



Interleukin-17 cytokine signalling in patients with asthma

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ABSTRACT Asthma remains a global health problem and, therefore, more effective pharmacotherapy is needed. This is particularly true for chronic and severe asthma. In these clinical phenotypes, chronic inflammation involving neutrophils is likely to play a pathogenic role, making it interesting to target cytokine signalling involved in the accumulation of neutrophils. Therefore, it is of interest that the archetype T-helper 17 cell cytokine interleukin (IL)-17A, perhaps also IL-17F, controls neutrophil accumulation, mucus secretion, macrophage mobilisation and smooth muscle reactivity in various experimental airway models. However, much less is known about the involvement of signalling *via* IL-17 cytokines in humans with asthma. Existing evidence suggests that these cytokines are released from several types of immune cells in asthma and, for IL-17A, there is a local increase associated with disease severity, with the mobilisation of neutrophils and smooth muscle cells locally in the airways. Even though the causative role of IL-17 cytokines remains unclear, there is potential for clinical utility in targeting IL-17A specifically in patients with moderate-to-severe asthma and high reversibility. There is a need for new and well-powered clinical investigations of signalling *via* IL-17 cytokines in this clinical phenotype.



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There is potential utility for drugs targeting IL-17 cytokine signalling in a sub-group of patients with asthma <http://ow.ly/wPw5p>

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Introduction

Asthma in humans constitutes a global health problem, affecting more than 300 million people of all ages [1]. Despite substantial progress in terms of optimising existing pharmacotherapeutic principles, therapy remains insufficient and few really novel principles have been developed during recent years. This is particularly true for severe asthma, which accounts for ~10% of all patients with this more or less reversible obstructive airway disease [1]. Thus, there is an unmet global need for improved pharmacotherapy and this means that there is a matching need for more research into pathogenic mechanisms and novel molecular targets.

The fact that not only eosinophils, but also neutrophils (synonymous to neutrophilic granulocytes or polymorphonuclear granulocytes), are mobilised in certain clinical phenotypes of asthma has been recognised [2–6]. This is also true for the fact that clinically important phenotypes among adult patients lack signs of true allergy. It is now also clear that the T helper (Th) 2 cell paradigm is insufficient to explain the whole variety of clinical phenotypes for asthma [2–6]. However, even though these more recent insights have taken the mechanistic understanding of asthma to a new level, they have not yet provided us with new pharmacotherapeutic tools.

Given the lack of sufficient therapy against asthma and due to recent and ongoing trials on drugs targeting the archetype Th17 cytokine interleukin (IL)-17A (synonymous to IL-17), this review will focus on summarising and scrutinising the accumulating body of evidence from (human) patients in whom IL-17A, and additional members of the IL-17 cytokine family (IL-17 cytokines), are actively involved in the pathogenesis of asthma. The preceding work in various animal and cell models, including the experimental evidence from models of allergic asthma, has been extensively reviewed and presented previously [7–14].

Original hypothesis on neutrophil accumulation

The original hypothesis that the heterodimeric cytokine IL-17A contributes to excess accumulation of neutrophils in the airways of patients with asthma was communicated more than a decade ago [7–9]. This hypothesis was initially based merely on plain experimental studies that documented the neutrophil mobilising potential of IL-17A protein in animal airway models *in vivo* plus models of human bronchial epithelial and venous endothelial cells *in vitro* [15–17]. In essence, the results of these early experimental studies indicated that IL-17A recruits and accumulates neutrophils indirectly into the airways by inducing the transcription and release of neutrophil mobilising cytokines from structural airway cells (fig. 1). These early studies and the subsequent studies on “pro-inflammatory effects” of IL-17A demonstrated that the IL-17-induced accumulation of neutrophils is also associated with an increase in proteolytic enzymes, including neutrophil elastase and matrix metalloproteinase-9 [17, 18]. The initial evidence for the actual involvement of IL-17A in patients with asthma was published several years later by MOLET *et al.* [19] and the observations in patients thereafter rapidly obtained support in two subsequent studies [20, 21].

Variety of functional effects

Neutrophils and macrophages

During the past 5 years, several experimental studies have indicated that IL-17A can do more than merely stimulate the accumulation of neutrophils and the release of their proteases in the airways (fig. 1). Thus, there is now experimental evidence from cell and animal models suggesting that IL-17A may stimulate the recruitment of macrophage precursor cells (*i.e.* monocytes), macrophage survival and phagocytosis of particles [22, 23]. Moreover, IL-17A can stimulate neutrophil apoptosis and macrophage phagocytosis of aged neutrophils [23]. In other words, IL-17A may exert “anti-inflammatory effects” that are important for ascertaining the efferocytosis of neutrophils and, thus, the resolution of the innate immune response (fig. 2).

It was recently demonstrated in cell and animal models that IL-17A is important for providing negative feedback on the release of the Th17 regulator IL-23 from macrophages (fig. 2). Thus, it seems feasible that there is a mammalian mechanism to avoid excess signalling through the IL-23/IL-17A axis. Hypothetically, this type of mechanism can serve to maintain a balanced control of neutrophil turn-over for the purpose of preserving an effective antibacterial host defence response without tissue damaging activity [24]. The proposed feedback mechanism is in line with the observation in an animal model that IL-17A also exerts negative feedback on the expansion of IL-17-producing T-cells [25].

