

Interleukin-9 and T helper type 9 cells in rheumatic diseases

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Introduction

The role of CD4⁺ T cells in the pathogenesis of rheumatic diseases has been well established [1,2]. The differentiation of naive CD4⁽⁺⁾ T cells into different cell subsets with distinct biological activities is determined largely by their interaction with dendritic cells (DCs) in lymphoid organs. In particular, the development of a particular subtype of functional T cells seems to be related to the interaction with the surrounding microenvironment, which include cellular interactions and the release of specific cytokines and transcription factors [3,4]. Different subsets of CD4⁺ T cells are defined usually by the specific expression of transcription factors, cytokine receptors and their released cytokines. T helper type 1 (Th1), Th2, Th17, T follicular helper and derived regulatory T cells are well-known CD4⁺ T cell subsets [4,5].

Interleukin (IL)-9, a cytokine with pleiotropic functions in the immune system [6,7], was identified originally as a T cell growth factor belonging to the common γ -chain-receptor cytokine family. IL-9 signals through a heterodi-

Summary

Interleukin (IL)-9 is a 28-30 kDa monomeric glycosylated polypeptide belonging to the IL-7/IL-9 family of proteins that bind to a composite receptor consisting of the private receptor IL-9R and the IL-2 receptor, gamma (IL-2RG), a common gamma subunit shared by the receptors of many different cytokines. The IL-9R is expressed widely and IL-9 impacts a number of effector cells, such as effector T cells, B cells, innate lymphoid cells, mast cells, polymorphonuclear cells, epithelial cells and smooth muscle cells, playing an important role in regulating inflammatory immunity. The critical role of IL-9 in promoting cellular and humoral immune responses makes it an important focus of potential therapeutic interventions. Recently, a defined subset of T helper type cells, Th9 cells, has been identified by the potent production of IL-9. The involvement of the Th9 cell subset has been described in many types of inflammatory diseases, namely atopic diseases, helminth infections, experimental autoimmune encephalomyelitis and ulcerative colitis. In this review, we summarize the IL-9 biological activities, highlighting roles for IL-9 and Th9 cells in rheumatoid and psoriatic arthritis, systemic vasculitis, systemic lupus erythematosus and systemic sclerosis.

Keywords: IL-9, psoriatic arthritis, rheumatoid arthritis, SLE, systemic sclerosis, vasculitis, Th9 cells

meric receptor composed by a specific IL-9 receptor chain (IL-9R α) and the common γ -chain (also known as IL-2R γ) [8]. IL-9R α is expressed on immune cells and its activation also promotes mast cell growth [8,9] and accumulation in inflamed tissues, innate lymphoid cells (ILC) survival and B cell immunoglobulin (Ig)E class-switch recombination [9–11] (Fig. 1 shows IL-9R-expressing cells). IL-9R α is also expressed on non-haematopoietic cells such as airway and intestinal epithelial cells, smooth muscle cells and keratinocytes [12,13].

IL-9 production was associated first with the Th2 phenotype and many of the preliminary functions of IL-9 were studied in models of Th2-associated immunity. Recently, a defined subset of T helper cells called Th9 has been identified by the potent production of IL-9 [14–16]. Th9 cells develop from naive T cells in the presence of transforming growth factor (TGF)- β , IL-4 and thymic stromal lymphopoietin (TSLP) (Fig. 1) and require PU.1 and the interferon regulatory factor-4 (IRF-4) as specific transcription factors [7,17].

Although a role of the Th9 cell subset has been described recently in many types of inflammatory diseases (namely, atopic diseases, helminthic infections, experimental autoimmune encephalomyelitis and inflammatory bowel diseases) [18,19], their role is still largely unknown in the pathogenesis of rheumatic disease. This paper aims to review the evidence indicating that Th9 cells may contribute to the pathogenesis of autoimmune-related diseases, due to the recent demonstration of a possible role of Th9 T cells and of their related cytokine IL-9 in rheumatoid arthritis (RA), psoriatic arthritis (PsA), systemic vasculitis, systemic lupus erythematosus (SLE) and systemic sclerosis (SSc) [20–28] (Fig. 1 shows the possible tissue target of Th9-mediated inflammation).

