Immunopathology of chronic rhinosinusitis

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Abstract

Chronic rhinosinusitis (CRS) is a heterogeneous disease characterized by local inflammation of the upper airways and sinuses which persists for at least 12 weeks. CRS can be divided into two phenotypes dependent on the presence of nasal polyps (NPs); CRS with NPs (CRSwNP) and CRS without NPs (CRSsNP). Immunological patterns in the two diseases are known to be different. Inflammation in CRSsNP is rarely investigated and limited studies show that CRSsNP is characterized by type 1 inflammation. Inflammation in CRSwNP is well investigated and CRSwNP in Western countries shows type 2 inflammation and eosinophilia in NPs. In contrast, mixed inflammatory patterns are found in CRSwNP in Asia and the ratio of eosinophilic NPs and non-eosinophilic NPs is almost 50:50 in these countries. Inflammation in eosinophilic NPs is mainly controlled by type 2 cytokines, IL-5 and IL-13, which can be produced from several immune cells including Th2 cells, mast cells and group 2 innate lymphoid cells (ILC2s) that are all elevated in eosinophilic NPs. IL-5 strongly induces eosinophilia. IL-13 activates macrophages, B cells and epithelial cells to induce recruitment of eosinophils and Th2 cells, IgE mediated reactions and remodeling. Epithelial derived cytokines, TSLP, IL-33 and IL-1 can directly and indirectly control type 2 cytokine production from these cells in eosinophilic NPs. Recent clinical trials showed the beneficial effect on eosinophilic NPs and/or asthma by monoclonal antibodies against IL-5, IL-4Rα, IgE and TSLP suggesting that they can be therapeutic targets for eosinophilic CRSwNP.

Introduction

Chronic rhinosinusitis (CRS) is a heterogeneous disease characterized by local inflammation of the upper airways and sinuses that persists for at least 12 weeks. It is a significant cause of morbidity in adults across the world, affecting over 10 million Americans.1–4 CRS is frequently divided into two groups based on the presence or absence of nasal polyps (NPs); CRS with NPs (CRSwNP) and CRS without NPs (CRSsNP). Inflammatory patterns in the two diseases are known to be different. Initial studies indicated that CRSsNP is characterized by type 1 inflammation and CRSwNP by type 2 inflammation.4–7 However, recent evidence suggests that this theory may not fit in Asia where mixed inflammatory patterns can frequently be found in Asian patients in contrast to those in Western countries. This review article will focus on the pathogenic role of immune cells in CRS and will clarify the current knowledge of inflammatory patterns in CRSsNP and CRSwNP in the Western and Asian populations.

CRS is also known to be characterized by tissue remodeling. However, the pattern of remodeling between CRSsNP and CRSwNP is different.5,8–11 Importantly, this variation is controlled by distinct inflammatory patterns, known to be different in the two diseases. Since these patterns are controlled by immune signaling, we will introduce remodeling as an example of differences between CRSsNP and CRSwNP that is controlled by immune signaling and then introduce the inflammatory patterns in CRS in the following sections.

Keywords:
Chronic rhinosinusitis
Eosinophilia
Nasal polyps
TSLP
Type 2 inflammation

Abbreviations:
BAFF, B cell-activation factor of the TNF family; CRS, chronic rhinosinusitis; CRSsNP, CRS without nasal polyps; CRSwNP, CRS with nasal polyps; CRTH2, chemoattractant receptor-homologous molecule expressed on Th2 cells; CysLT, cysteinyl leukotriene; DCs, dendritic cells; mDCs, myeloid DCs; DP2, P2X2 receptor 2; ILC2s, group 2 innate lymphoid cells; MC-T, mast cell-tryptase/chymase; NPs, nasal polyps; PG, prostaglandin; t-PA, tissue plasminogen activator; TSLP, thymic stromal lymphopoietin

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Remodeling

Tissue remodeling in CRS includes changes to the tissue structure and the deposition of extracellular matrix (ECM). Several factors that have been implicated in remodeling including matrix metalloproteinase (MMP) and TGF-β are well invested molecules in CRS. MMPs are a family of zinc-dependent proteolytic enzymes that degrade various components of ECM and mediate remodeling in both physiological and pathological processes. Several groups have investigated a subset of MMPs in CRS and shown that MMP-7 and MMP-9 were elevated in both CRSsNP and CRSwNP. This suggests that MMP-7 and MMP-9 might be remodeling factors common to both forms of CRS. TGF-β mediates anti-inflammatory effects by acting on many types of immune cells. TGF-β also affects fibrosis by increasing deposition of ECM such as collagen, fibronectin and proteoglycan. TGF-β has three isoforms, TGF-β1, TGF-β2 and TGF-β3. Van Brauene et al. reported in the Belgium population that TGF-β1, TGF-β2, and collagen were elevated in CRSsNP. The same group also investigated the presence of TGF-β in the Chinese population. Li et al. reported that TGF-β1 and collagen were elevated in CRSSNP. In contrast, Shi et al. found in the Chinese population that TGF-β1 was reduced but TGF-β2 and collagen were elevated in CRSsNP. Additionally, Sejima et al. reported that TGF-β3 was elevated in CRSsNP in the Japanese population although they did not state the isoform. Although there is some discrepancy, these studies suggest that TGF-β and collagen are elevated in CRSsNP across the world and may contribute to the remodeling process in this form of the disease.

