

## Review

## Targeting the IL17 Pathway for the Prevention of Graft-Versus-Host Disease

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## Article history:

Received 14 November 2013

Accepted 10 February 2014

## Key Words:

Allogeneic stem cell transplantation  
GVHD  
Interleukin 17  
Dendritic cell  
Danger-associated molecular patterns (DAMP)  
Pathogen-associated molecular patterns (PAMP)  
Th17 cell

## A B S T R A C T

Graft-versus-host disease (GVHD) is still a major complication of allogeneic stem cell transplantation (allo-SCT). The pathophysiology of GVHD is a multistep process initiated by tissue damage and proinflammatory cytokine cascades induced by the pretransplantation conditioning therapy. This eventually results in Th1-driven tissue damage. However, increasing evidence indicates the involvement of IL17-producing T cells in GVHD pathogenesis. Both CD4<sup>+</sup> and CD8<sup>+</sup> IL17-producing T cells are suspected of initiating the Th1 response and aggravating tissue inflammation, resulting in full-blown GVHD. In this review, we discuss the involvement of IL17-producing T cells in GVHD and the factors involved in their expansion, differentiation, and activation. Different dendritic cell (DC) subsets, such as plasmacytoid DCs and DC NK lectin group receptor 1<sup>+</sup> myeloid DCs have the capability to stimulate Th/Tc17 responses through the release of cytokines. Pivotal cytokines include IL1 $\beta$ , IL6, IL23, and TGF $\beta$ , which are known to drive differentiation and expansion of IL17-producing T cells, and these cytokines are highly elevated in patients after allo-SCT. Potent activators of these DC subsets are motifs that are released upon tissue damage and microbial exposure during allo-SCT. These motifs aggravate the Th/Tc17 response via the activation of various pathogen recognition receptors, thereby initiating and perpetuating GVHD. A more comprehensive understanding of the factors and DC subsets driving the IL17 pathway will result in developing and testing novel therapeutic approaches for the prevention of GVHD.

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## INTRODUCTION

Graft-versus-host disease (GVHD) is still a major complication and limitation of allogeneic stem cell transplantation (allo-SCT) [1,2]. The pathophysiology of GVHD is a multistep process involving tissue damage and proinflammatory cytokine cascades induced by the pretransplantation conditioning therapy [2,3]. Furthermore, innate immune activation caused by perturbed host-microbe interactions at epithelial barriers increases inflammation and antigen presentation [4]. This results in an excessive inflammatory environment in which donor-derived CD4<sup>+</sup> and CD8<sup>+</sup> T cells become activated by antigen-presenting cells. In addition, T cell trafficking towards inflamed GVHD-prone organs, such as skin, lung, the gastrointestinal tract, and liver, is strongly augmented. Further, tissue destruction in these organs occurs in the case of presentation of ubiquitous or epithelial expressed allo-antigens to infiltrating alloreactive T cells. Moreover, dendritic cells (DC), macrophages, and other T cell subsets are recruited, resulting in further enhancement of GVHD. In particular, T helper (Th) 1-type CD4<sup>+</sup> T cells and T cytotoxic (Tc)-1 type CD8<sup>+</sup> T cells play an important role in the effector phase of GVHD pathophysiology [1,2,5]. However, recent

studies indicate that other T cell subsets, such as Th/Tc17 cells, are involved in the initiation and aggravation of GVHD [6–9]. In this review, we will discuss the role of these IL17-producing T cells in GVHD development and the potential therapeutic strategies targeting the IL17 pathway to diminish or prevent this detrimental complication.

## REQUIREMENTS FOR DIFFERENTIATION OF IL17-PRODUCING CELLS

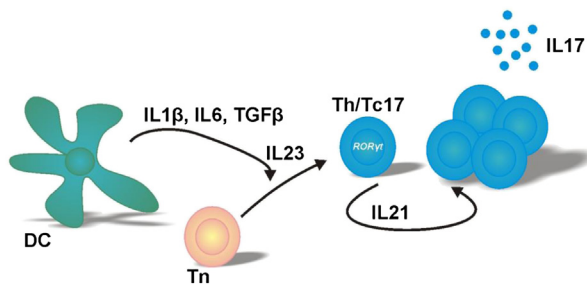
Generally, IL17-producing T cells differentiate from naïve T cells, though plasticity of other T cell subsets into IL17-secreting T cells has been shown [10]. Activation of the transcription factor retinoic acid-related orphan receptor (ROR) $\gamma$ t is essential for IL17-producing CD4<sup>+</sup> (Th17) and CD8<sup>+</sup> (Tc17) T cells [11] that can be achieved via various routes, though it generally occurs via the STAT3 signaling pathway. STAT3 is not only responsible for the activation of ROR $\gamma$ t, but it can also bind and function as a promoter to other genes involved in this process, such as *IL17* [12].