Other cells involved in asthma

In addition to its well-documented effects on macrophages and neutrophils, there is evidence from animal models and isolated human bronchi *in vitro* that IL-17A stimulates airway responsiveness [26, 27]. In line with an indirect role for IL-17A in bronchoconstriction, CHANG *et al.* [28] originally showed that IL-17A promotes the proliferation and survival of human airway smooth muscle cells *in vitro*, with a similar relative impact on cells from patients with asthma and control subjects. In further support of an indirect role for

IL-17A in bronchoconstriction and/or hyperreactivity, CHANG *et al.* [29] have also shown that IL-17A increases the migration of human airway smooth muscle cells *in vitro*. Compatible with a more specific role in allergy, SCANLON *et al.* [30] demonstrated that IL-17A stimulates the production of the B-cell chemokine CCL28 in human airway epithelial cells. These authors also demonstrated that the induced CCL28 release increases the chemotaxis of IgE-containing B-cells *in vitro*. In addition, IL-17A stimulates the gene expression for the mucin MUC5B in primary bronchial cells harvested from healthy donors and cultured *in vitro*, which is compatible with a role for IL-17A in airway hypersecretion, a hallmark of acute, severe asthma [31, 32]. Moreover, bronchial epithelial cells from patients with asthma and airway hyperresponsiveness cultured *in vitro* respond to IL-17A with secreted phospholipase A2 group X, indicating a feasible link between IL-17A and the production of lipid mediators in asthma [33]. However, even though the reported effects of IL-17A are intriguing from a pathogenic point of view, there is a general need to evaluate how relevant these effects are in causative terms for asthma patients with or without allergy.

Local involvement

A substantial body of clinical evidence now argues for the presence of IL-17A in the airways of patients with asthma, including data on this heterodimeric protein in bronchoalveolar lavage (BAL), sputum samples, exhaled breath condensate and bronchial tissue.

Bronchoalveolar lavage samples

In their seminal study, MOLET *et al.* [19] reported that the concentrations of soluble IL-17A protein in BAL were moderately increased in patients with mild asthma compared with healthy control subjects. In their study, a substantial fraction of the asthma patients displayed allergen sensitisation and some were ex-smokers but no patient had a history of heavy smoking, was treated with glucocorticoids or had a recent respiratory infection. Thus, it seems unlikely that the presence of IL-17A would be due to a mere activation

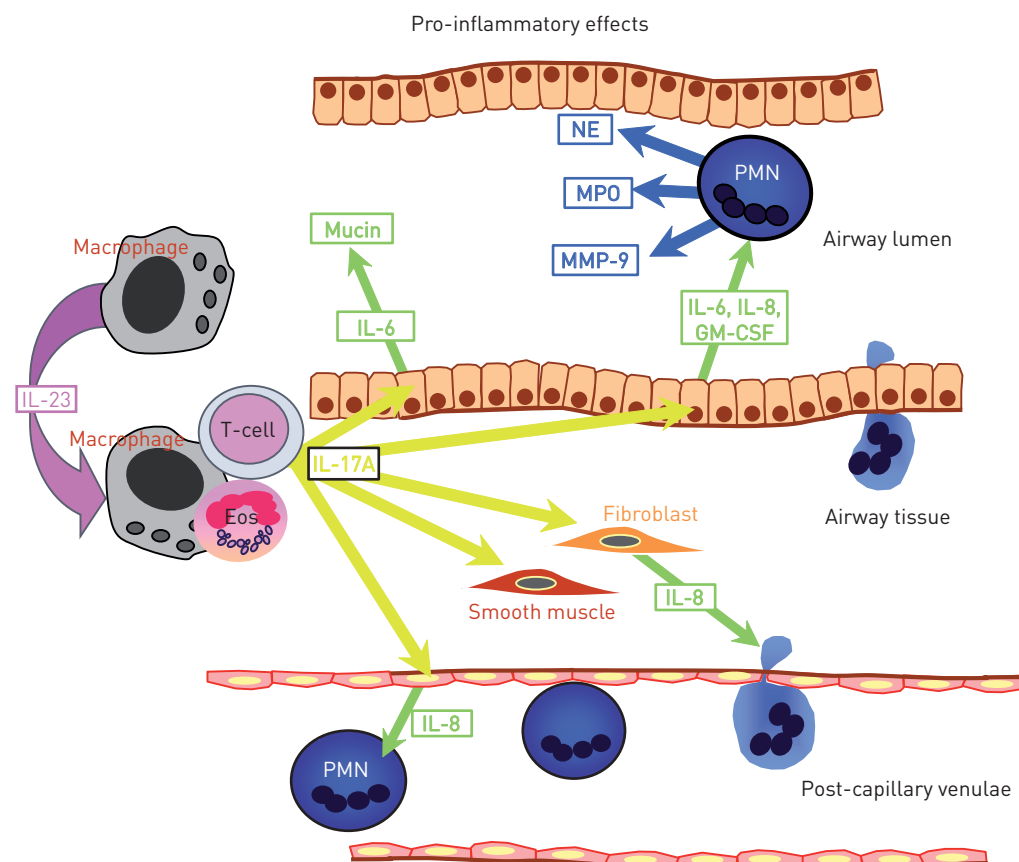


FIGURE 1 Overview of the pro-inflammatory effects caused by interleukin (IL)-17A in experimental studies that are of hypothetical relevance for the pathogenesis of asthma. These include the accumulation of neutrophils and macrophages. Inflammatory cells such as T-cells, macrophages, polymorphonuclear neutrophils (PMN) and eosinophils (Eos) are indicated, as are effector molecules such as matrix metalloproteinase (MMP)-9, myeloperoxidase (MPO), neutrophil elastase (NE) and mucins. GM-CSF: granulocyte-macrophage colony-stimulating factor.

of pulmonary host defence in this important study. MOLET *et al.* [19] also demonstrated a substantially more frequent immunoreactivity signal for IL-17A protein in BAL cells from patients with mild asthma, which was associated with a clear trend towards an increased fraction of neutrophils compared with control subjects.

