

ORIGINAL ARTICLE

The role of interleukin-33 in chronic rhinosinusitis

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► Additional material is published online only. To view please visit the journal online (<http://dx.doi.org/10.1136/thoraxjnl-2016-208772>).

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Received 13 April 2016

Revised 30 October 2016

Accepted 31 October 2016

Published Online First

24 November 2016



► <http://dx.doi.org/10.1136/thoraxjnl-2016-209655>



CrossMark

To cite: Kim D-K, Jin HR, Eun KM, *et al.* *Thorax* 2017;**72**:635–645.

ABSTRACT

Rationale Interleukin (IL)-33, a new member of the IL-1 family, is constitutively expressed in epithelial tissues and lymphoid organs and plays an important role in the pathogenesis of allergic disease. However, the role of IL-33 in chronic rhinosinusitis with nasal polyps (CRSwNP) remains unclear.

Objective To investigate the role of IL-33 in the pathophysiology of CRSwNP.

Methods We investigated IL-33 expression and its cellular origins in the nasal polyps (NPs) of human subjects by immunohistochemistry (IHC), quantitative reverse transcription PCR (qRT-PCR), and multiplex cytokine assays. Correlations between IL-33 expression and other inflammatory markers were also explored. To investigate the role of IL-33 in CRSwNP, anti-IL-33 antibody was used in a murine model of CRS.

Results Uncinate process tissues from control (19), CRSwNP (61), CRSwNP (69) and NP tissues (71) were used in this study. Increased expression of IL-33 mRNA and protein in patients with CRSwNP compared with controls was observed. The concentration of IL-33 protein in CRSwNP was positively correlated with the number of neutrophils and the expression of several Th1 and Th17 inflammatory markers, including interferon (IFN)- γ , IL-1 β , tumour necrosis factor (TNF)- α , IL-17A, IL-22, and various markers for neutrophil recruitment. However, protein levels of IL-5 and quantity of eosinophils were inversely correlated with levels of IL-33. The expression of tissue inhibitor of metalloproteinase (TIMP)-1 was negatively correlated with IL-33 protein levels, while the expression of matrix metalloproteinase (MMP)-2 and MMP-9 was positively correlated with IL-33 protein levels. In animal studies, IL-33 expression was upregulated in the CRSwNP group compared with controls. Anti-IL-33 treatment reduced the thickness of oedematous mucosa, subepithelial collagen deposition, and infiltration of neutrophils, but infiltration of eosinophils was not reduced. This treatment also inhibited the expression of neutrophilic inflammatory cytokines, but not IL-4. In addition, the expression of intracellular adhesion molecule 1, vascular adhesion molecule 1 and CXCL-2 in the nasal mucosa was suppressed in mice treated with anti-IL-33 antibody.

Conclusions Our data suggest a role for IL-33 in the pathogenesis of CRSwNP via neutrophil recruitment. Therefore, anti-IL-33 may provide a new treatment strategy to target infiltrating neutrophils in CRSwNP.

INTRODUCTION

Interleukin (IL)-33 is a novel member of the IL-1 superfamily and constitutively expressed at high levels in epithelial cells from human and mouse

Key messages**What is the key question?**

- Does interleukin (IL)-33 play an important role in the development of chronic rhinosinusitis with nasal polyp (CRSwNP)?

What is the bottom line?

- IL-33 expression is elevated in nasal tissues of patients with CRSwNP, and expression of this cytokine is associated with Th1/Th17 cytokines, neutrophil recruitment and remodelling proteins.

Why read on?

- IL-33 is one of the crucial mediators of pathogenesis in CRSwNP through neutrophil recruitment, and a promising target for the treatment of patients with CRSwNP.

tissues during homeostasis.^{1–3} IL-33 is produced by various types of immune cells such as macrophages and dendritic cells and by non-immune cells, including endothelial, epithelial, and smooth muscle cells.⁴ Biologically active full-length IL-33 can be released extracellularly by necrotic cells after tissue damage caused by airborne allergens, viruses and air pollutants.^{4–5} After release, IL-33 subsequently serves as an ‘alarmin’ by triggering inflammatory response by activating various types of immune cells, including eosinophils, Th2 cells and innate lymphoid cells type 2 (ILC2s).^{6–7} Thus, this cytokine is considered to play a crucial role in Th2 inflammation. However, besides promoting the Th2 response, IL-33 plays a diverse role in various inflammatory diseases. Some evidence suggests a role for IL-33 in recruiting neutrophils by increased production of the chemokine ligand 1 (CXCL-1) and preventing the downregulation of the receptor for CXCR-2.^{8–9} Other research shows that IL-33 can amplify macrophage polarisation towards an alternatively activated phenotype and can enhance production of tumour necrosis factor (TNF)- α .^{10–11} Moreover, in various rheumatological diseases, IL-33 appears to have proinflammatory effects, such as activating synovial fibroblasts and mast cells, and inducing neutrophil migration.^{12–14}

Chronic rhinosinusitis (CRS) is one of the most common chronic rhinological diseases and results in a decreased quality of life for affected people.^{15–16} CRS is characterised by accumulation of inflammatory cells and marked tissue

remodelling, and is currently divided into two clinical phenotypes: CRS without nasal polyps (CRSsNP) and CRS with nasal polyps (CRSwNP), based primarily on endoscopic findings.¹⁷ CRSwNP often show higher disease severity and risk of recurrence compared with CRSsNP.¹⁸

More recently, expression of IL-33 and its receptor ST2 was also investigated in patients with CRS,^{19–21} and two of these studies reported elevated expression of ST2,^{19 20} implying that the IL-33/ST2 pathway may function in nasal polypogenesis. Despite these findings, the specific role of IL-33 in the pathophysiology of patients with CRSwNP still remains unclear. Therefore, in this study we investigated the expression of IL-33 and the relationship between IL-33 and other inflammatory markers in sinonasal tissues from patients with CRSwNP. We also evaluated the therapeutic potential of blocking IL-33 in a murine model of CRS.

METHODS

Detailed methods can be found in the online supplementary data (<http://www.thorax.bmj.com>).