Various cytokines play a major role in the activation of STAT3, ROR $\gamma$ t, and Th/Tc17 differentiation (Figure 1). The importance of IL1 $\beta$  and IL6 has been shown in several studies, both in mouse models as well as in vitro cultures of human T cells [13,14]. The additional role of TGF $\beta$  in this process is still unclear, mainly due to differences in mouse and human studies. Most mouse studies showed that TGF $\beta$  is essential in Th/Tc17 differentiation. However, human Th/Tc17 cells can be generated in vitro without the presence of TGF $\beta$  [13].

Financial disclosure: See Acknowledgments on page 757.

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<http://dx.doi.org/10.1016/j.bbmt.2014.02.007>



**Figure 1.** Th/Tc17 cell differentiation. Cytokines produced by dendritic cells differentiate naive T cells (Tn) towards Th/Tc17 cells. DC indicates dendritic cell; Tn, naive T cell; Th/Tc, helper T cell/cytotoxic T cell.

Additional studies in humanized mouse models could give more insight in this process. Generally, in the presence of IL1 $\beta$  and IL6 (and possibly TGF $\beta$ ), while IFN $\gamma$  and IL4 are absent to induce Th1 or Th2 skewing, differentiation to IL17-producing T cells is initiated. In addition, IL23 plays a pivotal role in the commitment and maintenance of Th17 cells [14].

Furthermore, IL23 was shown to play a major role in the generation of pathogenic Tc17 cells, as IL23-treated Tc17 cells were potent inducers of autoimmune diabetes, whereas Tc17 cells treated with TGF $\beta$  and IL6 were not [15]. Finally, the cytokine IL21 is also involved in Th17 differentiation [16]. Upon initial activation of STAT3 by IL6, both IL21 and IL23, along with TGF $\beta$ , induced IL17 expression via the activation ROR $\gamma$ t. Moreover, the cytokine IL21 is also involved in self-maintenance, as this cytokine is produced by the Th/Tc17 cells themselves.

Multiple microenvironmental factors could also influence Th17 differentiation. For instance, the presence of retinoic acid in the microenvironment inhibits Th17 differentiation and induces regulatory T cell (Treg) formation [17]. On the other hand, several labs have described a role for the aryl hydrocarbon receptor in Th17 differentiation [18]. These studies showed that aryl hydrocarbon receptor ligands increase Th17 responses from naive T cells and promote autoimmunity in mouse models. The various cytokines required for Th/Tc17 differentiation are released from diverse immune (eg, myeloid DC [mDC] and plasmacytoid DC [pDC]) and nonimmune antigen-presenting cells on activation of pathogen recognition receptors (PRRs) by motifs released upon tissue damage and/or microbial exposure, including various danger-associated molecular patterns (DAMPs) and pathogen-associated molecular patterns (PAMPs), respectively. Because in the human body epithelial barriers are inhabited by multiple microbes, both Th17 and Tc17 cells are of particular importance in the protection against extracellular microbes and defense mechanisms in the epithelial barriers of skin, gut, and lung. The release of IL17 has, indeed, been shown to enhance barrier defense, for example, by the production of antimicrobial peptides. Consistently, Th/Tc17 cells have been described to play a major role in the initiation of autoimmune disorders and other inflammatory conditions that originate at epithelial barriers, such as Crohn's disease and psoriasis [19]. As GVHD predominantly affects the gut, skin, and liver, Th/Tc17 cells are also suspected to be major players in the onset of GVHD.

## TH/TC17 CELLS AND THEIR ROLE IN GVHD

### Indicative Data from Animal Models

Several mouse studies demonstrated the involvement of Th/Tc17 cells in the onset and persistence of GVHD [7,20,21].

Carlson et al. showed development of lethal GVHD upon injection of in vitro-generated Th17 cells in a bone marrow transplantation mouse model [7]. In addition, adoptive transfer of IL17 $^{-/-}$  T cells delayed the occurrence of GVHD compared with wild type (WT) T cells [21]. However, the precise role of Th/Tc17 cells in GVHD is still debated. For instance, Yi et al. reported initiation of Th1-driven GVHD in the absence of IL17 [22]. To the contrary, Yi et al. also observed IL17-driven GVHD when the Th/Tc1 cytokine IFN $\gamma$  was absent [23]. A joint event was described by Yu et al., who observed lower GVHD incidence while sparing the graft-versus-tumor (GVT) effect, but only when both the Th/Tc17 driving transcription factor ROR $\gamma$ t as well as Th/Tc1-inducing transcription factor t-bet were targeted simultaneously [24]. Targeting only ROR $\gamma$ t elevated IFN $\gamma$  levels, whereas IL17 was highly increased when t-bet was abolished. However, targeting both resulted in the lack of both cytokines and the absence of GVHD.