Sputum samples

In support of their original findings in BAL fluid and cells, MOLET *et al.* [19] demonstrated that the immunoreactivity for IL-17A protein is considerably more abundant in sputum cells from patients with mild asthma when compared with healthy control subjects. Similar to their findings in BAL fluid, MOLET *et al.* [19] also detected a trend towards an increase in sputum neutrophils that was associated with the increase in IL-17A.

In a follow-up study in a limited number of patients with mild-to-moderate asthma but without any documentation of atopy, BARCZYK *et al.* [34] demonstrated a trend towards an increase in luminal IL-17A protein when analysing induced sputum from patients with asthma compared with healthy control subjects. While conducting a pooled analysis of the patients with asthma plus a similarly limited number of patients with chronic bronchitis, BARCZYK *et al.* [34] detected a correlation between an increased concentration of IL-17A protein in induced sputum and the bronchial responsiveness to methacholine (measured as provocation concentration causing a 20% fall in forced expiratory volume in 1 s). However, when comparing the hyperresponsive subgroup of patients with a healthy control group without hyperresponsiveness, the average difference in sputum IL-17A emerged as modest, even though it was statistically significant. It cannot be ruled out that the inconsistent treatment with inhaled glucocorticoids among the patients with asthma contributed to this somewhat vague outcome, given what is known about the sensitivity of the production of IL-17A to glucocorticoids from animal and cell models [35]. Along these lines, in the study by BARCZYK *et al.* [34] the patients with asthma did not have an increase in sputum neutrophils.

SUN *et al.* [36] demonstrated a substantial increase in IL-17A protein in induced sputum from patients with severe asthma compared with healthy control subjects. Notably, this increase in IL-17A was also accompanied by a substantial and corresponding increase in the fraction of neutrophils among sputum cells. SUN *et al.* [36] also examined patients with mild-to-moderate asthma and, when analysing IL-17A

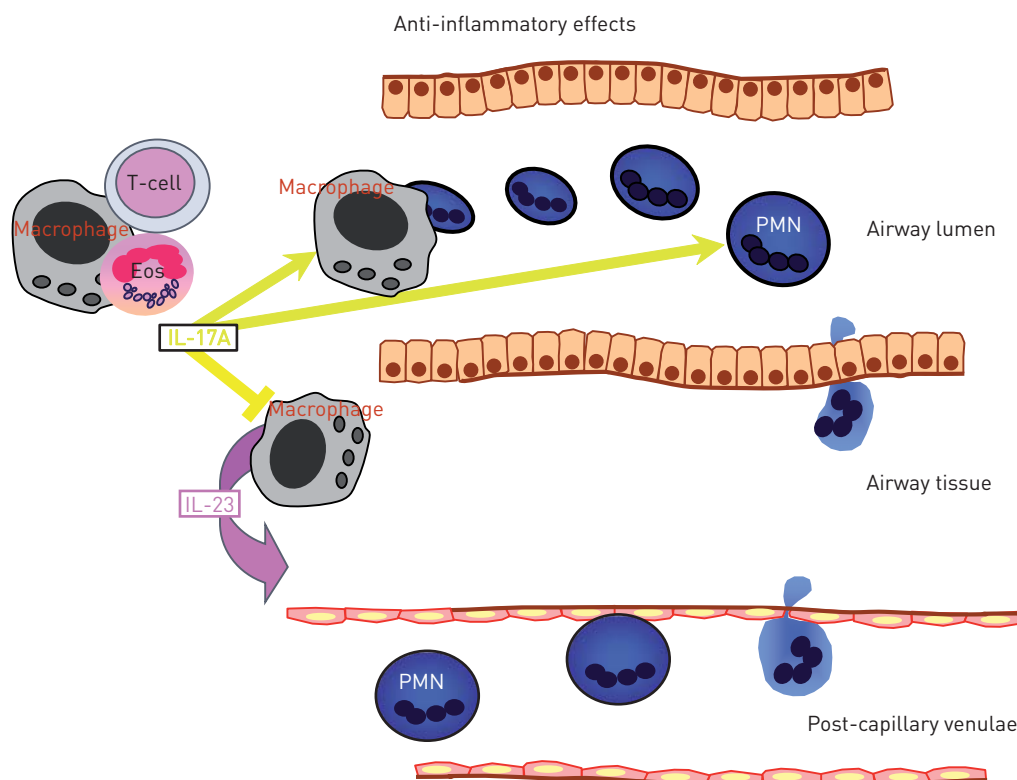


FIGURE 2 Overview of anti-inflammatory effects caused by interleukin (IL)-17A in experimental studies that are of hypothetical relevance for the pathogenesis of asthma. These include the induced apoptosis and phagocytosis of neutrophils. Inflammatory cells such as T-cells, macrophages and polymorphonuclear neutrophils (PMN) are indicated. Eos: eosinophils.

protein in all stages of disease, they found that the increase in sputum IL-17A clearly paralleled the severity of disease. The same was true for the protein concentrations of both the neutrophil-recruiting chemokine IL-8 and the activity marker myeloperoxidase, thus illustrating the association of IL-17A with neutrophil mobilisation in asthma. Importantly, none of the patients with asthma were treated with glucocorticoids at the time of the referred investigations, were recently infected or were a smoker [36]. Moreover, when patients with all three degrees of disease severity were subsequently treated with a moderate daily dose of an inhaled glucocorticoid, this treatment reduced the concentration of IL-17A protein. This suggests that the net release of IL-17A protein in the airways of patients with asthma is actually sensitive to glucocorticoids under these conditions, contrasting with previously published data indicating a very low sensitivity to glucocorticoids in animal models [35, 37]. However, and this may be important, the utilised glucocorticoid treatment did not markedly affect the relative number of neutrophils in induced sputum from the patients in this study.