Patients and tissue samples

Sinonasal and polyp tissues were obtained from patients with CRS during routine functional endoscopic sinus surgery. All patients provided written informed consent for study participation, and this study was approved by the institutional review board of Seoul National University Hospital, Boramae Medical Center. The diagnosis of CRS was based on personal history, physical examination, nasal endoscopy and CT findings of the sinuses according to the 2012 European position paper on rhinosinusitis and nasal polyps (EPOS) guidelines.¹⁷ Exclusion criteria were as follows: (1) younger than 18 years of age; (2) prior treatment with antibiotics, systemic or topical corticosteroids, or other immune-modulating drugs for 4 weeks before surgery; and (3) unilateral rhinosinusitis, antrochoanal polyp, allergic fungal sinusitis, cystic fibrosis or immotile ciliary disease (see online supplementary figure S1). Control tissues were obtained during other rhinological surgeries, such as skull base, lacrimal duct or orbital decompression surgery, from patients without any sinonasal diseases. In this study, we obtained unciniate process (UP) tissue from controls and patients with CRS, including those with CRSsNP or CRSwNP. Each sample obtained was divided into three parts: one third was fixed in 10% formaldehyde and embedded in paraffin for histological analysis, another third was immediately frozen and stored at -80°C for subsequent isolation of mRNA, and the final third of the tissue was submerged in 1 mL phosphate-buffered saline (PBS) supplemented with 0.05% Tween-20 (Sigma-Aldrich, St Louis, Missouri, USA) and 1% PIC (Sigma-Aldrich) per 0.1 g of tissue. This tissue was homogenised with a mechanical homogeniser at 1000 rpm for 5 min on ice. After homogenisation, the suspensions were centrifuged at 3000 rpm for 10 min at 4°C . The supernatants were separated and stored at -80°C for further analysis of cytokines and other inflammatory mediators.²² The atopic status of study patients was evaluated using the ImmunoCAP assay (Phadia, Uppsala, Sweden), which detects IgE antibodies against common aeroallergens (house dust mites, moulds, trees, weeds, grass, and animal dander). Patients were considered atopic if the allergen-specific IgE level was greater than 0.35 KU/L.²³ A diagnosis of asthma was performed by an allergist based on medical history and lung function analysis, including methacholine challenge tests. Nasal polyps (NPs) were classified into two groups: eosinophilic NP, which was defined as having an eosinophil count of more than 10% of the

inflammatory cell population, and non-eosinophilic NP, which did not fulfil this criterion.²⁴ Additional information and details of the subjects' characteristics are listed in [table 1](#).

Immunohistochemistry

Immunohistochemistry (IHC) staining was performed with Polink-2, polymerised horseradish peroxidase (HRP), and broad DAB Detection System (Golden Bridge International Labs., Washington, USA). Paraffin sections were treated with 3% hydrogen peroxide (H_2O_2) and then incubated with primary antibodies and biotinylated secondary antibodies. The numbers of positive cells in epithelia, glands and submucosa were determined in the densest part of the tissue in five high-power fields (HPF; $\times 400$) by two independent observers, and average values were determined. Detailed IHC procedures are described in the Methods section in the Online Repository at <http://www.thorax.bmj.com>.

Quantitative real-time PCR for inflammatory markers

The mRNA levels in UP and NP tissues were evaluated using quantitative real-time (qRT)-PCR analysis as previously described.²⁴ Detailed qRT-PCR conditions are described in the Methods section in the Online Repository (<http://www.thorax.bmj.com>).

Measurement of cytokine protein levels in tissue homogenates

The protein concentrations for tissue extracts were determined using the Pierce 660 nm Protein Assay Kit (Thermo Scientific Inc, New York, USA). Samples were thawed at room temperature and vortexed to ensure a well mixed sample. Multiplex cytokine analysis kits (IL-1 β , IL-5, IL-17A, IL-22, IL-33, interferon (IFN)- γ , CXCL-8, MPO, GRO- α , GRO- β , intercellular adhesion molecule (ICAM)-1, matrix metalloproteinase (MMP)-2, MMP-9, tissue inhibitor of metalloproteinase (TIMP)-1 and TNF- α) were obtained from R&D systems (Cat. No. LMSAHM) and data were collected using Luminex 100 (Luminex, Austin, Texas, USA). Data analysis was performed using the MasterPlex QT V2.0 (MiraBio, Alameda, California, USA). All assays were run in duplicate according to the manufacturers' protocol. Sensitivity of each cytokine is as follows: IL-1 β (0.8 pg/mL), IL-5 (0.5 pg/mL), IL-17A (1.8 pg/mL), IL-22 (11.7 pg/mL), IL-33 (1.8 pg/mL), IFN- γ (0.4 pg/mL), CXCL-8 (1.8 pg/mL), MPO (20.4 pg/mL), GRO- α (5.3 pg/mL), GRO- β (7.86 pg/mL), ICAM-1 (87.9 pg/mL), MMP-2 (108 pg/mL),

Table 1 Patient characteristics and study methodology

Total no. of subjects	Control	CRSsNP	CRSwNP	
	N=19	N=61	N=69	N=71
Tissue used	UP	UP	UP	NP
Age (years), mean (SD)	45 (19)	49 (13)	48 (14)	49 (14)
Atopy, N (%)	6 (31)	18 (29)	27 (39)	29 (40)
Asthma, N (%)	0	3 (5)	5 (7)	5 (7)
Lund-Mackay CT score	0 (0)	8.9 (5.3)	15.3 (6.1)	15.6 (5.1)
Methodologies used				
Tissue IHC	9 (47%)	25 (41%)	24 (35%)	58 (82%)
Tissue mRNA	17 (89%)	61 (100%)	69 (100%)	71 (100%)
Homogenate ELISA	9 (47%)	7 (11%)	27 (39%)	36 (51%)

CRSsNP, chronic rhinosinusitis without nasal polyps; CRSwNP, chronic rhinosinusitis with nasal polyps; IHC, immunohistochemistry; NP, nasal polyp; UP, unciniate process.

MMP-9 (13.6 pg/mL), TIMP-1 (3.42 pg/mL) and TNF- α (1.2 pg/mL). All the protein levels in the tissue homogenate were normalised to the concentration of total protein (mg/mL).

Murine model of chronic rhinosinusitis

All animal experiments were approved by the IACUC of Boramae Medical Center and were performed under strict governmental and international guidelines on animal experimentation. Nasal polypoid lesions in mice were induced according to a previously established protocol,²⁵ which has been confirmed by multiple studies.^{26–28} Thirty-two female BALB/c mice (4 weeks old, 20–25 g) were divided into three groups: PBS-treated (PBS, n=10), CRS model group (CRS, n=12), and CRS model group treated with anti-IL-33 (R&D systems, Minneapolis, Minnesota, USA; CRS+ α IL-33, n=10). Detailed animal experiments are described in the Methods section in the Online Repository at <http://www.thorax.bmj.com>.