IL-9 signalling

IL-9 is a protein of 144 amino acids with a secretory signal sequence of 18 amino acids. On target cells, IL-9 binds to IL-9R, a heterodimeric protein composed by IL-9R α belonging to the haematopoietin superfamily, and the IL-2R- γ , a common subunit shared by different cytokine receptors. Binding of IL-9 with the cognate receptors leads to the activation of signal transducer and activator of transcription-1 (STAT-1), STAT-3 and STAT-5 [29,30]. IL-9R is expressed on effector T cells but not naive T cells. Th2, Th17 and Th9 display the highest IL-9R expression. In asthmatic patients IL-9R is also expressed on mast cells and polymorphonuclear cells [31]. Several studies also suggest the expression of IL-9R on lung and intestinal epithelial cells and on the surface of smooth muscle cells (Fig. 1) [31].

Th9 polarization

The differentiation of Th9 cells requires the activation of IL-2/STAT-5 and IL-4/STAT-6 signalling and TGF- β , which acts by inducing redirection of naive T cells from a Th2 to Th9 cell differentiation pathway (in Fig. 1 it is summarized the differentiation pathway of Th9 cells) [7]. It has been demonstrated recently that, in primary cell cultures, the addition of TSLP led to an increase in IL-9 production from human and mouse Th9 cells, and induced an increase in STAT-5 activation and binding to the IL-9 promoter [31]. Th9 differentiation requires PU.1, IRF-4 and the B cell activating transcription factor-like (BATF), as specific transcription factors that bind to and activate the IL-9 locus [7]. Besides these cytokines, other signalling pathways seem to act by enhancing IL-9 production by Th9 cells, such as IL-25 and IL-33, by activating nuclear factor kappa B (NF- κ B), type I interferons and IL-1 β by inducing STAT-1/IRF-1 expression [7]. Furthermore, the interaction between T cells and antigen-presenting cells (APC) through the T cell receptor (TCR)/peptide–major histocompatibility complex (MHC) class II, CD28/CD80, OX40/OX40L,

NOTCH/DLL and Jagged may have an important role in Th9 differentiation [7].

IL-9 and Th9 cells in RA

RA is a chronic autoimmune and inflammatory systemic disease that primarily affects synovial joints. It is characterized by the aberrant expression of several proinflammatory cytokines and recruitment of autoreactive lymphocytes into inflamed tissues [32]. T and B cells are involved predominantly in the pathogenesis of RA, together with the orchestrated interaction of proinflammatory cytokines such as TNF- α , IL-6, IL-1 β and IL-17 [32].

IL-9 has been demonstrated to be increased significantly in the sera of RA patients not related to disease duration, disease activity score (DAS28), health assessment questionnaire (HAQ), rheumatoid factor positivity or erosions on radiography [25]. High levels of IL-9 have been also detected in patients' first-degree relatives' sera with RA or with asymptomatic RA-related autoimmunity [26,33]. Furthermore, IL-9, together with IL-6, is the most responsible in the cytokine score calculated by summing all cytokine/chemokine levels, weighted by their regression coefficients for RA–autoantibody association [26]. IL-9 levels are also elevated significantly in the synovial fluid of RA patients compared to osteoarthritis (OA) and IL-9 promotes proliferation and survival of synovial fluid CD3⁽⁺⁾ T cells of RA patients, the proliferation of T cells being dependent upon the PI3K/Akt/mTOR signalling pathway [34].