Alternatively, the observed effect on GVHD could also be explained by increased levels of Tregs in these mice. Interestingly, Fulton et al. attenuated GVHD by only targeting ROR $\gamma$ t, though in this mouse model no increase in IFN $\gamma$  was observed as a consequence of lacking IL17 [25]. In addition, targeting ROR $\gamma$ t in this mouse model only affected serum IL17, whereas Yu et al. showed disturbed levels in several GVHD target organs [24,25]. Taking into account the differences in human and mouse Th/Tc17 differentiation, more insight in this process could be revealed in humanized mouse models.

In addition to IL17, Th/Tc17 cells could also play a role in GVHD via production of IL22 [26]. IL22, which is related to the IL10-family, has a paradoxical role in inflammation with both protective and inflammatory effects. Though it was shown that IL22 produced by innate lymphoid cells protected intestinal stem cells from immune-mediated damage during GVHD [27], IL22 is also involved in the pathogenesis of several inflammatory diseases, such as rheumatoid arthritis and psoriasis [28,29]. Interestingly, a recent study revealed a possible role of IL22 in GVHD, as adoptive transfer of IL22 $^{-/-}$  instead of WT T cells reduced acute GVHD severity [30]. However, IL17 production of IL22 $^{-/-}$  T cells was not affected, and an increase in Tregs was observed in these mice, which could also contribute to the reduced GVHD. In conclusion, collective findings in experimental mouse models suggests that Th/Tc17 could contribute to the pathophysiology of GVHD through IL17, and possibly IL22, production.

### Studies in Allo-SCT Patients

In addition to their role in mice, a key role of Th/Tc17 cells in human GVHD pathogenesis has been reported [6,8]. Interestingly, increased levels of circulating Th17 cells have been described in GVHD patients [8,31]. For instance, Liu et al. showed increased circulating IL17-producing CD4 $^{+}$  T cells in patients at the onset of GVHD, compared with time-matched controls [31]. Furthermore, a single nucleotide polymorphism (SNP) in the IL23 receptor, important in Th/Tc17 differentiation, has shown a protective role in the development of acute GVHD in 2 independent studies [32,33]. However, as the functional consequence of the SNP is yet unknown, it cannot be stated whether IL17 differentiation would be inhibited or enhanced. Although the potential involvement of Th17 cells in GVHD has been proposed by multiple studies, Broady et al. found an expansion of Th1, rather than of Th17 cells, in GVHD-affected skin [34].

However, Th17 cells have been shown to coproduce IFN $\gamma$  or revert completely into Th1 cells [35]. Regarding this proposed plasticity of Th17 responses, other (surface) markers would be more reliable to study the involvement IL17-producing T cells in GVHD. Using CD161, the distinguishing surface marker for both CD4<sup>+</sup> and CD8<sup>+</sup> T cells capable of producing IL17 [36], we recently showed decreased levels of circulating Th/Tc17 cells in GVHD patients, although they infiltrated GVHD-affected skin and gut [9]. This was in concordance with data published by Bossard et al., who reported significantly higher absolute numbers of Th/Tc17 cells in GVHD-affected intestinal mucosa, using the markers CD161, ROR $\gamma$ t, and CC chemokine receptor (CCR)6 [6]. This suggests that Th/Tc17 cells could be involved in the onset of GVHD via their migration potential toward GVHD target organs, thereby exerting a proinflammatory effect, as well as increasing the recruitment of alloreactive Th/Tc1 cells and other immune cells. However, many factors, such as the conditioning regimen and different post-transplantation immune suppressive therapy, will have a prominent impact on expansion kinetics and involvement of Th/Tc17 and Th/Tc1 responses in GVHD in patients.

#### TH/Tc17 TRAFFICKING TO GVHD TARGET ORGANS

Migration towards GVHD target organs could take place via multiple chemokine axes, as Th/Tc17 cells display high levels of the chemokine receptors CCR6 and CXCR3 [26]. Interestingly, we showed that a SNP in CCR6 present in the donor resulted in lower GVHD incidence [37]. Its ligand, CCL20, is constitutively expressed in the GVHD target organs, such as the liver, colon, small intestine, lung, and skin [38].