Cytokines from nasal lavage fluid in murine model

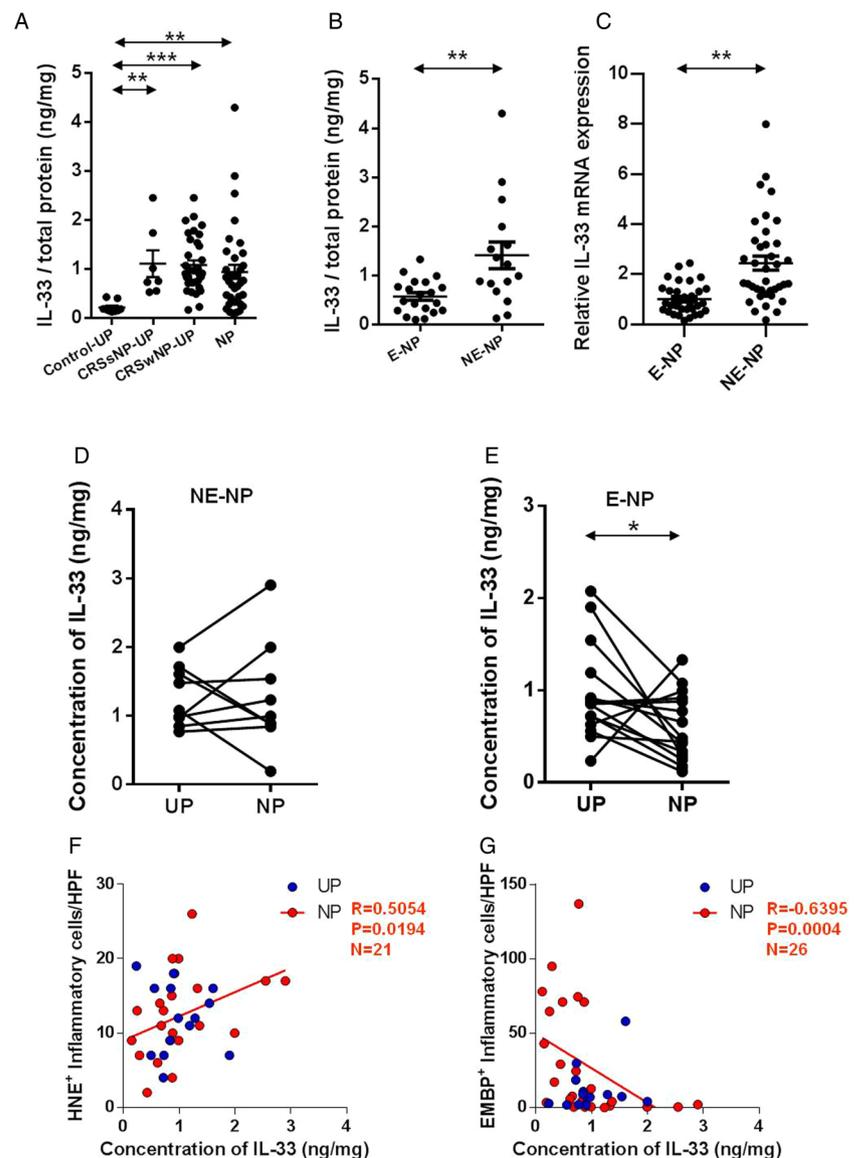
Nasal lavage was performed as previously described.²⁹ After partial tracheal resection under deep anaesthesia, a micropipette was inserted into the posterior choana through the tracheal

opening in the direction of the upper airway. Each nasal cavity was gently perfused with 200 μ L PBS, and the fluid from the nostril was collected and centrifuged. Supernatants were stored at -80°C . Levels of IFN- γ , IL-4, IL-17A and transforming growth factor (TGF)- β 1 in the nasal lavage fluids were measured using ELISA kits purchased from BioLegend (San Diego, California, USA). The lower detection limits of these ELISA kits were 8 pg/mL for IFN- γ , 0.5 pg/mL for IL-4, 2.7 pg/mL for IL-17A and 2.3 pg/mL for TGF- β 1.

Statistical analysis

Statistical analyses were performed using GraphPad Prism software V.6.0 (GraphPad software Inc, La Jolla, California, USA). In this study, the Kruskal–Wallis test and the Mann–Whitney U test with a two-tailed test for unpaired comparisons were used. For comparisons among groups, the Kruskal–Wallis test was used to establish the significant difference (human study groups: Control-UP, CRSSNP-UP, CRSwNP-UP, NP; animal study groups: Control, POLY, POLY+ α IL-33). The Mann–Whitney U test was then used for between-group comparisons with Bonferroni adjustment for multiple comparisons to

Figure 1 Expression of interleukin (IL)-33 in nasal tissues according to the different types of chronic rhinosinusitis. Comparison of IL-33 expression within the uncinata process (UP) tissues from control subjects, patients with chronic rhinosinusitis without nasal polyps (CRSSNP), and patients with chronic rhinosinusitis with nasal polyps (CRSwNP), and from nasal polyp (NP) tissues from patients with CRSwNP. (A) Protein levels of IL-33 among groups and (B and C) Difference between eosinophilic and non-eosinophilic NPs in the protein and mRNA levels. IL-33 expression was compared between UP and NP tissues in (D) non-eosinophilic CRSwNP (NE-NP) and (E) eosinophilic CRSwNP (E-NP). (F) Number of HNE-positive and (G) EMBP-positive inflammatory cells was counted from five of the densest areas (high power field; $\times 400$) and those were compared with the concentration of IL-33 homogenated. IL-33 antibody from R&D (accession #O95760) was used in multiplex cytokine analysis. * $p < 0.05$, ** $p < 0.01$, and *** $p < 0.001$.



control type I error rates. Adjusted p values were presented in this text. The Wilcoxon matched-paired signed rank test was also used to compare the expression of IL-33 between UP and NP tissues in the same patients. The Spearman correlation test was used to determine bivariate relationships. The significance level was set at α value of 0.05.

RESULTS

Expression of IL-33 and cellular origin in CRSwNP

To investigate expression of IL-33 in patients with CRSwNP, tissues were collected from patients with CRSsNP (UP tissue), patients with CRSwNP (NP and UP tissue) and control subjects (UP tissue). IL-33 protein levels were higher in UPs of CRSsNP