IL-9 has been reported to be over-expressed significantly in synovial tissue of RA patients and correlated with the degree of tissue inflammation [24]. IL-9 expression has been found mainly among synovial fibroblasts and infiltrating mononuclear cells. Th9 cells were demonstrated to be the main source of IL-9 in both synovial tissues (Fig. 1) and peripheral blood of RA, given the predominant co-expression of IL-9 with PU.1 [24]. Th9 polarization in RA synovium was accompanied by a significant up-regulation of IL-4, TGF- β and TSLP correlated directly with the number of IL-9-positive cells. Interestingly, Th17 also produced IL-9 in RA synovium, although to a lesser degree [24].

RA synovial tissue may display a different topographical distribution of the inflammatory infiltrates: diffuse infiltrates of inflammatory cells without aggregates of specific microstructures, T and B cell aggregates without germinal centre (GC) organization and aggregation of inflammatory cells in secondary follicles with GC formation [32]. IL-9 and IL-9R expression has been proved to be correlated directly with the degree of inflammatory infiltrate and lymphoid organization in RA patients [24]. The strong connection between IL-9/IL-9R axis activation and the induction of autoimmune responses has been highlighted recently due to the potential role of IL-9 in T cell-dependent B cell differentiation, expansion and antibody production [10], as well as by the evidence that IL-9R is