A true correlation between IL-17A and neutrophils was demonstrated by BULLENS *et al.* [38] when assessing mRNA for IL-17A in sputum cells from patients with mild and moderate-to-severe asthma. However, the increase in this mRNA was modest compared with healthy control subjects but the results did not suggest that the transcription of IL-17A is sensitive to treatment with inhaled glucocorticoids. In addition, there was a trend only towards a moderately higher level of mRNA for IL-17A in verified allergic compared with non-allergic asthma.

An unexpected link between IL-17A and a particular clinical phenotype of asthma was demonstrated by KAMINSKA *et al.* [39]. The authors assessed airway remodelling outcomes *versus* cytokine content in induced sputum and compared patients with chronic *versus* reversible phenotypes of severe asthma [39]. They found a higher content of luminal IL-17A protein when analysing sputum samples from patients with the reversible rather than with the chronic phenotype, but there was actually a luminal neutrophil accumulation of a similar magnitude in both phenotypes. Thus, it can be speculated that in addition to IL-17A, other mechanisms are involved in determining the luminal accumulation of neutrophils in patients with asthma, even though additional factors may have set the outcome of this particular study. First, there was inconsistent use of inhaled and oral glucocorticoids for the two study groups. Secondly, it is known that sputum sampling tends to yield higher neutrophil content in all samples and reflect more proximal conditions than BAL samples. Hypothetically, these factors may have shaded a more clear-cut difference in neutrophil accumulation in the lower airways for the patients with the reversible compared to the chronic phenotype.

DOE *et al.* [40] compared the concentration of IL-17A protein in the sputum of patients with asthma with that observed in patients with chronic obstructive pulmonary disease (COPD). These investigators detected higher concentrations of IL-17A protein in patients with COPD and no ongoing exacerbation compared with patients with asthma, even though the highest individual concentrations of IL-17A were detected in patients with asthma. Thus, this study indicated that there is a higher luminal concentration of IL-17A in COPD than asthma. However, the interpretation is complicated by the fact that current smokers were included among the patients with asthma in such a number that the average tobacco load exceeded that of the healthy control subjects. Moreover, former and never-smokers, but no current smokers, were included among the patients with COPD and the average tobacco load in the COPD group exceeded that in the group of smokers without COPD [40]. Finally, there was no information on the types of inhalation therapy being used by either of the patient groups and, thus, the confounding influence of glucocorticoids cannot be excluded. The key results from the study by DOE *et al.* [40] suggesting that there is more IL-17A in COPD than in asthma can, therefore, be questioned. These results are in need of confirmation by new studies with improved control of the confounding influence of tobacco load among the study groups.

Finally, given the question about IL-17 signalling and atopy, it is interesting to note the results from a recent study by MANISE *et al.* [41]. These investigators showed that the concentration of IL-17A protein in induced sputum is higher in patients with asthma plus high serum levels of IgE compared with low levels of this type [41]. As is expected, the former group also displayed a higher frequency of atopy, so a link to atopy cannot be excluded. However, the relevance of these findings in sputum remains to be verified in terms of their relevance for conditions in the peripheral airways.

Exhaled breath condensate

MATSUNAGA *et al.* [42] published a study on the concentrations of IL-17A protein in exhaled breath condensate from patients with asthma and healthy control subjects. The subjects were nonsmokers and who were not receiving any glucocorticoid treatment. In accordance with most of the studies on BAL and sputum samples, the results of this study suggest that IL-17A protein is moderately increased in patients with mild asthma, compared with healthy control subjects. It is of particular mechanistic interest that MATSUNAGA *et al.* [42] obtained indications that the increase in IL-17A among these patients with asthma is

associated with a corresponding increase in IL-8, tumour necrosis factor (TNF)- α and transforming growth factor (TGF)- β ; cytokines that are associated with the mobilisation of neutrophils and Th17 cells [7–13]. It is also of mechanistic interest that MATSUNAGA *et al.* [42] detected a trend towards a correlation between the concentration of “exhaled” IL-17A and airway obstruction in patients with asthma. Unfortunately, the utilised technique did not allow the investigators to relate exhaled IL-17A to the accumulation of luminal neutrophils [42].

Bronchial tissue

CHAKIR *et al.* [21] published early and important evidence for IL-17A being present in airway tissue from patients with asthma. By assessing immunoreactivity for IL-17A protein in bronchial tissue, the authors showed that patients with moderate-to-severe asthma have markedly more cells containing IL-17A protein in the sub-epithelial layer compared with healthy control subjects. Notably, there were no smokers among the patients or control subjects and none of the patients were treated with glucocorticoids [21]. The authors also found an increased IL-17A signal when analysing the intraepithelial bronchial layer [21]. The increase in IL-17A was also paralleled by a corresponding increase in TGF- β and type I collagen, showing that the basic prerequisite for IL-17A-producing cells being involved in remodelling were present. The major part of the immunoreactivity signal for IL-17A protein was reduced by systemic treatment with a glucocorticoid, in parallel with what has previously been shown in sputum samples [36]. AL-RAMLI *et al.* [43] subsequently demonstrated an increase in cells immunoreactive for IL-17A protein in the sub-epithelial layer of bronchial tissue from patients with various degrees of asthma severity. In their study, AL-RAMLI *et al.* [43] clearly demonstrated that the magnitude of the increase in IL-17A parallels disease severity and that this increase is substantial in patients with severe asthma. For IL-17A mRNA, the pattern was similar. Unfortunately, the referred publication included very little information on the current medication of the patients.

When DOE *et al.* [40] recently examined submucosal cells immunoreactive for IL-17A protein they detected a trend that IL-17A is most abundant in tissue from patients with mild-to-moderate asthma when compared with tissue from healthy control subjects and patients with truly severe asthma. However, the power of the utilised material was insufficient to prove this point in a statistically valid sense and it is unclear whether there were differences in medication or smoking habits between study groups.