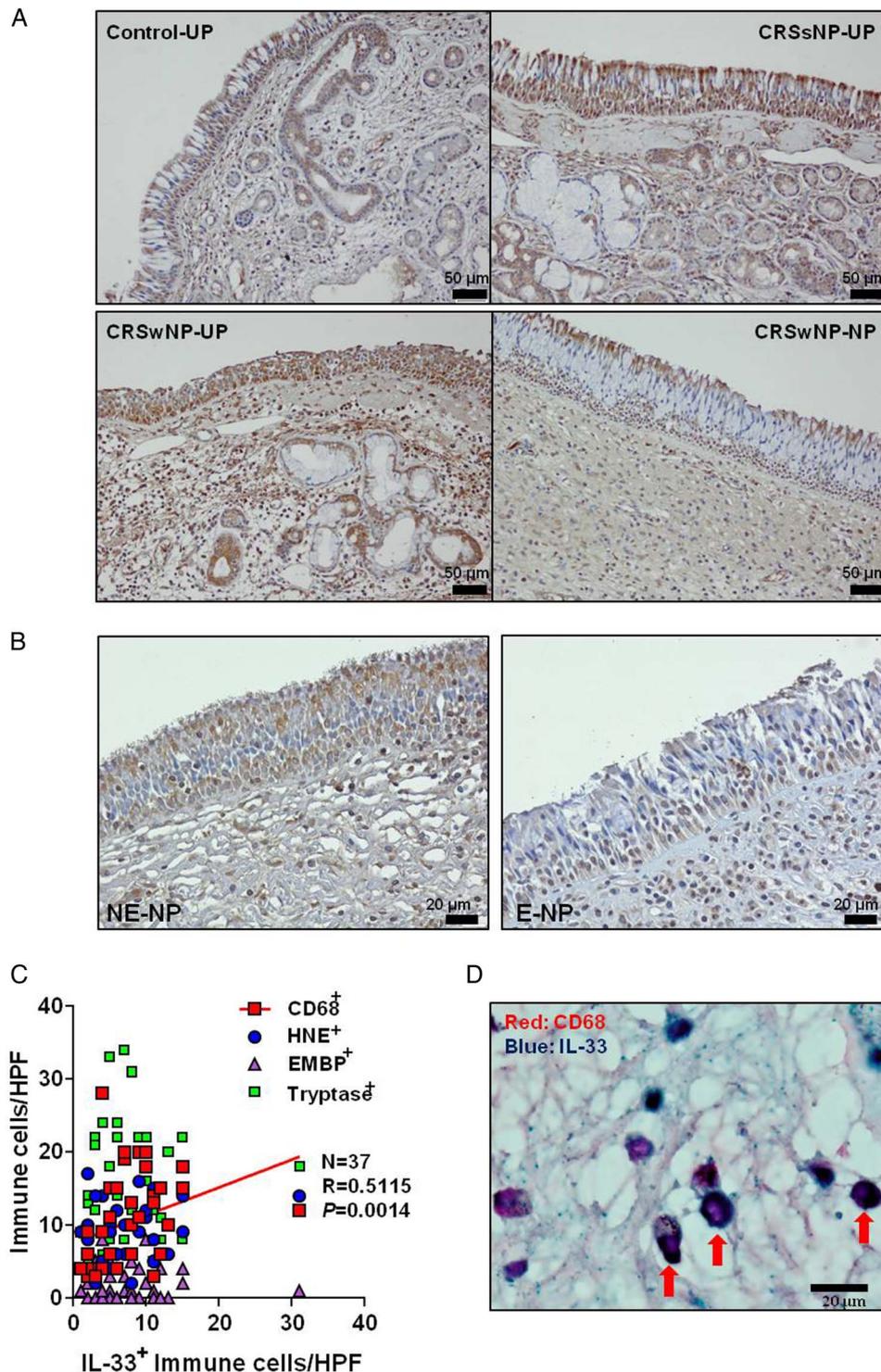


Figure 2 Expression of interleukin (IL-33) in different types of chronic rhinosinusitis and its cellular origin. Immunohistochemistry detection of IL-33 was performed. (A) Representative immunostaining for IL-33 within the nasal polyp (NP) tissues and uncinata process (UP) tissues from control subjects, patients with chronic rhinosinusitis without nasal polyps (CRSsNP) and patients with chronic rhinosinusitis with nasal polyps (CRSwNP). (B) Representative immunostaining for IL-33 within the NP tissues from non-eosinophilic CRSwNP (NE-NP) and eosinophilic CRSwNP (E-NP). (C) Double immunohistochemical stain for CD68, HNE, EMBP, tryptase, and IL-33 was performed and double-positive cells were counted. Immunostaining of representative cells with CD68 (red) and IL-33 (blue) in NP tissue. Arrows indicate double-positive immune cells ($\times 1000$). HPF, high power field.

(median: 0.760 ng/mg), UPs of CRSwNP (median: 0.917 ng/mg) and NPs (median: 0.772 ng/mg) than in those of UPs (medians: 0.187 ng/mg) in the controls (figure 1A). Interestingly, IL-33 protein levels were significantly increased in non-eosinophilic NP tissues (median: 1.112 ng/mg) compared with those of eosinophilic NP (median: 0.616 ng/mg; figure 1B). We also found that non-eosinophilic NP showed higher IL-33 mRNA levels than eosinophilic NP (2.53-fold, figure 1C). On the paired comparison of IL-33 protein levels between UPs and NPs from the same subjects, there was no significant difference in non-eosinophilic CRSwNP, whereas eosinophilic CRSwNP showed decreased concentration of IL-33 protein in NPs compared with UPs (figure 1D, E). Moreover, the concentration of IL-33 protein was positively correlated with the number of HNE-positive cells (neutrophils, $r=0.5054$, $p=0.0194$) and negatively associated with the number of EMBP-positive cells (eosinophils, $r=-0.6395$, $p=0.004$) in NPs (figure 1F, G).

Our IHC data confirmed that UP tissues from CRSwNP and CRSwNP, and NP tissues had increased expression of IL-33 compared with UP tissues from controls (figure 2A and see

online supplementary figure S2) and the expression of IL-33 was higher in epithelial cells of non-eosinophilic NPs versus those of eosinophilic NPs (figure 2B). To identify the cellular source of IL-33 in the subepithelial layer, IHC staining was performed using CD68 as a marker for macrophage, HNE for neutrophil, EMBP for eosinophil, and tryptase for mast cell. We observed a significant correlation between the number of IL-33-positive and CD68-positive cells in NPs ($r=0.5115$, $p=0.0014$; figure 2C). IL-33-positive and CD68-positive cells in NPs were also observed on double IHC staining (figure 2D).

Correlations between IL-33 expression and inflammatory markers in CRSwNP

To elucidate the role of IL-33 in CRS, we first examined the relationship between IL-33 expression and various inflammatory markers in UPs and NPs from patients with CRSwNP. Protein levels of IFN- γ ($r=0.6015$, $p=0.0001$) and IL-1 β ($r=0.5022$, $p=0.0018$) in NP tissues and IFN- γ , IL-1 β , TNF- α , IL-17A and IL-22 levels in UP tissues were positively correlated with the concentrations of IL-33. However, these IL-33 levels were

Figure 3 Correlation between interleukin (IL)-33 expression and inflammatory markers in nasal tissues according to the different types of chronic rhinosinusitis. The correlation of protein expression between IL-33 and inflammatory markers: (A) interferon (IFN)- γ , (B) IL-1 β , (C) tumour necrosis factor (TNF)- α , (D) IL-17A, (E) IL-22 and (F) IL-5. NP, nasal polyp; UP, uncinat process.