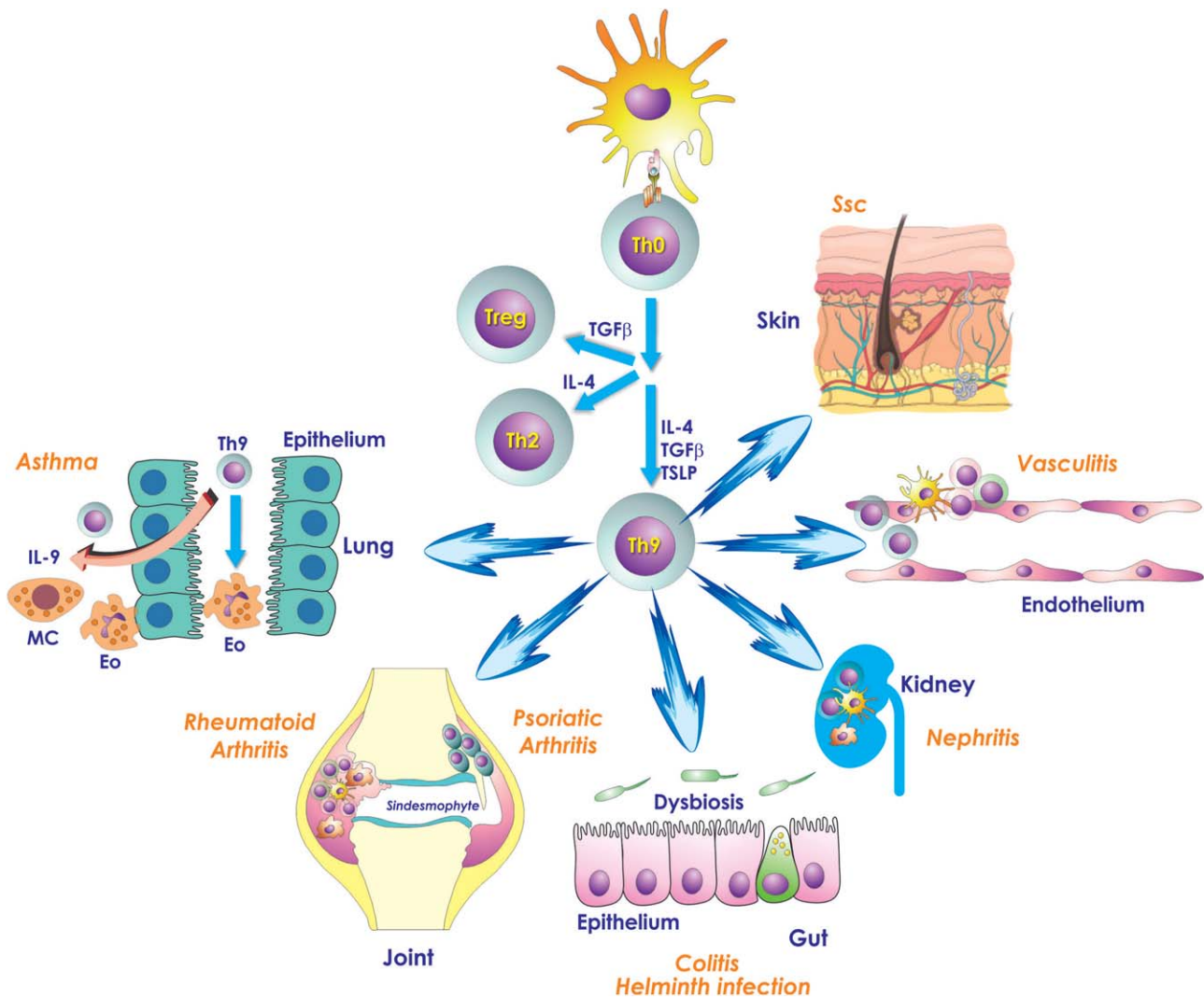


Fig. 1. T helper type 9 (Th9) cells develop from naive T cells in the presence of transforming growth factor (TGF)- β , interleukin (IL)-4 and thymic stromal lymphopoietin (TSLP). Aberrant Th9 activation may cause the onset and progression of autoimmune and chronic inflammatory diseases such as asthma, rheumatoid and psoriatic arthritis, chronic gut inflammation, kidney diseases, systemic vasculitides and systemic sclerosis. MC = mast cells; Eo = eosinophils; T_{reg} = regulatory T cells.

expressed in GC B cells where its stimulation induces STAT-5 activation [35]. The correlation between IL-9/IL-9R and lymphoid organization could confirm the importance of IL-9- and IL-9-producing cells in the induction of autoimmunity in RA.

Autoimmunity in RA is characterized by aberrant immune responses to autoantigens that have been modified post-translationally by citrullination [36,37]. Citrullinated arthritogenic aggrecan peptide has been considered to be a biomarker of RA, having been identified in the peripheral blood of RA patients, and has shown to stimulate a specific CD4 response [38–40]. It is considered a candidate autoantigen for RA, being one of the principle proteoglycans for cartilage extracellular matrix functioning in cushioning synovial joints, self-antigens that share a common sequence at position 70–74 of the human leucocyte antigen D-

related (HLA-DR) β -chain in 90% of RA patients [38,41]. In RA patients, stimulation with the arthritogenic aggrecan peptide resulted in a strong and significant expansion of Th9 cells. Together, these results suggest the existence of an association between the emergence of autoreactive Th9 cells and the occurrence of synovial inflammation and citrullination process [24].

IL-9 and Th9 cells in psoriatic arthritis

PsA is a chronic inflammatory disease affecting the spine or peripheral joints of patients with psoriasis or with a positive personal or familial history of skin manifestations [42]. PsA has long been considered as a Th1-mediated disease, with interferon (IFN)- γ and IL-12 as signature cytokines [43]. Recently, however, it has been shown that both

innate and adaptive immunity, leading to the activation of the IL-23/Th17 axis, may contribute to the initiation of tissue inflammation [44–46]. More recently, IL-9 has been also demonstrated to be involved in the pathogenesis of psoriasis [47]. In particular, Th9 cells that are increased in the skin lesions of psoriasis seem to have a specific tropism for the skin and their aberrant activation may contribute to skin inflammatory diseases [48].

Psoriasis and PsA have been demonstrated to be linked strongly to gut inflammation [49]. Several studies have highlighted the importance of subclinical gut inflammation in spondyloarthropathy pathogenesis, indicating the inflamed gut as the site where autoreactive and proinflammatory cells are activated and from which they are released in the systemic circulation, with consequent localization in extra-intestinal inflamed sites [50]. Few studies, however, have addressed specifically the immune responses in the gut of PsA patients.