As mentioned above, the pattern of remodeling in CRSwNP is known to be different compared to CRSsNP. Indeed, collagen was reduced in CRSwNP and TGF-β was either reduced or no difference compared to control subjects. In contrast to CRSsNP, the fact that fibrin deposition in the submucosa is especially high in NPs might represent a key event in the tissue remodeling process during CRSwNP. Takabayashi et al. recently reported that cross-linked fibrin was elevated and tissue plasminogen activator (t-PA) was reduced in NPs. The t-PA converts plasminogen to plasmin which mediates anti-inflammation by cross-linking fibrin and determines fibrinolysis. Importantly, type 2 cytokines, IL-4 and IL-13 which are elevated in NPs, reduced the expression of t-PA in epithelial cells. In a separate study, Takabayashi et al. reported that factor XIII-A (FXIIIa) was significantly elevated in NPs and it was produced by M2 macrophages (also known as alternatively activated macrophages). FXIIIa is a coagulation factor that can induce fibrin deposition by cross-linking fibrin and via the antifibrinolytic effect by mediating the binding of α2-plasmin inhibitor to fibrin. This data suggests that overexpression of FXIIIa and reduction of t-PA may facilitate the excessive deposition of fibrin in NPs via induction of fibrin cross-linking and the inhibition of fibrinolysis (Fig. 1). Very importantly, these factors are tightly controlled by the type 2 inflammatory milieu. Therefore they may not be a good fit to describe the mechanism of remodeling in non-eosinophilic Asian NPs (see following sections), which requires further study.

As we mention above, CRSSNP and CRSwNP are controlled by distinct inflammatory signals from various immune cells. Indeed, remodeling in CRSSNP might be controlled by TGF-β signaling. In contrast, remodeling in CRSwNP is regulated by type 2 inflammation. In addition, CRSwNP can be further divided into two phenotypes based on inflammatory patterns. Starting in the next section, we will describe the inflammatory patterns in CRSSNP and CRSwNP separately and also focus on the difference between the two types of CRSwNP, eosinophilic and non-eosinophilic.

Inflammatory patterns in CRSSNP

The mechanism of inflammation in CRSSNP is poorly understood. Van Zele et al. initially discovered that the type 1 cytokine IFN-γ was significantly elevated in CRSSNP in Belgium. Several groups confirmed the elevation of IFN-γ in CRSSNP in Belgium, China and Korea. In contrast, other groups could not find elevation of IFN-γ in CRSSNP in Japanese, Chinese and American populations. Another CRSSNP dependent cytokine could be TGF-β. Van Brauene et al. discovered that TGF-β was predominantly elevated in CRSSNP and several groups confirmed the elevation of TGF-β in CRSSNP in Asia although there is some discrepancy (see remodeling section). Since CRSSNP is a heterogeneous disease and is diagnosed by the phenotypic absence of NPs, the sample population may strongly affect the result. Future studies may require sub-classification of CRSSNP and investigation of the inflammatory patterns in each group.

Eosinophilic and non-eosinophilic CRSwNP

CRSwNP is well known to be characterized by type 2 inflammation and eosinophilia compared to CRSSNP or control sinus mucosa. However, the inflammation in CRSwNP varies based on race and regional differences. Several studies have indicated that 70–90% of CRSwNP in Europe and the United States showed eosinophilia in NPs. As mentioned above, the pattern of remodeling in CRSwNP is known to be different compared to CRSsNP. Indeed, collagen was reduced in CRSwNP and TGF-β was either reduced or no difference compared to control subjects. In contrast to CRSsNP, the fact that fibrin deposition in the submucosa is especially high in NPs might represent a key event in the tissue remodeling process during CRSwNP. Takabayashi et al. recently reported that cross-linked fibrin was elevated and tissue plasminogen activator (t-PA) was reduced in NPs. The t-PA converts plasminogen to plasmin which mediates anti-inflammation by cross-linking fibrin and determines fibrinolysis. Importantly, type 2 cytokines, IL-4 and IL-13 which are elevated in NPs, reduced the expression of t-PA in epithelial cells. In a separate study, Takabayashi et al. reported that factor XIII-A (FXIIIa) was significantly elevated in NPs and it was produced by M2 macrophages (also known as alternatively activated macrophages). FXIIIa is a coagulation factor that can induce fibrin deposition by cross-linking fibrin and via the antifibrinolytic effect by mediating the binding of α2-plasmin inhibitor to fibrin. This data suggests that overexpression of FXIIIa and reduction of t-PA may facilitate the excessive deposition of fibrin in NPs via induction of fibrin cross-linking and the inhibition of fibrinolysis (Fig. 1). Very importantly, these factors are tightly controlled by the type 2 inflammatory milieu. Therefore they may not be a good fit to describe the mechanism of remodeling in non-eosinophilic Asian NPs (see following sections), which requires further study.