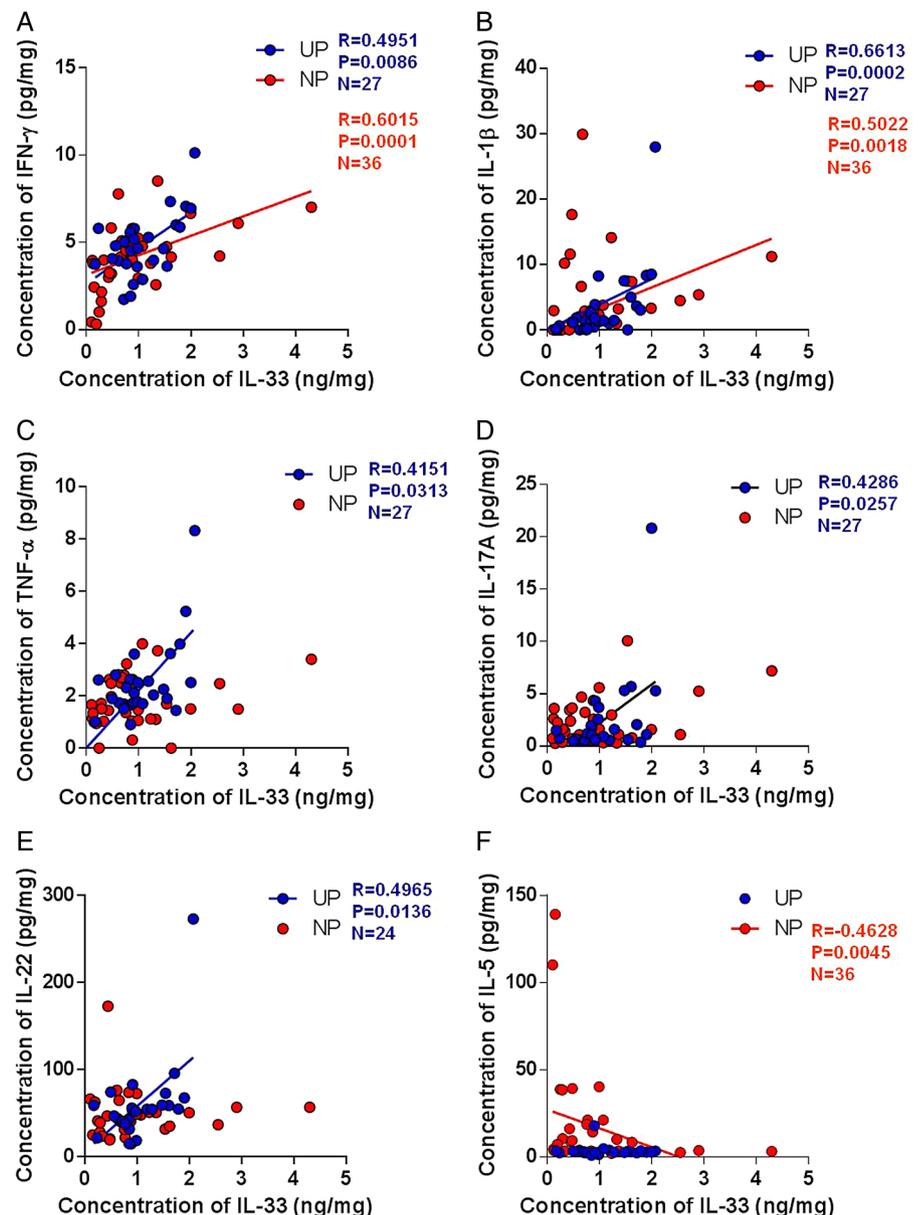
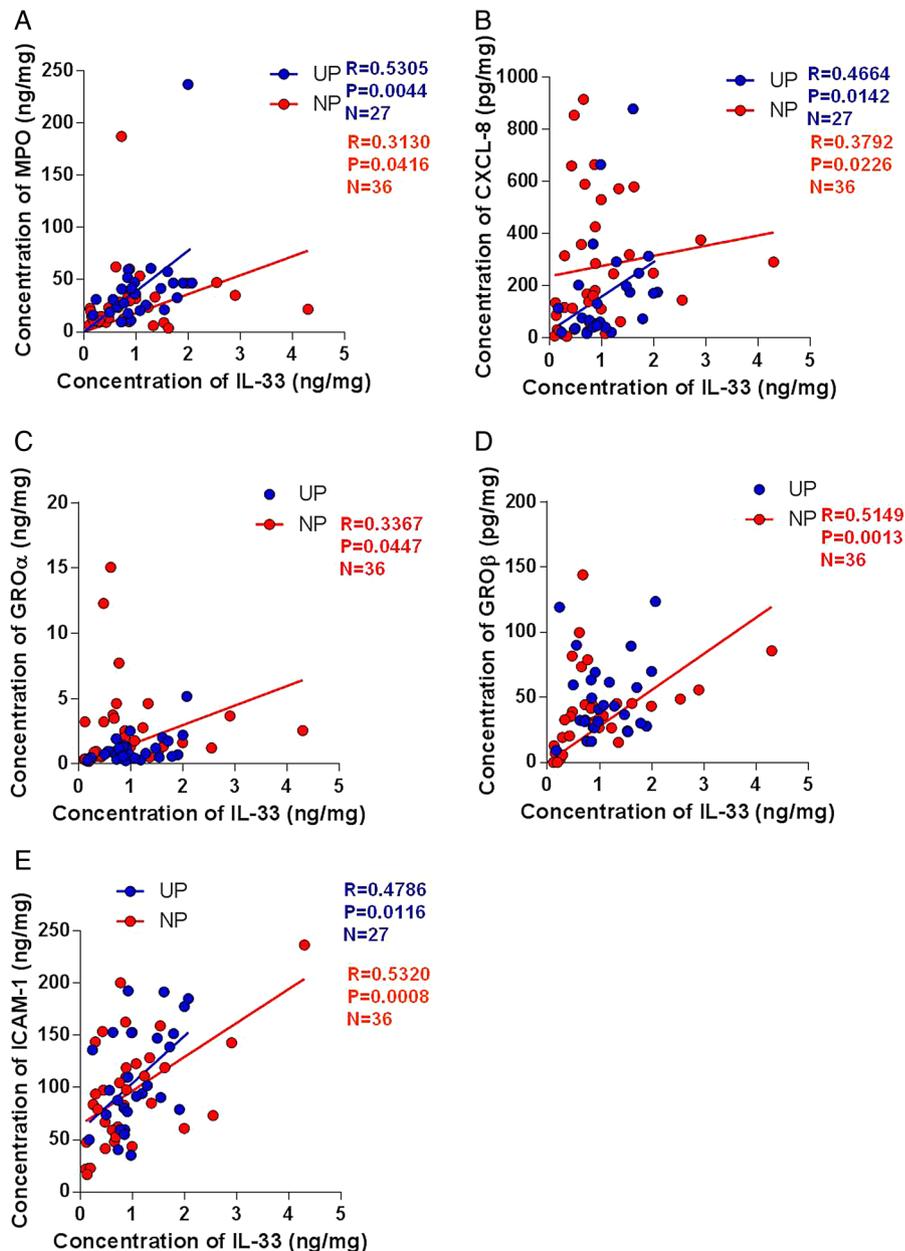


Figure 4 Correlation between interleukin (IL)-33 expression and recruitment markers for inflammatory cells in nasal tissues according to the different types of chronic rhinosinusitis. The correlation of protein expression between IL-33 and different markers: (A) MPO, (B) MCP4, (C) GRO α , (D) GRO β , (E) CXCL8 and (F) ICAM-1. NP, nasal polyp; UP, uncinat process.



negatively correlated with IL-5 ($r=-0.4628$, $p=0.0045$) protein levels in NP tissues (figure 3).