PsA subclinical gut inflammation is characterized histologically by a higher number of infiltrating inflammatory cells, which are organized frequently in lymphoid follicles. Immunologically, the ileum of PsA patients shows a clear and defined IL-23/Th17 response, as shown by both IL-23 and IL-17 strong up-regulation, with a defective Th1 polarization [50]. However, IL-9 seems to mark gut inflammation in PsA specifically, given the impressive IL-9 expression observed in the gut of PsA patients [27]. IL-9 is produced particularly by inflammatory cells and epithelial cells that were demonstrated to be Paneth cells (PC), as co-expression of IL-9 and α -defensin 5 (a specific marker of PC) has been observed. In PsA gut, PC also express the IL-9R and after *in-vitro* stimulation with recombinant IL-9, produce high levels of specific peptides such as α -defensin 5 and cytokines such as IL-23, indicating the occurrence of an autocrine loop involving IL-9 [27]. PC are highly specialized small intestine epithelial cells; they are located precisely at the bottom of Lieberkühn crypts and are involved in innate immune responses and anti-microbial host defence, thus contributing to the maintenance of the gastrointestinal barrier [51]. A specific microbiome signature has been demonstrated recently in PsA patients, indicating a role for dysbiosis in the pathogenesis of PsA [52]. The specific release of IL-9 by PsA PC could be relevant in this context, as it may represent a significant immune link between innate and adaptive responses.

Th9 polarization also characterizes the synovium (Fig. 1) and the peripheral blood of PsA patients, and the percentage of circulating Th9 cells is correlated significantly with disease activity [27]. Moreover, Th9 cells, isolated from both synovium and peripheral blood of PsA patients, express the intestinal homing receptor α 4 β 7, indicating that these cells are probably activated in the gut and recirculate into sites of inflammation [27]. Interestingly, we have recorded that circulating Th9 cells in PsA decreased after anti-TNF and ustekinumab treatment, leading us to

speculate that the clinical improvement observed in PsA patients treated with these classes of drugs could be, at least in part, referred to the modulation of Th9 response [27]. Together these findings indicate that PsA may be characterized by IL-9 and Th9 polarization, suggesting that these cells may represent a future therapeutic target.

IL-9 and Th9 cells in large-vessel vasculitis

Large-vessel vasculitides (LVV) are characterized by the autoimmune inflammation of medium and large arteries leading to occlusion of the lumen with ischaemic damage of dependent organs [53]. Several effector cytokines involved in both innate and adaptive immunity have been identified in vasculitic lesions [54]. However, among the different immune pathways characterizing immune responses, the IL-6–IL-17 axis and the IL-12–IFN- γ axis seem to play a fundamental role in LVV pathogenesis [54].

Giant cell arteritis (GCA) is the prototype of LVV and is characterized by a range of histological patterns of vascular wall injury [54]. Beyond the classic transmural inflammation (with or without giant cells), two other different histological aspects of GCA have been described [55]: small vessel vasculitis (SVV) defined as inflammation of the small vessels external to the temporal artery adventitia and vasa vasorum vasculitis (VVV), defined as isolated inflammation of temporal artery vasa vasorum [55].

Th1 and Th17 subsets of effector T cells have been demonstrated clearly to participate in the pathogenesis of GCA [54,56]. Although Th2-derived cytokines have been demonstrated previously to be consistently absent [54], the IL-33 pathway seems to be over-expressed in the inflamed arteries of GCA patients and accompanied by a strong M2 macrophage polarization [57]. Beyond the role in promoting the Th2 response, IL-33 has also been associated recently with the secretion of IL-9 by human CD4⁺ T cells isolated from peripheral blood [58–60], apparently indicating a potential role of Th9 cells in the pathogenesis of GCA.