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NPs had neutrophilia in Japan.33 In contrast, in the western country of Germany, Polzehl et al. showed that neutrophils were detected in early stage NPs but they were not elevated compared to control nasal mucosa.34 Van Zele et al. showed that neutrophils were significantly elevated in NPs in Belgium, although the degree of neutrophilia was significantly weaker than eosinophilia in those NPs.6 We also detected neutrophils and a subset of NPs showed elevation of neutrophils mixed with eosinophilia in the US (Fig. 2C, unpublished observation). This evidence indicates that neutrophilic CRSwNP is more commonly found in Asia. In contrast, neutrophil infiltration can be found in Europe and the US and it is generally mixed with eosinophilia. However, the degree of neutrophilia is weaker than eosinophilia.

Inflammatory patterns in eosinophilic CRSwNP

Since CRSwNP is characterized by eosinophilia in western countries, many researchers have investigated the role of eosinophil activation and survival factors in NPs. Hamilos et al. and others initially discovered that IL-5 was elevated in allergic NPs.35 Simon et al. and others confirmed the elevation of IL-5 in NPs and identified that IL-5 was a major eosinophil survival factor in NPs.46 After that, many investigators confirmed the elevation and importance of IL-5 in eosinophilic NPs across the world.5-7,8,23-25,29,31 Since IL-5 is a key factor of eosinophilia, Gevaert et al. performed clinical trials of anti-IL-5 in severe cases of NPs. Both mepolizumab and reslizumab, which are humanized anti-human IL-5 monoclonal antibodies, result in a significant reduction of NP size in 50–60% patients and IL-5 levels in nasal secretion can be predictive of response to anti-IL-5 therapy.

In addition to activation and survival, eosinophil recruitment into tissue should be another key event to induce eosinophilia. Chemokines are a large group of proteins that participate in recruitment of inflammatory cells into tissue sites by binding G protein-coupled receptors on their target cells. Among them, chemokine receptor CCR3 is predominantly expressed on eosinophils and 5 chemokines, MCP-4 (CCL13), RANTES (CCL5), eotaxin-1, -2 and -3 were all elevated in eosinophilic NPs.47 Furthermore, several investigators found a correlation between eotaxins and tissue eosinophilia in CRSwNP.47 These results indicate that eotaxins are key chemotactic factors for eosinophils in eosinophilic NPs.
Type 2 cytokines and epithelial cytokines in eosinophilic CRSwNP

Type 2 cytokines including IL-4, IL-5 and IL-13 are believed to control the inflammation in eosinophilic CRSwNP. As described above, IL-5 is a key activation and survival factor for eosinophils in NPs. In contrast, IL-4 and IL-13 are key factors that control mucus production in epithelial cells and IgE responses in B cells and plasma cells (Fig. 1). IL-4 and IL-13 also significantly contribute to several key inflammatory events in NPs including the activation of macrophages and remodeling. Very recently, a Phase 2a proof-of-concept study of dupilumab, which is a fully human monoclonal antibody against the α-subunit of the IL-4 receptor that blocks both IL-4 and IL-13 signaling, showed improvement of NP score as well as patient-reported symptom scores in moderate-to-severe CRSwNP (http://newsroom.regeneron.com/releasedetail.cfm?ReleaseID=873630). This reinforces the importance of type 2 inflammation in the pathogenesis of eosinophilic CRSwNP.

Type 2 cytokines, IL-4, IL-5 and IL-13, can be produced from several immune cells including Th2 cells, mast cells and group 2 innate lymphoid cells (ILC2s). Type 2 cytokines can be induced by both adaptive and innate signaling in these cell types. Recent evidence suggests that newly identified epithelial-derived cytokines, IL-25, IL-33 and thymic stromal lymphopoietin (TSLP), help to shape the local activation of type 2 immunity and exaggerated expression of these cytokines induces type 2 inflammatory diseases including atopic dermatitis and bronchial asthma.30-33 These epithelial-derived cytokines can have effects on both innate and adaptive type 2 immunity. Therefore many investigators have started to look at these epithelial cytokines in CRS.