Second, to determine whether IL-33 was associated with recruitment of neutrophils, markers for human neutrophil (MPO), neutrophil-recruiting chemokines (CXCL-8, GRO- α , GRO- β), and recruitment adhesion molecules (ICAM-1) were assessed in UPs and NPs from patients with CRSwNP. On the analysis of NP homogenate samples, the protein levels of MPO ($r=0.3130$, $p=0.0416$), CXCL-8 ($r=0.3792$, $p=0.0226$), GRO- α ($r=0.3367$, $p=0.0447$), GRO- β ($r=0.5149$, $p=0.0013$) and ICAM-1 ($r=0.5320$, $p=0.0008$) were positively correlated with the concentrations of IL-33 protein (figure 4A–E). Moreover, in UP tissues from CRSwNP, the protein concentrations of IL-33 had a significant positive relationship with MPO, CXCL-8 and ICAM-1, respectively (figure 4A, B, E).

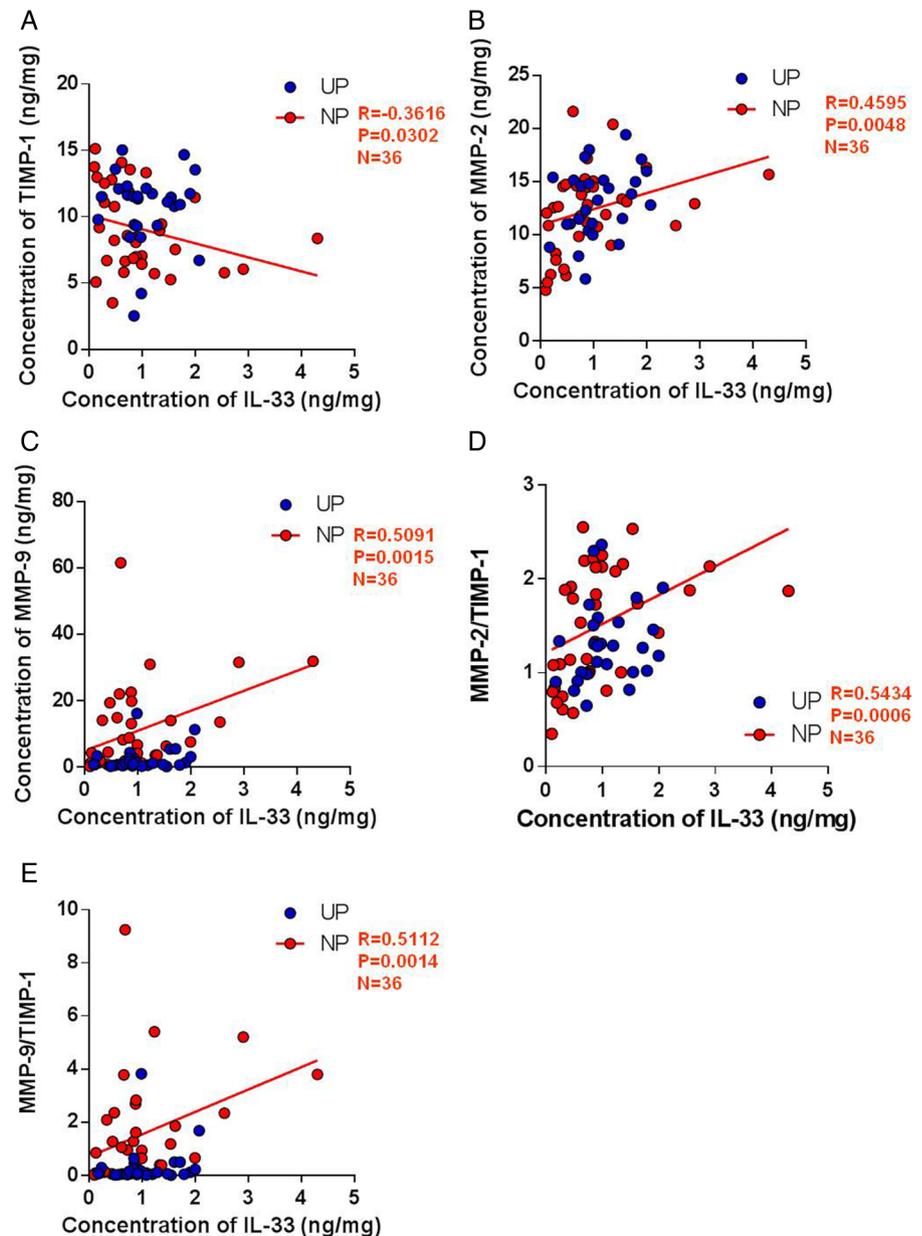
In this study, MMP-1 protein levels were significantly higher in tissue homogenates of eosinophilic NPs than those of controls, whereas increased production of MMP-2, MMP-9 and reduced levels of TIMP-1 were observed in tissue homogenates of non-eosinophilic NPs compared with controls (see online

supplementary figure S3). Interestingly, we found that IL-33 protein levels were correlated with remodelling markers such as TIMP-1, MMP-2 and MMP-9 in NP tissues (figure 5A–C). The ratios of MMP-2/TIMP-1 and MMP-9/TIMP-1 were positively correlated with the concentrations of IL-33 protein in NP tissues (figure 5D, E).

Anti-polyp effect of IL-33 neutralising antibody in a murine model of CRS

To investigate the therapeutic potential of blocking the effects of IL-33 in CRSwNP, we used a murine model of CRS (figure 6A). IHC staining was performed to examine the expression and localisation of IL-33 in this model before examining the effects of treatment with an anti-mouse IL-33 antibody. This murine model showed higher expression of IL-33 in the epithelial layer and in inflammatory cells compared with the control mice (figure 6B, C). In addition, the cytokine profiles of this murine model showed a mixed type of inflammatory response (see online supplementary figure S4) and were consistent with those of CRSwNP in Asian patients.^{6–8} The number of polypoid lesions,

Figure 5 Correlation between interleukin (IL)-33 expression and remodelling markers in nasal tissues according to the different types of chronic rhinosinusitis. The correlation of protein expression between IL-33 and different markers: (A) tissue inhibitor of metalloproteinase (TIMP)-1, (B) matrix metalloproteinase (MMP)-2, (C) MMP-9, (D) MMP-2/TIMP-1 and (E) MMP-9/TIMP-1. NP, nasal polyp; UP, uncinate process.



oedematous mucosal thickness, thickness of the subepithelial collagen layer, and number of goblet cells were apparently more severe in CRS mice than in control mice. Meanwhile, treatment with anti-IL-33 antibody (CRS+ α IL-33 group) significantly reduced those findings in the CRS model (figure 7A–E and see online supplementary figure 5). Moreover, cytokine profiles of nasal lavage fluids from the anti-IL-33-treated mice were in accordance with histological findings (figure 7F–I). After treatment with anti-IL-33 antibody, protein levels of IFN- γ , IL-17A and TGF- β 1 were significantly decreased compared with those of CRS mice (without anti-IL-33). However, there were no statistical differences in IL-4 protein levels between CRS and CRS+ α IL-33 groups.