Analysis of IL-9 and IL-17 expression in different histological subsets of GCA has demonstrated that a different immunological polarization characterizes different histological patterns of GCA [20]. The different expression of cytokines in different patterns of tissue pathology may be of immunological relevance. Multiple effector T cell subsets (such as Th17 and Th9) may induce GCA independently of each other, probably by overlapping but with distinct mechanisms. IL-9 and Th9 cells, in fact, immunologically mark the inflamed arteries of GCA patients with inflammation restricted to the peri-adventitial small vessels. Conversely, a more intense Th17 polarization, with weak IL-9 expression, predominates in the GCA arteries with inflammation restricted to the vasa vasorum. Interestingly, a concomitant Th9 and Th17 response is observed in arteries

displaying transmural inflammation and especially in those arteries with granulomatous reaction [20].

Inflammatory cells infiltrating the artery wall, giant cells, endothelial cells of vessels scattered through the inflammatory infiltrates and among vascular smooth muscle cells showed intense positivity for IL-9 [20]. IL-9 tissue expression was also correlated significantly with the intensity of the systemic inflammatory response, including erythrocyte sedimentation rate (ESR) and C-reactive protein (CRP) [20]. Infiltrating inflammatory cells producing IL-9 were demonstrated to be Th9 cells, considering the significant co-localization observed between IL-9 and PU.1 (the specific transcription factor for Th9 cells), suggesting the prevalent Th9 phenotype of IL-9 mononuclear-producing cells (Fig. 1). Notably, when present, giant cells scattered in the inflamed arteries showed an intense co-expression of IL-9 and IL-17 [20].

Th9 cells are a distinct subpopulation of CD4⁺ effector T cells that require TGF- β , IL-4 and TSLP for their differentiation [7]. GCA arteries with higher expression of IL-9 (transmural inflammation with granulomatous reaction and SVV) displayed intense over-expression of TGF- β , IL-4 and TSLP, suggesting that the upstream Th9 cytokine network is also up-regulated in GCA [20]. The biological effects of IL-9 are mediated by the functional IL-9 receptor complex, that is composed of this protein as well as the IL-2 receptor gamma (IL-2RG), a common gamma subunit shared by the receptors of many different cytokines [7]. The activation of this receptor leads to the activation of various Janus kinases (JAK) and STAT proteins [8] which connect to different biological responses [6]. In GCA, IL-9R positivity was observed among endothelial cells of vessels distributed in the inflammatory infiltrates and neutrophils [20]. Functional expression of IL-9R receptor by human neutrophils has been demonstrated in asthmatic patients, with an important role in IL-8 release [61]. IL-9/IL-9R over-expression in GCA was accompanied by a significantly high expression of IL-8 and IL-8R, especially in patients with transmural inflammation, providing a functional immunological link between IL-9R expression and neutrophil activity, thus highlighting the importance of neutrophils in the pathogenesis of GCA [20].

Studies conducted on small series have shown previously that IL-17A-expressing cells are reduced dramatically in specimens obtained from glucocorticoid-treated patients [56]. Differently from IL-17, IL-9 was modified only marginally by glucocorticoids. This apparent IL-9 low sensitivity to glucocorticoid inhibition may indicate a potential role for Th9 cells in steroid-resistant GCA [20].

IL-9 and Th9 cells in anti-neutrophil cytoplasmic antibodies (ANCA)-associated vasculitis

Granulomatosis with polyangiitis (GPA) is a systemic necrotizing vasculitis that affects small- and medium-sized vessels in many organs associated with granulomatous

inflammation and the presence of ANCAs directed against proteinase 3 (PR3) [62,63]. PR3 has been demonstrated to be expressed at the plasma membrane of neutrophils and may act by interfering with induction of anti-inflammatory mechanisms following phagocytosis of these cells by macrophages and/or by facilitating anti-PR3 ANCA binding, leading to neutrophil activation and consequent tissue inflammation [64]. Interestingly, in different murine models, membrane-associated PR3 on apoptotic cells triggered secretion of inflammatory cytokines and chemokines, thus inducing a specific microenvironment instructing plasmacytoid dendritic cells (pDC) to promote a specific effector T cell subset. In particular, pDCs exposed to apoptotic cells expressing membrane PR3 in a TGF- β -rich environment induce the generation of Th9 cells. Th9 and Th2 cell polarization [65]. Interestingly, PBMCs obtained from active, untreated systemic GPA patients also displayed skewed Th9 and Th17 responses, revealing a GPA-specific mechanism of immune polarization [65].