IL-25 and IL-33 mainly contribute to innate type 2 inflammatory responses.25,33 IL-25 is a member of the IL-17 cytokine family. Unlike other IL-17 family cytokines, IL-25 promotes type 2 inflammation including eosinophilia even in Rag2 deficient mice.35 IL-25R is expressed on memory Th2 cells, basophils and ILC2s and IL-25 enhances type 2 cytokine production in these cells.56 In the case of CRS, only a few groups have investigated the presence of IL-25. Lam et al. reported that IL-25 mRNA was significantly elevated in ethmoid sinuses of CRSwNP compared to controls and CRSsNP.57 In contrast, Miljkovic et al. reported that IL-25 mRNA was significantly decreased in NPs compared to ethmoid sinuses of controls and CRSsNP.58 Our data showed that expression of mRNA for IL-25 was very low in sinuses and the level of IL-25 mRNA was similar between NPs and control sinus tissues (unpublished observations). In addition, no one has yet shown the presence of IL-25 protein in CRS. On balance, IL-25 may not be a
key regulator of type 2 inflammation in CRS. However, further study will be required to investigate the presence of IL-25 protein in eosinophilic NPs.

IL-33 is the latest member of the IL-1 cytokine family that induces type 2 inflammation.\(^5\) The production of IL-33 is known to be controlled by innate immune signaling including via P2 purinergic receptors.\(^\_\) IL-33 signals through a heterodimeric receptor complex consisting of IL-1RL1 (ST2) and IL-1RaCP. IL-33R is expressed on memory Th2 cells, basophils, mast cells, NKT cells and ILC2s.\(^\_\) IL-33 induces type 2 inflammation in Rag deficient mice suggesting that IL-33 mainly contributes to innate type 2 inflammation.\(^\_\) In the case of CRS, several groups reported the presence of IL-33 in CRS. Shaw et al., Lam et al. and Miljkovic et al. reported that IL-33 mRNA was highly expressed in nasal mucosa but was not elevated in NPs or other inflamed areas of the sinuses in CRSwNP.\(^7\)\(^7\)\(^7\) Baba et al. also found that the concentration of IL-33 protein was not significantly different between eosinophilic NPs and control sinus tissue although it was highly detected.\(^7\) These results indicate that IL-33 may contribute to the inflammation in CRS but is not elevated in CRSwNP. TSLP is an IL-7 like cytokine that induces type 2 inflammation. Several animal studies indicated that TSLP may contribute to both innate and adaptive type 2 inflammation.\(^6\)\(^6\) The production of TSLP is known to be controlled by both innate and adaptive immune signaling including via activation of toll like receptors and cytokine receptors.\(^5\)\(^7\)\(^8\)\(^7\)\(^7\)\(^7\)\(^7\)\(^7\)\(^7\) In contrast to IL-25 and IL-33, TSLP was reproducibly reported to have a pathogenic role in CRSwNP. Indeed, several groups reported that TSLP mRNA was significantly elevated in eosinophilic NPs in the US, Canada, China, Japan and Italy.\(^6\)\(^7\)\(^7\)\(^7\)\(^7\)\(^7\) Although it is notably difficult to detect TSLP protein in clinical samples by commercially available detection systems, Nagarkar et al. and Allakhverdi et al. found that TSLP activity was significantly elevated in NPs compared to control sinus tissue.\(^7\)\(^7\) This indicates that TSLP protein is elevated in NPs. Interestingly, Nagarkar et al. discovered cleavage products of TSLP when TSLP protein was incubated with NP tissue extracts.\(^7\) More importantly, they also found that the cleaved product might have higher activity than full-length TSLP.\(^7\) These results suggest that regulation of induction and posttranslational modifications in tissue control local TSLP-mediated type 2 inflammation. Very importantly, a recent clinical trial showed that AMG 157 which is a fully human anti-TSLP monoclonal antibody reduced allergen-induced early and late asthmatic responses in patients with mild allergic asthma.\(^7\) This indicates that TSLP is a key factor and an important therapeutic target in asthma. Since TSLP may also control type 2 inflammation in eosinophilic NPs, anti-TSLP may become a therapeutic agent for CRSwNP in the future.

Based on the literature, TSLP may be a key controller of type 2 immunity in eosinophilic NPs. IL-33 may also be involved in the inflammation although it is not elevated in CRS. Starting in the next section, we will describe the presence of type 2 cytokine producing cells and the potential role of epithelial-derived cytokines on type 2 inflammation in eosinophilic NPs.

**T cells and dendritic cells**

T cells are known to be elevated in eosinophilic NPs and T cell accumulation can be found in both mucosal and epithelial areas of NPs (Fig. 2G). Th2 cells are known as the primary source of type 2 cytokines. Recently, two separate groups have provided an in-depth analysis of Th cell subsets found in NPs. Derycke et al. reported that NPs were characterized by mixed T helper cells and only Th2 cells were significantly elevated in NPs in Belgium.\(^7\)\(^7\) Shi et al. reported that increased Th2 cells were found only in eosinophilic NPs although Th1 cells and Th17 cells were elevated in both eosinophilic and non-eosinophilic NPs in China.\(^7\)\(^7\) These results suggest that Th2 cells are highly elevated in eosinophilic CRSwNP across the world.

