

A Recently Established Murine Model of Nasal Polyps Demonstrates Activation of B Cells, as Occurs in Human Nasal Polyps

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Abstract

Animal model systems are invaluable for examining human diseases. Our laboratory recently established a mouse model of nasal polyps (NPs) and investigated similarities and differences between this mouse model and human NPs. We especially focus on the hypothesis that B cell activation occurs during NP generation in the murine model. After induction of ovalbumin-induced allergic rhinosinusitis, 6% ovalbumin and *Staphylococcus aureus* enterotoxin B (10 ng) were instilled into the nasal cavity of mice three times per week for 8 weeks. The development of structures that somewhat resemble NPs (which we will refer to as NPs) was confirmed by hematoxylin and eosin staining. The mRNA and protein levels of various inflammatory cell markers and mediators were measured by real-time PCR in nasal tissue and by ELISA in nasal lavage fluid (NLF), respectively. Total Ig isotype levels in NLF were also quantitated using the Mouse Ig Isotyping Multiplex kit (EMD Millipore, Billerica, MA) on a Luminex 200 instrument (Life Technologies, Grand Island, NY). Similar to human NPs, there were significant increases in gene expression of inflammatory cell markers, such as CD19, CD138, CD11c, and mast cell protease-6 in nasal tissue samples of the NP group

compared with those of the control group. In further investigations of B cell activation, mRNA expressions of B cell activating factor and a proliferation-inducing ligand were found to be significantly increased in mouse NP tissue. B cell-activating factor protein concentration and IgA and IgG₁ levels in NLF were significantly higher in the NP group compared with the control group. In this study, the NP mouse model demonstrated enhanced B cell responses, which are reminiscent of B cell responses in human NPs.

Keywords: animal model; antibodies; B cells; chronic rhinosinusitis; nasal polyps

Clinical Relevance

The present study demonstrates that the nasal polyp (NP) mouse model revealed enhanced B cell responses, reminiscent of human NPs. This mouse model may enhance our understanding of the pathophysiology of NPs and provide a model to test therapeutic targets *in vivo*.

Chronic rhinosinusitis (CRS) is one of most common chronic diseases worldwide. CRS can be categorized into two types: CRS with nasal polyps (NPs; CRSwNPs) and CRS

without NPs (CRSsNPs). The pathogenesis of NPs is incompletely understood, and lack of an *in vivo* animal model for NPs has been a major hurdle in investigating NP

pathogenesis and testing new treatment modalities. Animal model systems are invaluable *in vivo* models for examining a variety of human diseases and aid in the

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development of new therapeutic targets. Our laboratory recently established a mouse model of NPs using a modified protocol generated by Kim and colleagues (1), and investigated similarities and differences between this mouse model and human NPs.

There have been several reports that local activation of B cells and production of antibodies are associated with pathogenic mechanisms in airway diseases (2–7). In addition, our group has previously shown that B cell-activating factor (BAFF) of the TNF family, a key B cell survival factor, is highly expressed in NP tissue from patients with CRSwNPs (8). We also reported increased levels of autoantigen-specific antibodies in NP tissue (9, 10). Several reports have also demonstrated elevated levels of various isotypes of Igs, including IgG, IgE, and IgA, in sinus tissue from patients with CRS (11, 12). Taken together, these findings suggest that B cell activation and antibody production may play important roles in the pathogenesis of CRSwNPs. We therefore focused this investigation on B cell activation in the mouse NP model.

Materials and Methods

Generation of the Mouse Model of NPs

Mice were divided into three groups and each group consisted of 10 mice (control group, ovalbumin [OVA] group, NP group; Figure 1). The procedure for the mouse CRS model is summarized in Figure 1. Briefly, 4-week-old female BALB/c mice were systemically sensitized with an intraperitoneal injection of PBS or 25 µg of OVA (grade V; Sigma, St. Louis, MO) plus 2 mg of aluminum hydroxide gel on Days 0 and 7. After general sensitization, mice were nasally challenged with PBS or 6% OVA daily from Day 14 to Day 20. For the NP group, 6% OVA with *Staphylococcus aureus* enterotoxin B (SEB; 10 ng) was instilled into the nasal cavity of mice three times per week for 8 weeks after induction of OVA-induced allergic rhinosinusitis. Development of NP-like tissue was confirmed by staining with hematoxylin and eosin (H&E). Negative control mice received neither OVA nor SEB. The OVA group of mice were challenged intranasally with only 6% OVA without SEB. The whole experiment was repeated three times. All