Systemic involvement

WONG *et al.* [20] were actually earlier than MOLET *et al.* [19] in specifically addressing the presence of IL-17A protein in a study in patients with asthma. However, WONG *et al.* [20] examined blood cells and plasma and, thus, IL-17A at the systemic level. They found a trend towards a moderate increase in soluble IL-17A in the plasma of patients with allergic asthma when compared with healthy control subjects without allergy [20]. Unfortunately, the study material was relatively small and thus the statistical power was insufficient to prove significance. The preliminary finding by WONG *et al.* [20] was subsequently confirmed by the results of the study by MOLET *et al.* [19]; including a statistically significant increase in blood eosinophils immunoreactive for IL-17A protein in patients with mild asthma compared with healthy control subjects.

Interestingly, AGACHE *et al.* [44] recently forwarded evidence for high IL-17A protein in serum constituting an independent risk factor for severe asthma. These investigators found that patients with severe asthma, some treated with a high dose of an inhaled glucocorticoid, have substantially more IL-17A protein in serum compared with patients with mild-to-moderate asthma. In this study, however, patients with or without atopy were not consistently separated in the analysis. The most interesting finding by AGACHE *et al.* [44] was that a serum IL-17A concentration exceeding a threshold ($20 \text{ pg}\cdot\text{mL}^{-1}$) is associated with a more than three-fold increased risk for severe disease. Here, there was evidence that serum IL-17A protein is negatively correlated with the fraction of blood neutrophils, as well as with airflow in the small airways according to spirometry. Based upon this, it can be speculated that IL-17A produced and released in the airways does recruit neutrophils from the blood, thereby contributing to small airway obstruction. Thus, the study supports the idea that IL-17A is critically involved in the pathogenesis of severe asthma.

When analysing plasma, ZHAO *et al.* [45] obtained evidence that IL-17A protein at the systemic level is markedly (more than five-fold) increased in patients with allergic asthma in general and even further increased in allergic patients with severe asthma in comparison to healthy control subjects. A corresponding and clear increase in the release of IL-17A protein was detected in conditioned media from peripheral blood mononuclear cells cultured and activated *in vitro*. From a mechanistic point-of-view, it is interesting that ZHAO *et al.* [45] were able to demonstrate a positive correlation between IL-17A and IL-23 protein in the group of patients with allergic asthma, since this is compatible with IL-23 constituting a trigger of IL-17A production in circulating Th17 cells.

In a recently published study on blood Th cells from healthy control subjects and patients with moderate-to-severe asthma who had been without treatment with glucocorticoids for 4 weeks, NANZER *et al.* [46] found that the intracellular immunoreactivity for IL-17A and IL-22 was enhanced in patients with “steroid-refractory” but not “steroid-sensitive” clinical disease. The blood Th cells co-expressing IL-17A and IL-22 were enhanced in all patients with asthma, as was the corresponding release of extracellular protein *in vitro*. For patients with steroid-refractory clinical disease, the release of IL-17A protein was higher than in the patients with steroid-sensitive disease. Moreover, treatment with a glucocorticoid enhanced the extracellular release of IL-17A protein in blood Th cells from healthy control subjects but not in the corresponding cells from patients with asthma, albeit the latter cells displayed more release of IL-17 in absolute terms. Clearly, this provides evidence that cytokine production in Th17 cells may be less sensitive to the inhibitory effect of glucocorticoids and, possibly, the Th17 phenotype may even be driven by this class of anti-inflammatory drugs.

Multiple cellular sources

Sources in the lungs

MOLET *et al.* [19] published the initial evidence for IL-17A in eosinophils harvested from BAL and sputum samples. This evidence was obtained using three different techniques: immunocytochemistry, Western blot and *in situ* hybridisation [19]. Until recently, there was no functional evaluation of this potentially important finding. However, there is now a study from KOBAYASHI *et al.* [47] suggesting that isolated human blood eosinophils harvested from healthy volunteers can release IL-17A protein, albeit in response to crystals of monosodium, a stimulus of unclear relevance for human asthma.

Most evidence from human cell models, as well as from animal *in vivo* and *in vitro* models, suggests that CD3-positive mononuclear cells constitute important IL-17A-producing cells, thus pointing out T-cells [7–14, 48–50]. However, the corresponding evidence from human lungs is much more limited. In an early study, IVANOV *et al.* [49] demonstrated immunoreactivity for IL-17A protein in BAL cells with a lymphocyte-like morphology after exposure to organic dust in healthy human volunteers. IVANOV *et al.* [49] also demonstrated a substantial increase in detectable mRNA for IL-17A among BAL cells after exposure to organic dust in the same study on humans.

More recently, GLADER *et al.* [50] demonstrated the presence and increase of CD3- and CD4-positive but CD8-negative cells containing IL-17A in BAL samples harvested after exposure to endotoxin in a bronchial segment of healthy volunteers. These IL-17A-containing Th cells were detected in a mixed population of BAL cells; one that also contained mRNA for RORC variant 2, the archetype human Th17 transcription factor. GLADER *et al.* [50] also showed that the level of mRNA for RORCvariant2 correlates with the level of mRNA for IL-17A after local endotoxin exposure. The findings clearly support the idea that Th17 cells constitute an important source for IL-17A in human airways, even if this type of evidence for a Th17-like phenotype of cells has not yet been generated in patients with asthma.