Although TSLP directly promoted Th2 differentiation from naïve T cells in the absence of antigen presenting cells (APC) in mice,\(^7\)\(^7\) TSLP dependent-Th2 differentiation required APC in humans.\(^7\)\(^8\) TSLP-R was not expressed on naïve T cells and TSLP did not directly induce Th2 cytokine production in naïve T cells in humans, although expression of TSLP-R was weakly induced by TCR-dependent activation on naïve T cells and by allergen plus TSLP on a subset of Th2 cells in atopic subjects.\(^7\)\(^7\)\(^7\) Therefore TSLP dependent-Th2 differentiation is believed to be controlled by dendritic cells (DCs) in humans.

Dendritic cells are known to be important in skewing Th responses in the mucosa. Human DCs can be divided into two major subsets, myeloid DCs (mDCs) and plasmacytoid DCs (pDCs).\(^7\) Myeloid DCs act as strong antigen presenting cells and one of their key functions is to control polarization of T helper cells. In contrast, pDCs are less effective at antigen presentation than mDCs but strongly promote antiviral immunity. Recently, three groups identified DC subsets in NPs and found that mDCs were significantly elevated in NPs in the US, Belgium and China.\(^7\)\(^7\)\(^7\)\(^7\) Myeloid DCs can be classified into two subsets in humans, mDC1 and mDC2, which can be differentiated by cell surface expression of CD1c and CD141, respectively. Poposki et al. and Pezato et al. further differentiated mDC phenotypes and found that both mDC1s and mDC2s were significantly elevated in NPs although mDC2s were a minor population of mDCs in NPs.\(^7\)\(^7\)\(^7\)\(^7\)\(^7\)\(^7\) Although it is notoriously difficult to detect TSLP protein in clinical samples by commercially available detection systems, Nagarkar et al. and Allakhverdi et al. found that TSLP activity was significantly elevated in NPs compared to control sinus tissue.\(^7\)\(^7\) This indicates that TSLP protein is elevated in NPs. Interestingly, Nagarkar et al. discovered cleavage products of TSLP when TSLP protein was incubated with NP tissue extracts.\(^7\) More importantly, they also found that the cleaved product might have higher activity than full-length TSLP.\(^7\) These results suggest that regulation of induction and posttranslational modifications in tissue control local TSLP-mediated type 2 inflammation. Very importantly, a recent clinical trial showed that AMG 157 which is a fully human anti-TSLP monoclonal antibody reduced allergen-induced early and late asthmatic responses in patients with mild allergic asthma.\(^7\) This indicates that TSLP is a key factor and an important therapeutic target in asthma. Since TSLP may also control type 2 inflammation in eosinophilic NPs, anti-TSLP may become a therapeutic agent for CRSwNP in the future.

In contrast to mDC2, the mechanisms of mDC1-related Th2 polarization are partially identified. Shi et al. evaluated DC subsets from NPs of eosinophilic and non-eosinophilic patients. They found that mDC1s isolated from either eosinophilic or non-eosinophilic NPs could skew naïve T cells toward Th1 and Th17 phenotypes, but only DCs from eosinophilic NPs were able to skew naïve T cells toward a Th2 phenotype.\(^7\) They also investigated the mechanism of Th2 skewing by mDC1s from eosinophilic NPs. They found that the frequency of mDC1s expressing either OX40 ligand (OX40L) or programmed death ligand-1 (PD-L1) was significantly increased in eosinophilic NPs.\(^7\) Furthermore, they also found that blockade of either OX40L or PD-L1 suppressed the skewing of Th2 cells to produce type 2 cytokines.\(^7\) TSLP-R is known to be highly expressed on mDC1s and TSLP-stimulated mDCs induce naïve CD4+ T cells to differentiate into Th2 cells.\(^7\) Importantly, this response is known to be via OX40L and TSLP induces expression of OX40L on mDCs.\(^7\) This evidence indicates that mDC1s in eosinophilic NPs may be primed by TSLP and thereafter control Th2 responses in NPs. Since TSLP is known to strongly induce a Th2-polarized response although the mechanisms were not clear.\(^7\) This suggests that the elevation of mDC2s in NPs may contribute to the local Th2-polarized environment. However, the mechanism needs further investigation.

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Since epithelial cells are one of the main sources of TSLP, we initially hypothesized that mDC1s might be accumulated in the epithelium of NPs, yet we found that accumulation of mDC1s can be detected in lamina propria but not in epithelium.\(^7\) However, CD1c+ mDC1s were frequently found just below the surface of the epithelium in NPs (Fig. 2F). This suggests that it may be easy for mDC1s to interact with epithelial cells and epithelial cell derived cytokines including TSLP in NPs. This evidence suggests that TSLP
may control adaptive type 2 inflammation through activation of mDCs and induction of Th2 differentiation (Fig. 1).