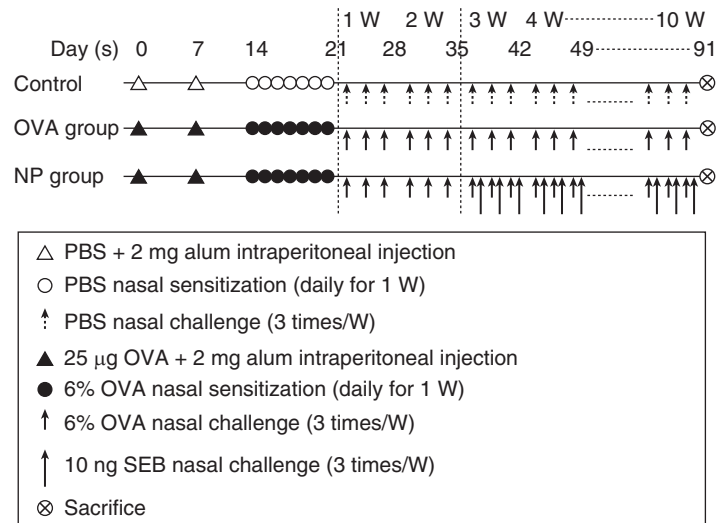


Figure 1. Experimental protocol. BALB/c mice were sensitized with PBS or ovalbumin (OVA) plus 2 mg of aluminum hydroxide gel (Alum) on Days 0 and 7 (general sensitization) and then received intranasal PBS or OVA from Day 14 to Day 20. For the nasal polyp (NP) group, 6% OVA with *Staphylococcus aureus* enterotoxin B (SEB; 10 ng) was instilled into the nasal cavity of mice three times per week for 8 weeks after 2 weeks of the OVA-alone challenge.

animal experiments conducted in this study followed the guidelines and ethics of the Institutional Animal Care and Use Committee at Northwestern University (Chicago, IL).

Tissue Preparation for H&E Staining, Real-Time RT-PCR, and Nasal Lavage Fluid Collection

For the preparation of nasal cavity samples, we decapitated mice and then, with their heads immobilized, removed the lower jaw together with the tongue. Using the hard palate as a guide, we then used a large scalpel to remove the snout with a transverse cut behind the back molars. After removing the skin and any excess soft tissue, we flushed the external nares with PBS to wash out any blood within the nasal cavity.

For H&E staining, nasal tissues were fixed in 4% paraformaldehyde, decalcified, embedded in paraffin, and sectioned coronally (4 µm thick) approximately 5 mm from the nasal vestibule.

For real-time RT-PCR, the nasal mucosa was taken out meticulously using a small curette after bisecting the nasal tissue sagittally along the nasal septum. Nasal mucosa was immediately immersed in liquid nitrogen and stored at -70°C until use for real-time RT-PCR.

To collect nasal lavage fluid (NLF), 200 µl of PBS was flushed through the nasal

cavity from the posterior choanae to the anterior nostrils using a pipette tip after the lower jaw was resected. This was performed twice, and approximately 400 µl of NLF was collected.

Measurement of mRNAs in Nasal Tissue, and BAFF Protein and Total Ig Isotype Levels in NLF

The mRNA levels of various cell markers and inflammatory mediators were measured by real-time RT-PCR in nasal tissue (8). Total RNA was prepared from the nasal mucosa with TriZol reagent (Invitrogen, Carlsbad, CA). cDNA was synthesized using Superscript reverse transcriptase (Invitrogen) and oligo(dT) primers (Fermentas, Burlington, ON, Canada). For analysis of CD19 (Mm00515420_m1), CD138 (Mm00448918_m1), CD11c (Mm00498701_m1), mast cell protease-6 (MCP-6) (Mm01301240_g1), BAFF (Mm00446347_m1), a proliferation-inducing ligand (APRIL; Mm03809849_s1), and glyceraldehyde 3-phosphate dehydrogenase (GAPDH; Mm99999915_g1), and Pre-Developed Assay Reagent kits of primers and probes were purchased from Applied Biosystems (Foster City, CA). Amplification of CD19, CD138, CD11c, MCP-6, BAFF, APRIL, and GAPDH cDNA was performed in MicroAmp optical

96-well reaction plates (Applied Biosystems). The reaction was performed using an ABI PRISM 7,000 Sequence Detection System (Applied Biosystems). The average transcript levels of genes were then normalized to GAPDH.

The protein level of BAFF was quantitated by ELISA in NLF. Total Ig isotype levels in NLF were also quantitated using the Mouse Ig Isotyping Multiplex kit (EMD Millipore, Billerica, MA) on a Luminex 200 instrument (Life Technologies, Grand Island, NY), and values were normalized to total protein.

Results

Histologic Confirmation of NP Generation

The H&E staining (Figure 2) of nasal tissue revealed that mice challenged with OVA plus SEB (NP group) developed multiple edematous, polyp-like lesions with heavy eosinophilic infiltration, whereas mice challenged with only OVA (OVA group) showed eosinophilic infiltrations, but no polypoid lesions. We streamlined the original protocol of the mouse NP model (1) by changing the OVA-challenge period from 4 weeks to 2 weeks (Figure 1). Even after omitting 2 weeks of the OVA-challenge period, we still obtained a similar pattern of polypoid lesions by H&E staining (data not shown). This finding can contribute to shortening the duration of the NP generation.