Additionally, damage to the epidermal permeability barrier, as well as stimulation with IL1 $\beta$ —both known to be elevated after conditioning of allo-SCT patients—results in upregulation of CCL20 expression [9,39,40]. We, indeed, showed that GVHD-affected skin and gut display high expression of CCL20, both by keratinocytes and infiltrating immune cells [9]. Furthermore, increased expression of the CXCR3-ligands, CXCL9 and CXCL10, has also been described in serum of GVHD patients and in their affected tissues [41]. These chemokines, which are strongly upregulated upon inflammation, can attract migrating Th/Tc17 cells, resulting in their recruitment into GVHD-affected tissues, and therefore decreased levels of circulating CXCR3<sup>+</sup> T cells as described in GVHD patients [41,42]. This endorses the various migration routes of Th/Tc17 cells towards GVHD-target tissues.

#### CROSSTALK OF TH/Tc17 WITH TH/Tc1 CELLS

Upon recruitment of Th/Tc17 cells into GVHD tissues, other T cell subsets are attracted. Crosstalk between Th/Tc1 and Th/Tc17 cells in GVHD development is very likely, as has also been suggested in other immune-related diseases [43,44]. In this regard, we suggest that shortly after allo-SCT, Th/Tc17 cells could play an important role in initiating the GVHD response, which is thereafter overshadowed by a tremendous Th/Tc1 response, causing the main damage in the tissues. This hypothesis has been underscored by histological data showing that in early-stage mild gut GVHD, higher numbers of Th17 cells were found than in patients with severe GVHD after allo-SCT [45]. Possibly, Th/Tc17 and Th/Tc1 responses act either sequentially or simultaneously together in GVHD. This was also observed in inflammatory bowel disease, another IL17-driven disease, in which a synergy of Th1 and Th17 cytokines was observed [43].

The dependence of these subsets upon each other has been clearly demonstrated in different studies. In the absence of IFN $\gamma$  (with or without IL4), differentiation towards IL17-producing T cells was increased, resulting in augmented GVHD [23]. However, *visa versa*, lack of IL17 increased IFN $\gamma$  production, resulting in Th1-driven GVHD [22]. Taken together, crosstalk between Th/Tc17 and Th/Tc1 cells is apparent in the setting of GVHD, in which upon initiation of Th/Tc17 cells, Th/Tc1 cells will be potently activated and aggravate the disease. However, no hard evidence yet exist for this sequential T cell response and further research is warranted to understand dynamics and crosstalk of these 2 cooperating T cell subsets in GVHD.

#### FACTORS PROMOTING TH/Tc17 ACTIVATION AND EXPANSION AFTER ALLO-SCT

##### Role of DC Subsets

The role of DCs in GVHD development was established more than a decade ago by Shlomchik et al. [46]. Although it has been demonstrated that both DCs from the host as well as from the donor can contribute, less is known about the role of the different DC subsets in GVHD. The most common known DCs, the mDCs, have been shown to play a role in GVHD in several mouse models [46,47]. Especially, depletion of host mDCs in recipient mice has been shown to prevent GVHD. In addition, low levels of circulating mDCs in allo-SCT patients have been described to correlate with the development of severe acute GVHD, suggesting migration of these cells to GVHD sites [48].

Another DC subset, which has recently been associated with GVHD, are pDCs. Notably, increased levels of pDCs were detected in affected gut tissue of acute GVHD patients [6]. Also, a decrease in circulating pDCs was correlated with acute GVHD, suggesting active migration from the periphery into inflamed tissue [48]. Furthermore, pDCs are capable of producing high levels of IL1 $\beta$ , a key cytokine in the development of GVHD and the IL17 differentiation pathway [49,50]. Interestingly, it was shown that presence of IL1 $\beta$  resulted in increased IL17 production by CD161-expressing CD8<sup>+</sup> Tc17 cells [51]. Moreover, pDC leukemia is often associated with isolated cutaneous lesions due to skin accumulation of leukemic pDCs [52], suggesting their involvement in inflammatory diseases of the skin, such as GVHD.