Changes in neutrophil recruitment after IL-33 inhibition

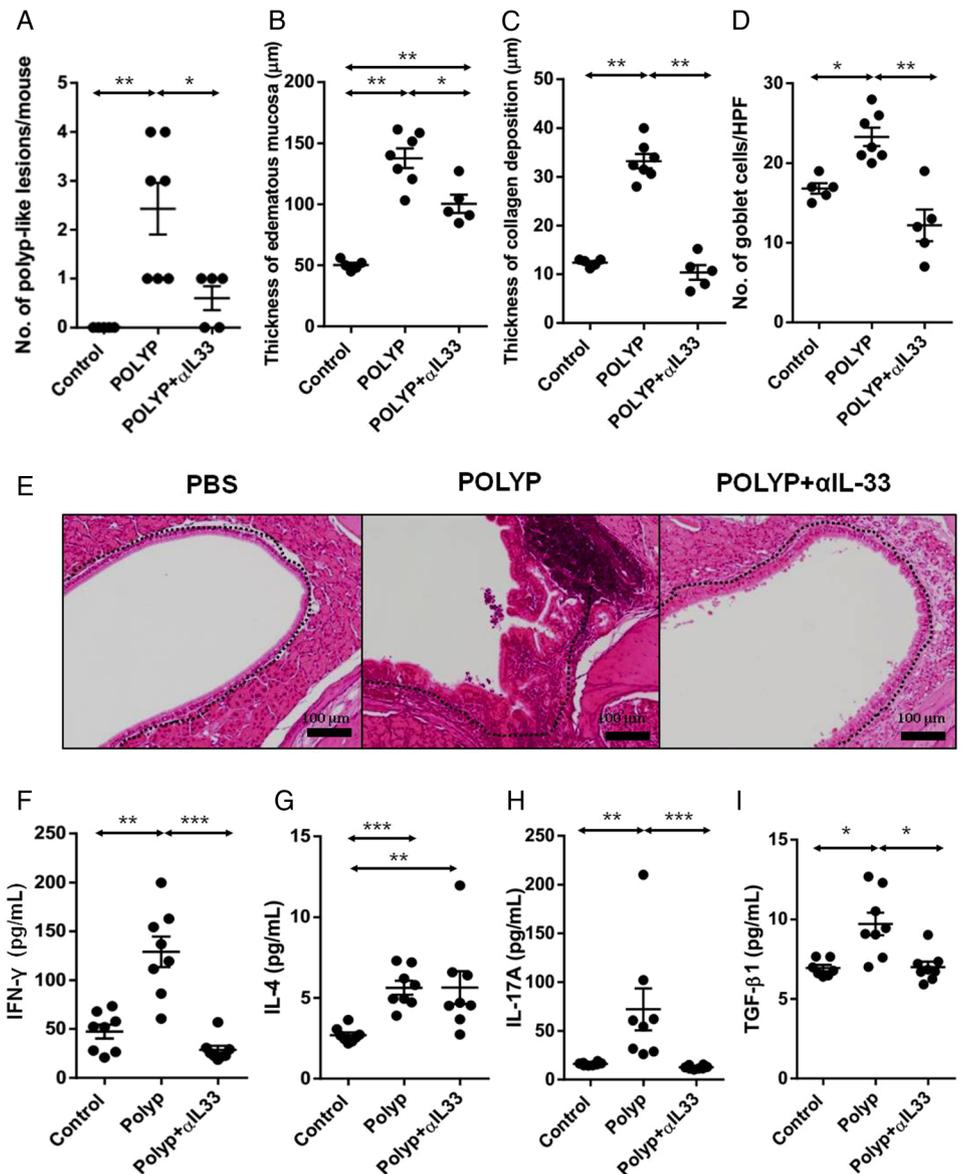
To verify the change in cytokine profiles after IL-33 inhibition, we investigated the infiltration of inflammatory cells. The degree of neutrophilic infiltration in the CRS+ α IL-33 group was decreased compared with that in the CRS group (figure 8A and

see online supplementary figure 5A), whereas anti-IL-33 treatment had no inhibitory effect on the number of eosinophils in these mice (figure 8B and see online supplementary figure S6B). To verify that inflammatory cell recruitment was inhibited by anti-IL-33 treatment, semi-quantitative RT-PCR was used to assess the levels of recruitment adhesion molecules (ICAM-1 and vascular endothelial adhesion molecule (VCAM)-1), eosinophil chemotactic cytokines (CCL-11 and CCL-24), and neutrophil-recruiting chemokines (CXCL-1 and CXCL-2) in this experimental model (figure 8C–H). These adhesion molecules and chemokines were unregulated in CRS mice compared with controls, while anti-IL-33 treatment led to downregulation of ICAM-1, VCAM-1 and CXCL2 (figure 8C, D, H). However, there were no significant differences in CCL-11 and CCL-24 levels with or without anti-IL-33 treatment.

DISCUSSION

IL-33 is a nuclear cytokine from the IL-1 family constitutively expressed in airway epithelial cells.⁶ Although IL-33 is

Figure 7 Effect of anti-interleukin (IL)-33 on nasal inflammation in the murine model. (A) Number of polypoid lesions, (B) oedematous mucosal thickness, (C) subepithelial collagen deposition, and (D) the number of goblet cells were determined in 10 different high-power fields (HPF; $\times 400$) and compared among the groups ($n=5$ for control; $n=7$ for chronic rhinosinusitis (CRS); and $n=5$ for CRS + α IL-33). (E–H) Cytokine profiles from nasal lavage fluids were also compared among groups ($n=8$ for each group). (I) Representative photograph for polypoid lesion in the control group (phosphate-buffered saline (PBS)), CRS model (POLYP), and CRS treated with anti-IL-33 (POLYP+ α IL-33). We performed the murine model three times (* $p<0.05$, ** $p<0.01$ and *** $p<0.001$). IFN, interferon; TGF, transforming growth factor.