PURWAR *et al.* [51] published evidence for Th cells producing IL-17A protein in human lungs; more precisely, CD4-positive cells containing IL-17A protein in lung tissue from patients undergoing surgery for lung cancer. The study by PURWAR *et al.* [51] also demonstrates that whereas the IL-17A-containing CD4-positive cells constitute around 1% of all CD4-positive cells, IL-17A-containing CD8-positive cells constitute 0.8% of all CD8-positive cells in lung tissue from patients undergoing surgery for lung cancer. Thus, these findings are compatible with Tc17, as well as Th17 cells constituting sources of IL-17A in human lungs, but it is not known whether asthma is linked to both or either of these T-cell subsets.

Given the limited data on the phenotype of T-cells producing IL-17A in the airways of patients with asthma, it is important to pursue more elaborate studies in this field. The study by BULLENS *et al.* [38] demonstrated a positive and relatively strong correlation between the levels of mRNA for IL-17A and CD3- γ in sputum harvested from atopic and non-atopic patients with moderate-to-severe asthma. This correlation is compatible with the observation of AL-RAMLÍ *et al.* [43] who described immunoreactivity for IL-17A protein in mononuclear inflammatory cells, mainly residing in the bronchial submucosa of patients with asthma. It remains to be evaluated whether Th17 or Tc17 cells in the airways of patients with asthma share characteristics with the corresponding blood cells [12, 13, 48, 51–57]. Moreover, the documentation of expression for IL-17A in invariant natural killer T-cells and in mucosal-associated invariant T-cells from human subjects is intriguing but further study is needed given that there are no published studies addressing these cell types in the airways of patients with asthma [11, 13, 42, 44, 48, 50–66].

Finally, immunoreactivity for IL-17A protein has been detected in macrophages from the airways of patients with allergic asthma and this motivates more in-depth investigations [58]. This is also true for the recent detection of immunoreactivity for IL-17A in neutrophils from the airways of patients with cystic fibrosis [59].

Sources in the blood

There is support from several studies that Th17 and other immune cells in the blood from patients with allergic asthma can produce IL-17A protein under certain conditions.

HASHIMOTO *et al.* [60] demonstrated that isolated mononuclear cells harvested from the blood of patients with allergic asthma produce IL-17A protein in response to stimulation with anti-CD3 plus anti-CD28 antibodies and house dust mite extract. Examining peripheral blood mononuclear cells from patients with allergic asthma, ZHAO *et al.* [64] subsequently showed that there is a subset of CD3-positive and CD8-negative cells with immunoreactivity for IL-17A, arguing that it is the Th17 and not the Tc17 population that is involved in asthma. According to the referred study, the percentage of these Th17 cells is moderately increased in patients with allergic asthma, compared with healthy control subjects. Of note, the percentage of these presumed Th17 cells seems to increase with disease severity.

COSMI *et al.* [61] generated T-cell clones from the CCR6- and CD161-positive fraction of CD4- and CD3-positive peripheral blood cells and examined the cytokine content of these cells. After stimulation, these cells produced IL-17A and IL-4 as well as other archetype Th17- and Th2-cytokines. WANG *et al.* [62] have published similar findings. However, in an unorthodox manner, WANG *et al.* [62] claimed that the subset of CD4-positive cells containing IL-17A protein, as well as the Th2 cytokines IL-4, -5 and -13, is a CD45-, CCR6- and CRTH2-positive subset. According to WANG *et al.* [62], this subset contains both the archetype transcription factors for Th17 and Th2, RORCvar2 and GATA-3. This actually indicates the existence of a Th17/Th2 subset of memory Th cells, whether a committed lineage or not. According to the study by COSMI *et al.* [61], this subset of Th17/Th2 cells causes the secretion of IgE from autologous human B-lymphocytes (B-cells) *in vitro*, a mechanism indirectly linking IL-17A to true allergy. In their study, ZHAO *et al.* [64] demonstrated that there is an increased frequency of CD4-positive cells containing IL-17A protein in the blood of patients with allergic asthma in comparison to healthy control subjects. This was confirmed in the recent study by RAMIREZ-VELAZQUEZ *et al.* [63].

To date, there is very limited documentation of blood $\gamma\delta$ T-cells as sources of IL-17A in patients with asthma. However, a study by ZHAO *et al.* [64] reports evidence that the relative expression of IL-17A is increased among CD4-, CD8- and $\gamma\delta$ T-cells in patients with allergic asthma compared with healthy control subjects. The study did not contain any specific information on the relationship between extracellular concentrations and intracellular expression of IL-17A.

The fact that blood eosinophils may be capable of producing IL-17A protein has been discussed previously [19, 20, 45]. In addition to this evidence of granulocytes constituting a source of IL-17A, there is now evidence that a subset of blood neutrophils expressing the adhesion molecule CD177 contain IL-17A protein as well [63]. Even more interesting, this CD177-positive subset of neutrophils is markedly increased in patients with allergic asthma when compared with healthy subjects, and it is more abundant in moderate asthma than mild asthma. It is intriguing that the most pronounced signal for IL-17A in CD177-positive neutrophils among patients with allergic asthma was detected among those allergic to fungal allergens, again emphasising the link between IL-17A and pulmonary host defence [50].

Receptor signalling

It has now been established in a range of cell and animal models that IL-17A acts by stimulating a receptor complex constituted by an IL-17 receptor A (IL-17RA) and IL-17 receptor C (IL-17RC) receptor sub-unit, signalling *via* the adaptor protein nuclear factor (NF)- κ activator (Act1) downstream to more generic intracellular signalling compounds [13, 28, 56, 65–67]. These signalling compounds may include TNF receptor-associated factor-2, -3 and -6, TGF-activated kinase-1 and mitogen-activated protein kinases such as c-jun N-terminal kinase, extracellular-regulated kinase and p38 [13, 28, 56, 67–69]. This signalling cascade leads to the activation of NF- κ B-activating kinase and the transcription factors NF- κ B, CCAAT/enhancing-binding protein- β and - γ [13, 28, 56, 67–69]. In addition, there is an Act1-independent pathway signalling *via* Janus kinase-1 and phosphatidylinositol 3-kinase including inactivation of glycogen synthase kinase-3 β .