Moreover, pDCs could be potentially activated via TLR9 by self-DNA, released upon pretransplantation conditioning. Although viral DNA can gain access to the intracellular TLR9 receptor, self-DNA does not gain direct entry to endosomes and TLR9 under normal conditions. However, in the presence of the endogenous cationic antimicrobial peptide LL37, pDCs can be also activated by self-DNA [53]. The main function of LL37 peptide is to prevent microbial invasion of damaged epithelial surfaces by directly killing a broad spectrum of pathogens, including bacteria, protozoa, fungi, and viruses. Besides this, it has the ability to bind self-DNA, released by, for instance, dying cells after the conditioning regimen, to form aggregates and condensed particles that are transported into pDCs, allowing TLR9 signaling [53,54]. LL37 is normally not expressed in healthy skin, but it is transiently produced by keratinocytes and released by infiltrating neutrophils in response to skin wounds or infections [55]. In psoriatic skin, LL37 has been identified as a key mediator in pDC activation [53]. As the conditioning regimen, containing chemotherapy and total body irradiation, probably results in increased LL37 and self-DNA exposure, and thereby robust

pDC activation, a Th/Tc17 response will be triggered, starting the inflammatory response of GVHD.

Another relevant DC subset possibly involved in the initiation of GVHD could be DC NK lectin group receptor 1 (DNNGR1)<sup>+</sup> CD141<sup>+</sup> mDCs. This DC subset, capable of producing IL6, IL12, TNF, and IFN, has the capacity to induce Th17 responses [56,57]. Recently, it was shown that these DNNGR1<sup>+</sup> DCs are activated upon interaction with filament actin, which has been shown to be strongly exposed upon cell damage [58]. Because the pretransplantation conditioning regimen of allo-SCT patients results in host tissue damage and, thereby, possible exposure of filament actin, DNNGR1<sup>+</sup> DCs could be activated and, thereby, promote IL17 responses.

Therefore, both pDCs as well as DNNGR1<sup>+</sup> DCs could play a key role in the activation of IL17-producing T cells and, thereby, the onset of GVHD.

### Role of Microbe and Damage Association Patterns

The role of innate immune activation by host microbe interactions at epithelial barriers in the pathogenesis of GVHD is increasingly recognized [4,59,60]. In 1971, it was shown by studies in chimeric mouse models that germ-free or gut-decontaminated mice showed reduced GVHD-related mortality [61,62]. Recently, the composition of the gut microbiota and changes that occurred therein after allo-SCT were shown to be pivotal in the pathogenesis of GVHD in both mice and humans, supporting previous observations [63].

Bacteria and fungi, and also viruses, have been associated with the occurrence of GVHD [2,64,65]. In addition, the use of antibiotics resulted in reduced anaerobic bacteria and a lower incidence of acute GVHD [66]. The fungus *Candida albicans* has also been shown to result in increased incidence of acute GVHD [67]. This was in line with the study of Marr et al. who showed that prophylactic fluconazole use prevented *Candida* infections and resulted in a lower GVHD incidence [68].

These microbes and their PAMPs, that potentially contribute to GVHD, are mainly present early after allo-SCT, which is during the initiation phase of GVHD [2]. Generally, DC activation by microbes can induce Th1, Th2, and/or Th17 responses, dependent on the type of antigen. However, most of these commensal microbes and pathogens can elicit strong Th17 responses. As several microbes elicit plastic immune responses during the progress from mucosal to systemic infection, with Th17 responses preceding Th1, it can be imagined that Th/Tc17 might be an early event in GVHD that precedes massive Th/Tc17-driven tissue damage. Microbes could influence Th17-mediated GVHD indirectly by activating innate immune cells via PRRs and, more directly and specifically, by eliciting CD4<sup>+</sup> and CD8<sup>+</sup> T cell responses through the several DCs subsets. The role of the latter in GVHD is unknown, but it is suggested by a role for microbe-specific CD4<sup>+</sup> Th17 and Th1 responses in colitis and syngeneic GVHD of the gastrointestinal tract [69,70]. In addition, Jankovic et al. showed that activation of the PRR NLRP3 by PAMPs and DAMPs, ie, uric acid, contributed significantly to the occurrence of intestinal GVHD [71]. The observation that this was mediated by IL1 $\beta$ , followed by a Th17 response especially indicates the role of Th17 cell as early reactor after activation signals.

