T, CD4⁺CD25⁺CD127^{low} Treg⁺, CD4⁺CD8⁺ T, CD8⁺ T, CD19⁺ B, CD56⁺ NK cells, and CD14⁺ monocytes were quantified. Data

are presented as mean \pm SEM (n=3). *P*<0.05 was considered statistically significant. *P*<0.05; ****P*< 0.001; *****P*<0.001 by Student *t*-test.

SUPPLEMENTARY FIGURE 3

Addition of Ruxolitinib improves spleen damage in SLE xenografts. (A) Impact of UCB-Tregs in combination with ruxolitinib on spleen tissue in SLE

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mean \pm SEM (n=5). *P*<0.05 was considered statistically significant. **P*<0.05; *****P*<0.0001 by two-way ANOVA with Tukey's multiple comparison tests. (C) Quantification analysis of human CD20+ B cells in spleen tissues. Data are presented as mean \pm SEM (n=5). *P*<0.05 was considered statistically significant. **P*<0.05; ***P*<0.01 by Student *t*-test.

xenografts. (B) Quantification analysis of total positive cells and H-scores

for human CD3, CD4, CD8, and Ki-67 in spleen tissues. Data are presented as

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