Remarkably little is known about human pathology *versus* aberrations in signalling *via* IL-17 cytokines and this is particularly true for patients with asthma. PARK *et al.* [70] recently demonstrated that there is an association of single nucleotide polymorphisms on the IL-17RA gene and aspirine hypersensitivity in patients with asthma, thereby suggesting a pathogenic involvement of IL-17RA. Moreover, it was recently demonstrated that Act1 is a client protein of ATPase and the chaperone protein heat shock protein (Hsp)90 in an animal *in vivo* model of psoriasis [71]. When the Act1-Hsp90 interaction is disrupted, Th17 signalling through IL-17A becomes attenuated and the corresponding signalling through IL-22 becomes hyperresponsive. Given the high mammalian conservation of IL-17A-related mechanisms, it can be

speculated that a malfunction in the critical interaction between Act1 and Hsp90 contributes to altered IL-17 signalling in human pathology, such as in asthma [71].

Involvement of additional IL-17 cytokines

There are two known members of the IL-17A cytokine family that are very similar to IL-17A from a structural and functional point of view, which have been detected in patients with asthma, namely IL-17F and IL-25 (formerly known as IL-17E) [10, 12, 13]. Of these two cytokines, it is IL-17F that can currently be linked to neutrophil mobilisation whereas IL-25 mainly seems to be linked to eosinophil mobilisation.

Interleukin-17F

The initial evidence that IL-17F is relevant for asthma in humans was published by KAWAGUCHI *et al.* [72]. They demonstrated that IL-17F protein stimulates the production and release of neutrophil-mobilising cytokines in human bronchial epithelial cells. They also demonstrated that local allergen challenge increases the signal for IL-17F mRNA among BAL cells in patients with allergic asthma [72].

Whereas IL-17F seems to act *via* the same IL-17RA/receptor C complex as IL-17A, it is conceptually interesting that IL-17F may be produced and released by a wider range of cells than is the case for the archetype Th17 cytokine IL-17A [10, 12, 13, 72]. Thus, the production of IL-17F appears to take place in structural cells as well as in immune cells, more specifically in epithelial cells, basophils, mast cells, monocytes and T-cells.

More recently, AL-RAMLÍ *et al.* [43] reported an increase in subepithelial immunoreactivity for IL-17F protein in patients with verified asthma but with no information on atopy or current medication. The referred increase in IL-17F protein followed disease severity, as did the level of the corresponding mRNA. Interestingly, the study by AL-RAMLÍ *et al.* [43] indicated IL-17F protein in the epithelial layer and even within epithelial cells in the very same patients that expressed IL-17A in the submucosa. Thus, IL-17F and IL-17A may be expressed simultaneously.

In a recently published study, DOE *et al.* [40] reported an increased signal for IL-17F protein in the submucosa of bronchial tissue from patients with asthma compared to healthy control subjects. Even though their material failed to prove a statistically significant difference in this sense, the data from DOE *et al.* [40] suggested a trend that this expression of IL-17F protein increases with disease severity.

Interleukin-25

Whereas an early experimental study on a mouse model indicated that IL-25 may trigger steroid-resistant Th2 signalling *via* IL-4 and IL-13, there is still limited information on the relevance of IL-25 for the pathogenesis of patients with asthma [73].

When measuring protein concentrations in serum from patients with severe asthma compared with healthy control subjects, NADI *et al.* [74] found no clear difference in extracellular IL-25 protein. However, a trend towards somewhat higher IL-25 concentrations in patients with asthma was detected in whole blood. Unfortunately, no information on atopy was provided in this study. However, SEYS *et al.* [75] obtained evidence that high levels of mRNA for IL-25 in sputum cells, just like high mRNA levels for IL-17A, are increased in patients with stable asthma compared with healthy control subjects. However, both groups included atopic subjects so the role of allergy is difficult to evaluate for this study but, of clinical interest, a pattern of particularly high levels of mRNA for IL-25, IL-5 and IL-17A was associated with poorly controlled asthma. Moreover, TANG *et al.* [76] have recently shown that the extracellular concentration of IL-25 in plasma is increased in patients with allergic asthma compared with non-atopic healthy subjects but not when compared with atopic non-asthmatic subjects. In their study, the concentration of IL-25 in plasma displayed a modest, negative correlation with lung function assessed as forced expiratory volume in 1 s [76].

Tentatively, IL-25, with its proven link to the mobilisation of eosinophils rather than neutrophils, and its unique signalling *via* the IL-17receptor A/receptor B complex, remains of interest for further studies in patients with asthma [77, 78]. Based upon the few and relatively small studies conducted to date, it is possible, but not proven, that IL-25 is involved in the pathogenesis of certain phenotypes of asthma. If this turns out to be the case, the next goal should be to evaluate whether phenotypes with allergy are of particular interest for intervention with IL-25 signalling.

How should we target IL-17 cytokines in asthma?

The general strategy for pharmacotherapeutic development may appear an easy one. First, asthma is a disease where Th cells are believed to play a key role and there is evidence for an excess mobilisation of neutrophils in certain clinical phenotypes of asthma [2–9]. Secondly, there is evidence for an increase in the archetype Th17 cytokine IL-17A and its sibling IL-17F locally in the airways of patients with asthma

[19, 70]. Finally, at least for IL-17A, there is also evidence for a corresponding increase in blood cells [19]. Given that IL-17A and IL-17F contribute to the mobilisation of neutrophils, these two cytokines, as well as the receptor complex they share, emerge as promising targets. However, it is not trivial to decide whether the cytokine signalling should be targeted at the protein or at the receptor level and this is for several reasons.