Furthermore, it has been observed that Th17 cells respond to several microbial antigens [72,73]. More specifically, CD161-expressing T cells can be activated by *Escherichia coli* resulting in an inflammatory cytokine response [72]. In addition, IL17 secretion by CD4<sup>+</sup> T cells was observed upon stimulation with *Candida albicans* [73]. Plausibly, pDCs could

play a role in the activation of Th17 cells by microbes via TLR 7 and 9 [74]. In this regard, Yu et al. found increased IL17 production by T cells upon coculture with TLR7 stimulated pDCs [50]. In addition, similar data were observed when pDCs were activated with Influenza A or HIV-infected cells. Also, as viral infections by herpes simplex virus and cytomegalovirus have been associated with the occurrence of GVHD, activation of pDCs by these viruses could contribute to the initiation of the disease [64,65]. Furthermore, pDCs are activated by extracellular and intracellular bacterial DNA and RNA, supposedly via TLR7 and TLR9 after CD32-mediated uptake [75,76]. In an animal model for the Th17-driven disease multiple sclerosis, initiated by exposure to *Mycobacterium tuberculosis* and pertussis toxin, depletion of pDCs resulted in less severe clinical symptoms as well as less Th17 cells [49]. All together, this supports the notion that pDCs can initiate a Th/Tc17 response via activation by microbial antigens.

### Role of Cytokines

Pretransplantation conditioning regimen results in tissue damage with a subsequent rise of various (proinflammatory) cytokines [77,78]. Within the first weeks after allo-SCT, levels of IL1 $\beta$ , IL6, IL8, IL10, IL21, IL23, and TNF $\alpha$  are increased, which can remain high up to 12 weeks after allo-SCT [31,77]. Increased levels of IL6 have been reported in several studies and have been correlated with the occurrence of infections and GVHD [79,80]. Importantly, TGF $\beta$  appears also elevated during the first weeks after allo-SCT in patients with acute GVHD [78]. As described in previous paragraphs, these cytokines are mainly produced by DCs upon activation due to tissue damage and microbe exposure. Several of these cytokines are involved in the differentiation and expansion of Th/Tc17 cells, which could result in increased levels of Th/Tc17 cells after allo-SCT, as previously described by us and others [8,9].

Additionally, a trend towards higher IL17 serum levels has been observed in patients with acute GVHD [81]. This is in alliance with data published by Espinoza et al., who found that patients receiving a transplant from donors with a SNP in the promoter region of the IL17 gene were more prone to developing GVHD [82]. Interestingly, T cells with this allele produced more IL17 than those without. Furthermore, Lui et al. showed increased IL17 in the serum of GVHD patients; however, IL22, also produced by Th/Tc17 cells, was decreased in these patients [31].

In addition to IL17 and IL22, Th/Tc17 cells also produce IL21 to support self-renewal [16]. The lack of IL21R on T cells prevented GVHD while sparing the GVT effect [83]. Moreover, antibody blockade of IL21 also protected mice from the development of GVHD [84,85]. However, 2 out of these 3 models showed minimal responsive IL17 levels, and the observed effect could also be explained by the increased levels of Tregs [83,85]. Nevertheless, Hippen et al. showed decreased levels of IL17 upon blockade of IL21, although also in this study an increase of Tregs was observed [84]. In addition, given the importance of IL21 in T cell proliferation, blockade of IL21 also resulted in decreased numbers of alloreactive T cells, which could also contribute to the reduced GVHD response in this model. Taken together, the increased levels of cytokines induced by the conditioning regimen results in expansion and activation of Th/Tc17 cells after allo-SCT. This could be an important, initial driver of the development of GVHD, though the cytokines produced by the Th/Tc17 cells might not all contribute to GVHD development.



## TARGETING TH/TC17 RESPONSES TO PREVENT GVHD

Currently, GVHD treatment options rely largely on the suppression of the whole immune system, using immunosuppressive drugs such as cyclosporine A and mycophenolate mofetil. Though these treatments can reduce GVHD intensity in a proportion of the patients, treatment success remains modest, especially in patients with severe GVHD [86]. In addition, suppression of the whole immune system predisposes patients to an increased risk of infections and tumor relapse. Therefore, selective prevention of GVHD would be a far more favorable strategy. With more comprehensive understanding of which factors and T cells subsets are involved in GVHD development, improved targeting strategies could be exploited to prevent GVHD. Selective GVHD prevention will likely reduce the nonrelapse-related mortality, and, importantly, increase quality of life.

### Manipulation of T Cells in Graft and Donor Lymphocyte Infusion

Complete or partial depletion of T cells from the graft has the ability to reduce the occurrence of GVHD [87]. However, crude T cell depletion strategies will also reduce the therapeutic GVT effect. By more specific T cell depletion strategies, such as the depletion of only CD161<sup>+</sup>/Th/Tc17 cells, the GVT effect could be sustained by the non-GVHD-initiating T cells. Targeting the tissue homing properties, such as CCR6 or CXCR3, on these cells is a possibility, either by depletion or blockade of the homing capacity. However, this strategy will be less specific for Th/Tc17 depletion. In addition, depletion of the different DC subsets, such as pDCs and DNGR1<sup>+</sup> DCs, could also be a target, as specific depletion of DCs from the graft could inhibit the initiation of GVHD. However, to spare the GVT effect, depletion of specific DC subsets might be more favorable.