Th9 cells in systemic lupus erythematosus

SLE is a disorder of generalized autoimmunity characterized by the dysregulation of cellular and humoral immune responses, the break in immune tolerance to self-antigens, polyclonal B cell activation and the production of autoantibodies. Autoantibody deposition and inflammatory cell infiltration in target organs such as kidneys and brain characterizes SLE. It has been reported that aberrant T lymphocyte activation and altered cytokine production are important contributors to SLE pathogenesis. However, so far the exact mechanisms that lead to the development of SLE remain undefined.

Only few studies have been carried out on Th9 cells and their related cytokine, IL-9, in human SLE. These studies have shown an increase in the Th9 cell subset and IL-9 serum levels on active SLE. In lupus-prone mice Th9 cells accumulated in the spleen and in the kidneys (Fig. 1) and their number is related closely to the presence of germinal centres [23]. In addition, *in-vivo* treatment with neutralizing anti-IL-9 antibodies was able to reverse serum anti-dsDNA antibody levels [23].

IL-9 can be produced by different T cell subsets that include CD8⁺ cells, invariant natural killer T (NK T) cells, $\gamma\delta$ T cells, Th17 cells and innate lymphoid cell group 2 (ILC2) [50–53]. IL-9 is also known to induce TH17 differentiation and IL-17 production, and these cytokines may work together synergistically in promoting SLE pathogenesis. Despite these observations, the role of Th9 and IL-9 in SLE patients remains unclear.

IL-9 and Th9 cells in systemic sclerosis

Expression of IL-4, TGF- β and TSLP has been implicated in fibrosis and remodelling in SSc [66–68]. Furthermore,

Th9 and IL-9 seem to increase tissue fibrosis, as demonstrated in the liver, where IL-9 induces hepatic fibrosis and an exacerbated disease end-point [69]. Serum IL-9 level has been demonstrated to be increased in patients with SSc and associated with a lower frequency and severity of pulmonary fibrosis, indicating IL-9 as a protective factor in SSc [28]. However, the presence of high IL-9 levels and the strong expression of Th9 polarizing cytokines in SSc raise the question of whether Th9 cells might participate in SSc pathogenesis. In this regard, strong expression of IL-9 and IL-9R has been demonstrated recently in skin tissue of patients with diffuse SSc and correlated with the modified Rodnan skin score and the presence of pulmonary fibrosis [70]. In particular, IL-9 expression was observed mainly in the context of infiltrating mononuclear cells and in the keratinizing squamous epithelium of skin from SSc patients [70]. The majority of IL-9-producing cells in the skin were identified as Th9 and Th17 cells. Among peripheral blood mononuclear cells, Th9 cells were the major source of IL-9, being expanded significantly in SSc patients compared to controls [70]. The participation of IL-9 and Th9 in the process of inflammation and fibrosis action in SSc, however, is still not clear, and further studies are required to better clarify their role in modulating tissue fibrosis.

Conclusions

IL-9 and Th9 cells seem to be involved in the immunopathology of several human diseases. In systemic rheumatic diseases this axis is activated and involved potentially in the triggering and/or maintaining inflammation. The targeting of IL-9/Th9 cells could be considered a productive strategy in treating systemic autoimmune diseases. However, further studies are required to understand more clearly the specific contribution of this axis in the pathogenesis of different systemic autoimmune diseases.

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Disclosure

The authors declare no conflicts of interest.

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