The mechanism of accumulation of mDCs in eosinophilic NPs is not clear. We recently reported that the chemokines CCL18 and CCL23 were significantly elevated in NPs. CCL23 is known to recruit mDCs via the receptor CCR1. We also found that levels of CCL23 significantly correlated with CCR1 and CD1c in NP tissues. CCL18 is known to be chemotactic for immature mDC1. These results suggest that overproduction of CCL18 and CCL23 may contribute to the accumulation of mDC1 in NPs. CCL18 is also known to recruit Th2 cells via CCR8. Interestingly, we also identified that CCL18 and CCL23 were mainly produced by M2 macrophages and eosinophils, respectively. CCL23 also potently recruits CCR1 monocytes and macrophages. Since eosinophilic NPs are in a type 2 microenvironment, recruited macrophages can be differentiated into the M2 phenotype by type 2 cytokines. CD163 M2 macrophages are known to be accumulated in NPs (Fig. 2E). They are known to produce eosinophil chemokines including eotaxins and MCP-4. This data indicates that production of CCL23 from eosinophils, recruitment of macrophages, differentiation into M2 macrophages, and production of eosinophil chemokines from M2 macrophages may constitute a positive feedback loop to amplify the eosinophilia in NPs (Fig. 1).

Mast cells

Mast cells play key roles in host defense, homeostasis, tissue repair, and mechanisms of allergic inflammation. Mast cells express the high-affinity IgE receptor, FcεRI. Upon cross-linking of the FcεRI via IgE-antigen, mast cells release several pre-stored mediators, including histamine and proteases and synthesize many mediators including prostaglandins (PGs), cysteinyl leukotrienes (CysLTs) and type 2 cytokines especially IL-5 and IL-13. Recent reports suggest that a subset of patients with asthma and eosinophilic esophagitis that displayed a type 2 cytokine high-phenotype have elevated mast cells. Interestingly, mast cells primary accumulated in the epithelium. In addition, intraepithelial mast cells in these patients showed a mast cell-tryptase (MC-T) but not mast cell-tryptase/chymase (MC-TC) phenotype. In the case of CRS, our group found that mast cells were significantly elevated in mucosal and glandular epithelium but not within the lamina propria in NPs (Fig. 2D). Interestingly, mast cell phenotypes were different in mucosal and glandular epithelium. Takabayashi et al. found that mucosal epithelial mast cells showed an MC-T phenotype. In contrast, glandular epithelial mast cells showed an MC-TC phenotype in the US. Cao et al. also found significant elevation of MC-T in the mucosal epithelium of eosinophilic NPs in China. These results suggest that mast cells are highly elevated in the epithelium of eosinophilic CRS/wNP across the world. However, functional differences between MC-T and MC-TC in the pathogenesis of CRS will need to be further investigated.

Mast cells are known to express IL-1R, IL-33R (ST2) and TSLPR on the cell surface and to respond to these ligands. Although TSLP alone isn’t sufficient to stimulate mast cells, TSLP synergizes with IL-1 and IL-33; potently activating mast cells to produce type 2 cytokines. As described, mast cell numbers were increased only within the NP epithelium in patients with eosinophilic CRS/wNP. However, the specific role of the epithelium in the direct activation of mast cells was not clear. We hypothesized that epithelial cells can produce a combination of cytokines that directly activates intraepithelial mast cells to produce type 2 cytokines. We therefore investigated the effect of viruses, TLR ligands and cytokines on the interaction of mast cells and epithelial cells in vitro. We found that airway epithelial cells directly promoted production of type 2 cytokines in mast cells during viral infection through the production of IL-1 and TSLP and that this occurred only in a type 2 microenvironment. We initially expected that IL-33 released from epithelial cells contributes to this reaction. However, we could not detect IL-33 protein in poly(I:C) stimulated or influenza infected epithelial cells and ST2-Fc fusion protein could not block this response. This suggests that intraepithelial mast cells in eosinophilic NPs may contribute to the production of type 2 cytokines via interaction with epithelial cytokines TSLP and IL-1 in response to pathogens on epithelial cells. Interestingly, Ikura et al. found that IL-1 and IL-33 enhanced IgE receptor mediated type 2 cytokine production in mast cells. Since mast cells produce type 2 cytokines through epithelial-derived cytokines and IgE-mediated reactions, mast cells may contribute to both innate and adaptive type 2 inflammation in eosinophilic NPs (Fig. 1).