### In Vivo Manipulation of Th/Tc17 Cell Differentiation

In addition to manipulation of the graft and donor lymphocyte infusion, in vivo targeting of Th/Tc17 cell populations could additionally lower the generation of these cells in time. As upregulation of ROR $\gamma$ t is crucial in the generation of Th/Tc17, this could be used as a target to inhibit Th/Tc17 generation. This possibility has been shown in different mouse models, where less GVHD was observed upon adoptive transfer of ROR $\gamma$ t<sup>-/-</sup> compared with WT T cells [24,25]. However, whether this is a direct effect of lacking Th/Tc17 cells, or whether this is due to the observed increase of Tregs should be investigated further. A possible clinical application to apply this strategy in patients is a small molecule inhibitor for ROR $\gamma$ t, as described by Huh et al. [88]. However, possible side effects should be taken into account, as studies of ROR $\gamma$ t<sup>-/-</sup> mice have shown the role of ROR $\gamma$ t in processes other than Th/Tc17 development, resulting in defects in lymphoid development. A putative safer option with similar effects could be induced by isoforms of retinoic acid, which inhibit Th17 formation and enhance Treg generation [17]. Together, targeting ROR $\gamma$ t could be an option to prevent differentiation of Th/Tc17 cells, and thereby prevent the occurrence of GVHD.

### PRR Inhibitors

Prevention of GVHD could also be achieved by preventing the activation of DCs by PAMPs and DAMPs. As these molecules have been implicated in GVHD pathobiology extensively, especially during the initiation phase, they could be

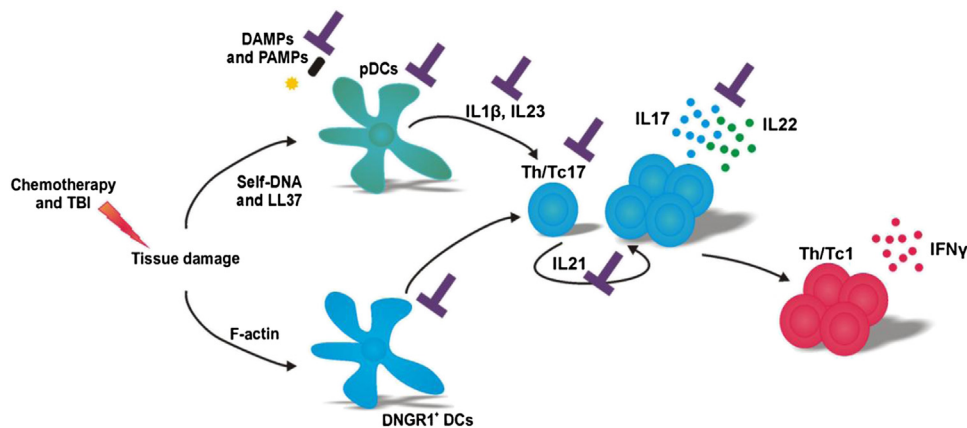
promising cascade targets in the prevention of GVHD [38]. By inhibiting the PRRs, activation of pDCs and DNGR1<sup>+</sup> mDCs could be prevented, thereby suppressing the Th/Tc17 activation.

### Cytokine Pathway Inhibition

An alternative approach to inhibit Th/Tc17 responses would be to influence cytokine cascades involved in the IL17 pathway. Blocking IL1 $\beta$ , which is a major player in GVHD, with canakinumab (IL1 $\beta$  antibody) or its receptor with anakinra (recombinant IL1R antagonist), could interfere with the activation of Th17 cells [89]. However, the blockade of IL1 $\beta$  has shown some contradictory results in GVHD treatment [90]. Blocking IL6 signaling could also contribute to preventing GVHD [91,92]. Nevertheless, blockade of IL6 has not resulted in clear effects on Th/Tc17 responses. A probable mechanism by which blockade of IL6 could contribute to the prevention of GVHD is the increased levels of Tregs observed, with the supplementary suppression of Th/Tc1 and Th/Tc17-mediated tissue damage [92,93]. On the other hand, promising results have been obtained by anti-IL17 (ixekizumab) and anti-IL17 receptor (brodalumab) in the treatment of the IL17-driven skin disease, psoriasis [94,95]. Although blocking IL17 using secukinumab (IL17-blocking antibody) was associated with increased infections in Crohn's disease patients, only local and no systemic fungal complications were observed, showing the feasibility to safely interfere with Th17 responses [96]. Promising results have also been found by depleting the p40 subunit of IL12 and IL23 with ustekinumab (IL12/IL23-blocking antibody), which is approved for the treatment of psoriasis. A recent case report showed the potency of ustekinumab in GVHD treatment, as a complete remission was observed after treatment of steroid-resistant GVHD with this blocking antibody [97]. However, this also coincided with an increase in Tregs, again suggesting that the effect is not solely Th/Tc17-lineage specific. Depletion of IL12 (p70) and IL23 could also be achieved using a small molecule inhibitor, which also for psoriasis, resulted in lower IL17 levels [98]. Moreover, molecules downstream of the cytokine signaling, such as STAT3, could also be considered possible targets. Combined, blockades of signaling pathways could be used for preventing and/or treating GVHD, blocking the Th/Tc17 initiation.