There is now a growing understanding that, just like neutrophils *per se*, IL-17A and IL-17F are critically involved in mammalian host defence against bacteria, fungi and, possibly, viruses [7–9, 11–13, 50]. Naturally, this latter fact constitutes a potential complication when it comes to targeting IL-17A and IL-17F, regardless of whether the protein or the shared receptor complex is targeted. Further indication for potential complications is brought forward by the novel experimental evidence from cell and animal models that IL-17A is important for the resolution of the innate immune response and for providing negative feedback to avoid excess signalling through the IL-23-IL-17A axis (fig. 2) [23–26]. This may also constitute a therapeutic problem regardless of whether the cytokine signalling is targeted at the protein or the receptor level.

Even though we still lack corresponding data on the more exact cellular phenotype from patients with asthma, we know from the study by GLADER *et al.* [50] that a “Th17-like” population of immune cells is involved in the pulmonary host defence of healthy human subjects. This study clearly provides a rationale for characterising the immune cells producing IL-17A and IL-17F in the airways of patients with asthma more in detail. Moreover, the recently demonstrated involvement of “Th17-like” cells in community-acquired pneumonia, both at the local and the systemic level, underlines the need for caution when targeting cytokine signalling *via* IL-17A and IL-17F in patients at risk of infection [79]. It seems reasonable to assume that local inhibition of this cytokine signalling is a safer approach than systemic inhibition, but this remains to be proven.

Clinical trials on the inhibition of IL-17 signalling in asthma

To date, there is very little information on the clinical utility of targeting IL-17 cytokines in patients with asthma. BUSSE *et al.* [80] recently published a report of a clinical trial of an IL-17RA monoclonal antibody in patients with various types of asthma. Interestingly, a beneficial clinical effect was claimed for patients with inadequately controlled asthma who displayed high reversibility in response to a bronchodilator. Of note, this is in line with the finding of a higher level of luminal IL-17A protein among patients with more reversible airflow obstruction in the recent study by KAMINSKA *et al.* [39]. The clinically beneficial effect was defined as an improvement in the scoring of the Asthma Control Questionnaire, whereas the effect did not reach statistical significance in terms of lung function (defined as forced expiratory volume in 1 s), for which a trend towards improvement was observed. Given that the main group of interest was only one out of nine sub-groups investigated, new, better-powered studies of the particular sub-group (*i.e.* phenotype) are warranted. Such studies may also clarify whether the postulated beneficial effect of the anti-IL-17RA antibody in highly reversible asthma really relates to an impact on neutrophil mobilisation or to other mechanisms. Another important question to address is whether the dosing of the anti-IL-17RA antibody was optimal for the patients that were studied.

In further support of conducting new studies, an anticipated increase in local infections was observed in the clinical trial by BUSSE *et al.* [80], and this side-effect is in need of further evaluation prior to developing this new pharmacotherapeutic principle.

Conclusions and questions for the future

The evidence from an increasing number of studies, albeit conducted on relatively few patients, suggests that cytokine signalling *via* IL-17A and IL-17F is involved in the pathogenesis of asthma. However, due to the fact that most studies have been conducted on mixed clinical phenotypes of asthma, the significance of this involvement remains uncertain. In particular, due to the lack of critical mechanistic evidence from patients it is not yet possible to make a firm conclusion with reference to the causative involvement of cytokine signalling *via* IL-17A and IL-17F in the mobilisation of neutrophils in asthma. This being said, it is very promising that a recent interventional [39] and a recent descriptive study [80] identified patients with moderate-to-severe asthma and high reversibility as a clinical phenotype of particular interest. Even though these two important studies leave several important questions unanswered, this particular finding warrants further study.

It is also of interest to note that the growing body of evidence for not only pro- but also anti-inflammatory effects of IL-17A in experimental studies on neutrophil mobilisation, indicates a dual role of this archetype Th17 cytokine that questions the point of inhibiting it. This leaves us with the question of whether cytokine signalling *via* IL-17A and IL-17F controls the turnover of neutrophils in certain patients with asthma by increasing accumulation, as well as efferocytosis, apoptosis and the phagocytosis of neutrophils.

Additional critical questions to address in future studies of well-characterised patients with asthma include whether cytokine signalling *via* IL-17A and IL-17F contribute to the mobilisation of macrophages or the

development of airway hyperresponsiveness in patients with particular phenotypes of asthma. This may be particularly interesting to address in patients with severe asthma [81]. The putative involvement and role of IL-25 is also in need of further study. A very interesting question for future studies is whether allergic and non-allergic asthma differ in terms of the specific involvement of IL-17A, IL-17F and IL-25 [13, 22, 26, 82, 83]. Another, more practical, question for future studies is whether the signalling pathways mediated by IL-17A and IL-17F are truly sensitive to drugs that are already in clinical practice, including glucocorticoids [36]. This question has now become even more interesting due to recent findings on the involvement of IL-17A in induced glucocorticoid insensitivity in human cells cultured *in vitro* and in transformed bronchial epithelial cells, as well as in primary monocytes [84, 85]. Furthermore, given what is now known about the downstream intracellular signalling caused by IL-17A and IL-17F via the IL-17A/C receptor complex, it is imperative to establish whether this aspect of signalling is altered in patients with certain phenotypes of asthma [13, 67–69]. The same is true for IL-25 and its signalling via the IL-17RA/RB complex [77, 78].

Finally, given the now established role of IL-17A and IL-17F in pulmonary host defence, it is important to design future trials on the concept of inhibiting cytokine signalling via IL-17 cytokines so that immunosuppressive side-effects can be carefully monitored.

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