ILC2

The last population of type 2 cytokine producing cells to discuss is innate lymphoid cells (ILCs). ILCs are a recently identified minor population of immune cells that lack antigen receptors and lineage (Lin) markers. ILCs are activated by innate immune signaling, including pathogens and innate cytokines. Like T helper cells, ILCs can be classified into 3 subsets, ILC1, ILC2 and ILC3 dependant on the production of cytokines. Among them, ILC2s produce large quantities of type 2 cytokines (including IL-5 and IL-13). The first evidence for ILC2-like cells was reported by Fort et al. in 2001. An ILC2 population was identified by several groups in 2010 in mice and variously named nuocytes, natural helper cells, innate helper 2 cells and others. In 2013, Spits et al. organized the nomenclature of the ILC family members and defined group 2 ILCs (ILC2s) as ILCs that produce type 2 cytokines and are dependent on GATA3 and RORα for their development and function.

Human ILC2s were discovered by Mjosberg et al. in 2011 and were defined as CD45-, Lin-, CD127+, CRTH2+ cells. Importantly, they also identified an ILC2 population in NPs of Dutch patients and found that ILC2s were significantly elevated in NPs compared to control sinus mucosa. Several groups have now confirmed the elevation of ILC2 in NPs in the US and Australia. In my knowledge, the presence of ILC2s in eosinophilic NPs in the Asian population has yet not been investigated. Future study is required to identify ILC2s in the Asian population and investigate whether ILC2s are significantly elevated in eosinophilic NPs compared to non-eosinophilic NPs and control sinus mucosa.

ILC2s are known to express IL-33R and TSLP-R on the cell surface. IL-33 and TSLP are known to induce type 2 cytokine production in ILC2s, although the effect by individual cytokines is minimal. However, the combination of TSLP and IL-33 synergistically enhances the production of type 2 cytokines including IL-5 and IL-13 in human ILC2s isolated from peripheral blood and NPs. As described above, TSLP is significantly upregulated in eosinophilic NPs. IL-33 is present but not elevated in NPs. This suggests that ILC2s may contribute to the local innate type 2 inflammation in eosinophilic NPs by responding to TSLP and IL-33 with the production of type 2 cytokines.

CRTH2 is now known to be one of the key markers of human ILC2. CRTH2 is a chemokine receptor homologous molecule and is known as the PGD2 receptor 2 (DP2). PGD2 is known to recruit Th2, eosinophils and basophils via CRTH2. Chang et al. recently discovered that PGD2 also induced chemotaxis of human ILC2s. Xue et al. found that PGD2 dependent migration of ILC2 was via CRTH2. Importantly, Xue et al. also found that PGD2 strongly induced the production of type 2 cytokines including IL-4, IL-5 and IL-13 in ILC2s via the activation of CRTH2. Mast cells are the one of the major sources of PGD2. Xue et al. also looked at the role of
mast cell products on the activation of ILC2s and found that supernatants of degranulated mast cells induced ILC2 migration and production of type 2 cytokines in ILC2. Mast cells also produce other pro allergic arachidonic acid metabolites called CysLTs including LTC4 upon degranulation. Doherty et al. found that CysLTR1 was expressed on ILC2s, and LTC4 and LTD4 strongly induced the production of type 2 cytokines in ILC2s. Since mast cells are highly accumulated and many of them are activated and degranulated in eosinophilic NPs, degranulation of mast cells may further enhance the recruitment of ILC2s and the production of type 2 cytokines in ILC2s via release of PGD2 and LTC4/D4 from mast cells and activation of CRTH2 and CysLTR1 on ILC2s in eosinophilic NPs (Fig. 1).

B-lineage cells

We have described immune cells that produce type 2 cytokines and cells that control type 2 cytokine production in these cells. In the last section, we will focus on the B-lineage cells that can be activated by type 2 cytokines in eosinophilic NPs. In general, B cell maturation and expansion occur in germinal centers of the lymphoid organs. However, there are compelling reasons to believe that local proliferation and activation of B-lineage cells, B cells, plasmablasts and plasma cells, are of central pathogenic importance in several airway inflammatory diseases. The production of IgA and IgE from B-lineage cells is known to be critical to allergic disease via the activation of airway eosinophils and mast cells, respectively, in response to antigen exposure. Several groups have shown that extensive class switch recombination and antigen specific IgA and IgE production can occur in the nasal and bronchial mucosa as part of inflammatory diseases such as allergic rhinitis and asthma. Due to the recognized importance of IgA and IgE in the airway, the local activation of B-lineage cells in the airway has been proposed to be an important event in the pathogenesis of allergic airway diseases.

In the case of CRS, B cell accumulation can be found in the lamina propria of NPs (Fig. 2H). Van Zele et al. found that CD20+ B cells and CD138+ plasma cells are elevated in NPs by immunohistochemistry. Several groups also found B cell clusters and B cell follicle-like structures in NPs. However, unlike the germinal center, the structure of these local follicles was not well organized (Fig. 2I). In addition, we also found B cell clusters and follicles in the nasal mucosa of CRSsNP and control subjects and found that they were not elevated in NPs. The role of B cell clusters and follicles in nasal mucosa needs further investigation. Our group also characterized B-lineage cells in CRS by flow cytometry and found that B-lineage cells, B cells, plasmablasts and plasma cells, were all elevated in NPs compared to uncinate process tissues from control subjects and patients with CRSsNP.