### CONCLUSION AND FUTURE PERSPECTIVES

In recent years, new and exciting insights have been gained in the pathogenesis of GVHD, and increasingly, a role for IL17-producing cells and the Th/Tc17 pathway has been acknowledged. In the future, it might be proven that the specific context of allo-SCT determines whether IL17-producing T cells play an important role or not. As IL17-producing cells are mainly involved in epithelial barrier defenses of gut, lung, and skin, organs where GVHD occurs, it can be anticipated that the role of the Th/Tc17 pathway in GVHD is greatest in the presence of tissue damage (eg, resulting from myeloablative and reduced-intensity conditioning), microbial exposure (eg, infection or gut translocation), and the release of proinflammatory cytokines including IL1 $\beta$ , IL6, and IL23. Consequently, IL17-producing cells emerge and activate early on; that is, during or directly after the conditioning-induced tissue damage and innate immune activation. Thereafter, they set the stage for full-blown GVHD by inducing aggravated tissue inflammation, immune cell recruitment, and priming alloreactive Th1/Tc1 responses. Being an early event in GVHD, the Th/Tc17



**Figure 2.** Th/Tc17 pathway involved in GVHD and its potential targets. TBI indicates total body irradiation; F-actin, filament actin; DNGR1, DC NK lectin group receptor; DC, dendritic cell. DAMP, danger-associated molecular patterns; PAMP, pathogen-associated molecular patterns, Th/Tc, helper T cell/cytotoxic T cell.

pathway seems to be both a logical and suitable target for preventive interventions, and this is most appealing as treatment of manifested acute GVHD remains difficult, with 50% being steroid refractory, and resulting in a high mortality.

Many factors are involved in the differentiation, expansion and activation of IL17-producing T cells in allo-SCT recipients. Different DC subsets, such as pDCs and DNGR1<sup>+</sup> mDCs, have the capability to activate Th/Tc17 cells through the release of cytokines. Pivotal cytokines include IL1 $\beta$ , IL6, IL23, and TGF $\beta$ , which are known to drive the differentiation and expansion of IL17-producing T cells, and these cytokines have been shown to be elevated in patients after allo-SCT. Potent activators of these DC subtypes are motifs that are released on tissue damage (and microbial exposure) during allo-SCT. These motifs aggravate the Th/Tc17 response via activation of various PRRs, thereby initiating and perpetuating GVHD.

At all these different levels, interventions are possible (Figure 2). Attractive approaches are preventing the occurrence of tissue damage by using less toxic conditioning, and inhibiting Th/Tc17 generation by scavenging PAMPs and DAMPs that activate DCs, modulating the PRR pathways, and/or inhibiting key cytokines with monoclonal antibodies, eg, IL1 $\beta$  with anakinra and canakinumab and IL23 with ustekinumab.

In addition, Th/Tc17 can be targeted more specifically by depleting them from the graft, inhibiting their tissue-homing properties, or inhibiting their effector cytokines and related receptors, especially IL17, with ixekizumab and brodalumab, for example.

Although many new preventive strategies seem promising, further investigations are warranted. Still, many questions remain to be answered on the role of different subsets of lymphocytes, including Th/Tc17, in acute GVHD. Nevertheless, the insights we now gain rapidly on the IL17-producing T cells open up new ways that can, and should be, explored for the prevention and treatment of GVHD. This way, we could enhance the outcome of allo-SCT and improve the quality-of-life of hematology patients.

#### ACKNOWLEDGMENTS

*Conflict of interest statement:* There are no conflicts of interest to report.

*Financial disclosure:* The authors have nothing to disclose.

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