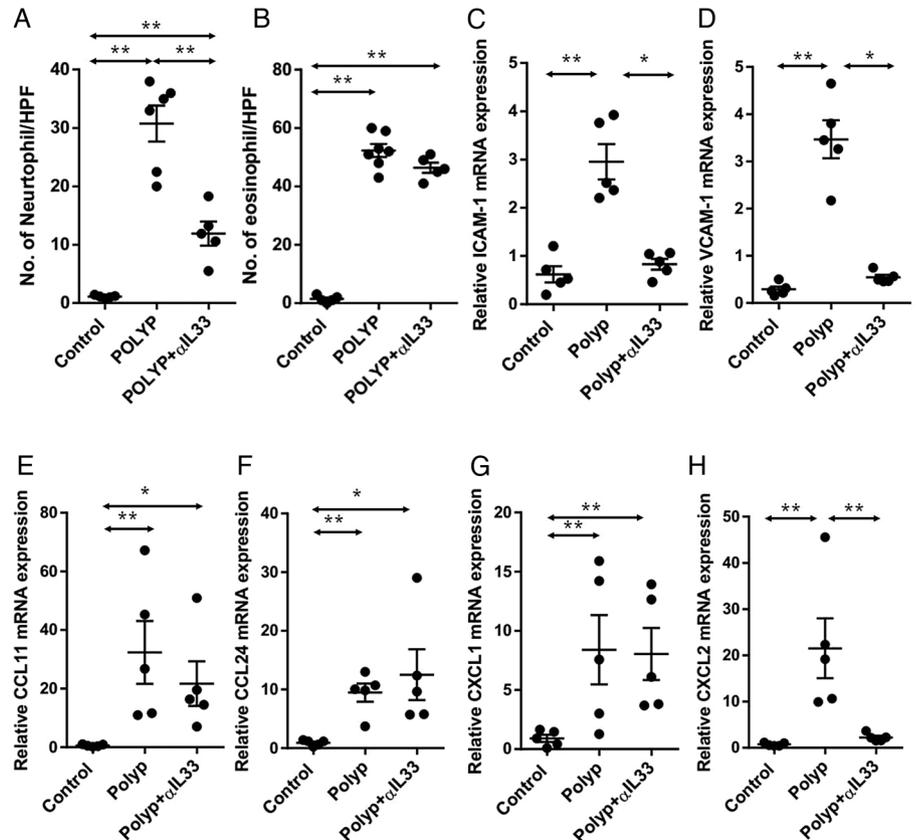
from CRSwNP. Other studies have reported a similar result regarding the association between IL-33 and neutrophils,^{8, 9} implying that IL-33 may enhance neutrophil recruitment to sites of infection. The number of neutrophils in an allergen-induced murine model of allergic rhinitis and asthma was significantly decreased after anti-IL-33 treatment,^{32, 33} whereas in a murine model for arthritis, neutrophil migration to the knee was induced by IL-33.¹³ Consistent with these studies, our results demonstrated that IL-33 was correlated with neutrophil recruitment.

The combination of MMPs, extracellular matrix (ECM) degrading enzymes, and TIMPs contributes to oedema formation in CRSwNP. During nasal polypogenesis, the balance between these proteases and protease inhibitors is important for degradation of ECM.^{34, 35} In this study, the concentrations of MMP-2 and MMP-9 protein were positively correlated with IL-33 levels in NPs. In addition, we found a negative correlation between TIMP-1 and IL-33 in NPs. Interestingly; a recent *in vivo* study revealed that MMP-2 and MMP-9 may promote neutrophil migration,³⁶ which is consistent with our results.

We also utilised the murine model of CRS,²⁵ and this model has been used to evaluate epithelial remodelling, the

therapeutic benefits of certain treatment, and various immunological host characteristics.^{26–28} In this model, polypoid lesions are defined as distinct mucosal bulges with inflammatory cell infiltration with underlying microcavity formation. This animal model showed mixed inflammation (see online supplementary figure S2) involving both eosinophilic and neutrophilic activity, which is consistent with NPs with mixed inflammation. Thus, the animal model using ovalbumin (OVA)/staphylococcal enterotoxin B (SEB) was well suited to determine whether anti-IL-33 antibody exerted a therapeutic effect on chronic inflammation and NP formation. While mice treated with anti-IL-33 showed reduced eosinophilic inflammation in OVA-induced acute allergic airway diseases,^{32, 33} the effect of IL-33 blockade on chronic sinonasal inflammation remains unknown. In this study, we found increased production of IL-33 in epithelial and inflammatory cells in OVA/SEB-induced CRS mice. These findings suggest that IL-33 may contribute to OVA/SEB-induced sinonasal inflammation. In addition, we found that the degree of average mucosal thickness was diminished by administration of anti-IL-33. Similarly, the CRS + α IL-33 group showed a significant decrease in the thickness of the subepithelial collagen and in the number of goblet cells

Figure 8 Anti-IL-33 therapy suppresses neutrophil infiltration, expression of neutrophil chemotactic cytokines and intercellular adhesion molecules in a murine model. (A) The numbers of infiltrated neutrophils and (B) infiltrated eosinophils were counted from 10 different high-power fields (HPF; $\times 400$) and compared among groups ($n=5$ for control, $n=7$ for chronic rhinosinusitis (CRS), and $n=5$ for CRS+ α IL-33). Relative mRNA expression levels of the intercellular and vascular adhesion molecules (C) ICAM-1 and (D) VCAM-1, the eosinophil-recruiting chemokines (E) CCL11 and (F) CCL24, and the neutrophil-recruiting chemokines (G) CXCL1 and (H) CXCL2 were compared among groups ($n=5$ for all groups). We performed the murine model three times. (* $p<0.05$, ** $p<0.01$ and *** $p<0.001$).



compared with the CRS group. Interestingly, in our animal study, we observed that the infiltration of neutrophils was decreased in the CRS+ α IL-33 group compared with the CRS group. However, no significant differences in eosinophilic infiltration were observed between these groups. Moreover, we demonstrated decreased expression of cytokines (IFN- γ /IL-17A/TGF- β 1), adhesion molecules (ICAM-1/VCAM-1), and neutrophil-recruiting chemokine (CXCL2) with anti-IL-33 treatment. These cytokines, adhesion molecules and chemokines are closely related to neutrophil recruitment. Considering these results, we speculate that the anti-inflammatory effect of anti-IL-33 antibody may stem from suppression of neutrophil recruitment. However, the exact subset(s) of IL-33-responsive innate immune cells involved in reducing neutrophil recruitment should be further investigated.

In conclusion, we found that UP or non-eosinophilic NP tissues of CRSwNP were characterised by elevated expression of IL-33, and these levels correlated with levels of Th1/Th17 cytokines. Additionally, IL-33 protein levels were associated with various markers for neutrophil recruitment and remodelling in NP tissues. Moreover, anti-IL-33 antibody exerted an inhibitory effect on chronic sinonasal inflammation by reducing the recruitment of neutrophils in a murine model of CRS. These findings suggest that IL-33 plays a crucial role in the pathogenesis of neutrophilic inflammation in patients with CRS and that IL-33 inhibition may provide a new treatment strategy for CRS, particularly in non-eosinophilic NPs.

Contributors Study supervision: DWK. Critical revision of the manuscript for important intellectual content: D-KK, HRJ and KME. Conceived and designed the experiments: D-KK and DWK. Performed the experiments and analysed the data:

D-KK, J-HM, SHC, SO and DWK. Contributed reagents/materials/analysis tools and wrote the paper: D-KK, DC and DWK.

Funding This research was supported by a clinical research grant-in-aid from the Seoul Metropolitan Government Seoul National University (SMG-SNU) Boramae Medical Center (03-2013-7), by SK Telecom Research Fund (01-2014-10).

Competing interests None declared.

Patient consent Obtained.

Ethics approval This study was approved by the institutional review board of Seoul National University (Boramae Medical Center).

Provenance and peer review Not commissioned; externally peer reviewed.

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