In addition to accumulation of B-lineage cells in NPs, many groups reported the elevation of IgE in NPs. Gevaert P et al. found that total IgE, Staphylococcus aureus enterotoxin-specific IgE and IgE positive cells were elevated in NPs. Our group also evaluated the immunoglobulin isotypes, IgM, IgG, IgA and IgE in the nasal mucosa of CRSsNP and control subjects and found that they were not elevated in NPs. The role of B cell clusters and follicles in nasal mucosa needs further investigation. Our group also characterized B-lineage cells in CRS by flow cytometry and found that B-lineage cells, B cells, plasmablasts and plasma cells, were all elevated in NPs compared to uncinate process tissues from control subjects and patients with CRSsNP.

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Several factors are involved in B-lineage cell responses. As described above, type 2 cytokine IL-13 is known to be elevated in eosinophilic NPs. IL-13 is a critical inducer of IgE class switch recombination and IgE production in B cells. We also found that B cell-activation factor of the TNF family (BAFF) was significantly elevated in NP tissues and nasal lavage from patients with CRSsNP. Gevaert et al. confirmed elevation of BAFF in NPs in Belgium. BAFF is an essential factor for B cell maturation and survival. BAFF also induces T cell-dependent and independent immunoglobulin class switching and production. Importantly, expression of BAFF positively correlated with B cell marker CD20 and one of the BAFF receptors, TACI, in sinus mucosa suggesting that BAFF may be a key B cell activation factor in NPs. Since BAFF cannot induce migration of B-lineage cells, we also looked at chemokines and found elevation of the B cell chemokine CXCL12 (SDF-1) and CXCL13 (BCL) in NPs. This data indicates that chemokines CXCL12 and CXCL13 may contribute to the initial recruitment of B-lineage cells. BAFF may be involved in the proliferation and activation of B-lineage cells and IL-13 is a key factor for the local production of IgE in NPs.

IgE induces allergic inflammation by activation of mast cells and basophils and plays an important role in allergic diseases including asthma. Omalizumab, which is a humanized anti-human IgE monoclonal antibody, is approved and known to be effective in the treatment of patients with severe allergic asthma. As described above, mast cells and IgE are highly elevated in eosinophilic NPs. We also found that basophils were elevated in NPs in the US. Gevaert et al. performed clinical trials of anti-IgE in patients with CRSsNP who were comorbid with asthma. They found that Omalizumab showed significant reduction in NP score and had a beneficial effect on QOL scores. This suggests that IgE can be an important therapeutic target for patients with eosinophilic NPs.

As mentioned above, the cytokine BAFF is elevated in NPs. BAFF is now known as a pathogenic factor for autoimmune diseases. More importantly anti–BAFF (Belimumab) has very recently been approved for the treatment of lupus. We therefore looked at the presence of autoantibodies in CRS. We found that anti-dsDNA antibody and anti-BP-180 antibody were significantly elevated in NPs. Since we could not find elevation of autoantibodies in the serum of polypoid patients, this should be a local phenomenon. Interestingly we found that anti-dsDNA antibodies were more elevated in NPs who had revision surgery. This indicates that autoantibodies may be a pathogenic factor and/or biomarker of severe NPs. Further study will be required to identify the pathogenic role of autoantibodies in NPs.

Conclusions

This review primary focused on immune cells and cytokines that control initiation and amplification of inflammation in CRS. Although CRSsNP is known to be characterized by type 1 inflammation with elevated levels of IFN-γ, CRSsNP is a heterogeneous disease and sub-classification of this disease and further investigation of the inflammatory patterns in each group might be required in the future. Although the majority of CRSsNP cases in western countries show type 2 inflammation and eosinophilia, about 50% of CRSsNP in Asian countries show a non-eosinophilic phenotype. However, the prevalence of eosinophilic NPs has increased in Asia in the past 10–20 years and may continue to increase in the future. Inflammation in eosinophilic CRSsNP is controlled by type 2 cytokines including IL-5 and IL-13 that can be produced from Th2, mast cells and ILC2 in eosinophilic NP. Epithelial cytokines including TSLP might control production of
type 2 cytokines in these cell types. Recent clinical trials showed the beneficial effect on eosinophilic NPs and/or asthma by monoclonal antibodies against IL-5, IL-4Rα, IgE and TSLP suggesting that they can be novel therapeutic targets for eosinophilic CRSwNP.

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Conflict of interest

The author has no conflict of interest to declare.

References