Increased Expression of Inflammatory Cell Markers in CRS and the NP Model

As in human NPs, there were significant increases in expression of inflammatory

cell marker genes, such as CD19 (B cell marker, twofold; Figure 3A), CD138 (plasma cell marker, 3.5-fold; Figure 3B), CD11c (dendritic cell marker, ninefold; Figure 3C), and MCP-6 (mast cell marker, 250-fold; Figure 3D), in the nasal tissue samples from the NP group compared with those of the control group ($P < 0.05$). However, CD3e, a T cell marker, was not significantly different between the groups (data not shown). In the case of CD19, CD138, CD11c, and MCP-6, mRNA expression levels were also significantly increased in the OVA group compared with those seen in the control group (Figures 3A–3D). These results suggest that these markers are not specific to NPs, but rather indicators of allergic inflammation.

Increased Production of BAFF mRNA and Protein in the NP Model

Upon further analysis of B cell activation, mRNA expression levels of BAFF (threefold; Figure 3E) and APRIL (2.5-fold; Figure 3F) were found to be significantly increased in mouse NP tissue compared with those seen in control ($P < 0.05$). However, the OVA group did not show significant differences in mRNA expression of either BAFF or APRIL compared with control. Because we previously found that BAFF, but not APRIL, is significantly elevated in CRS and associated with elevation of Igs in the airways (8, 13), we measured BAFF protein concentration in NLF. BAFF levels were significantly higher in the NP group than those in the control group (1.5-fold, $P < 0.05$; Figure 3G).

Increased Levels of IgA and IgG₁ in the Mouse Model of NPs

Although B cells can play various roles in ongoing inflammation, an important

activity is the production and secretion of antibodies (14). We therefore evaluated Ig isotype levels in NLF using the Mouse Ig Isotyping Multiplex kit. In accordance with our previous results of increased IgA, IgG₁, and IgG₄ in human NPs (13), NLF from the NP mouse group contained significantly higher levels of IgA (Figure 4A) and IgG₁ (Figure 4B) than those seen in the control group ($P < 0.05$). IgG₁ levels were also significantly increased in the NP group compared with the OVA group ($P < 0.05$). On the other hand, IgG_{2a}, IgG_{2b}, and IgG₃ levels were significantly lower in the NP group compared with those in control group ($P < 0.05$; Figures 4C–4E). Regarding the inflammatory changes in mice, IgG₁ is typically associated with Th2 immune responses, whereas IgG_{2a}, IgG_{2b}, and IgG₃ are associated with Th1 immune responses (15).

Discussion

Because one of the most representative signs of B cell activation is enhanced production and secretion of antibodies, our group and others have previously evaluated antibody secretion in human NPs (13, 16). Increased levels of several specific autoantibodies were found in NP tissue compared with levels seen in tissues of control subjects and in patients with CRSwNPs. In addition, nuclear-targeted autoantibodies, such as anti-double-stranded DNA IgG and IgA antibodies, were found at significantly increased levels in NPs, particularly in NPs from patients requiring revision surgery for recurrence (9). In addition, we previously measured the levels of various antibody isotypes in NP and uncinatate tissue, as well as in serum. NPs had significantly higher levels of all antibody isotypes (except IgG3) compared with normal uncinatate tissue, accompanied by enriched infiltrates of B cells and plasma cells in NPs. In contrast, there were no differences in circulating antibody levels (13). Taken together, activated B cells and plasma cells in NPs vigorously produce antibodies locally, and may be involved in chronic inflammation in patients with CRSwNPs. Our mouse NP model appears to reproduce major elements of findings in the B cell compartment observed in human NPs, showing significantly increased levels of IgA and IgG₁ in NLF. Although we could not measure IgE level in NLF due to the

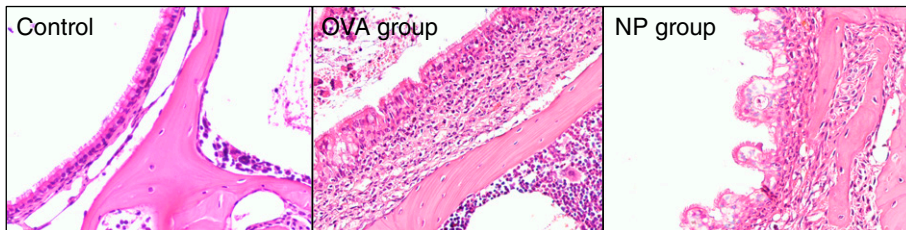


Figure 2. Histology of a mouse model of NPs. Representative hematoxylin and eosin staining images show that mice challenged with OVA plus SEB developed multiple edematous polypoid lesions with heavy eosinophilic infiltrations (NP group), whereas control mice and mice challenged with only OVA (OVA group) did not. Magnification, $\times 200$.

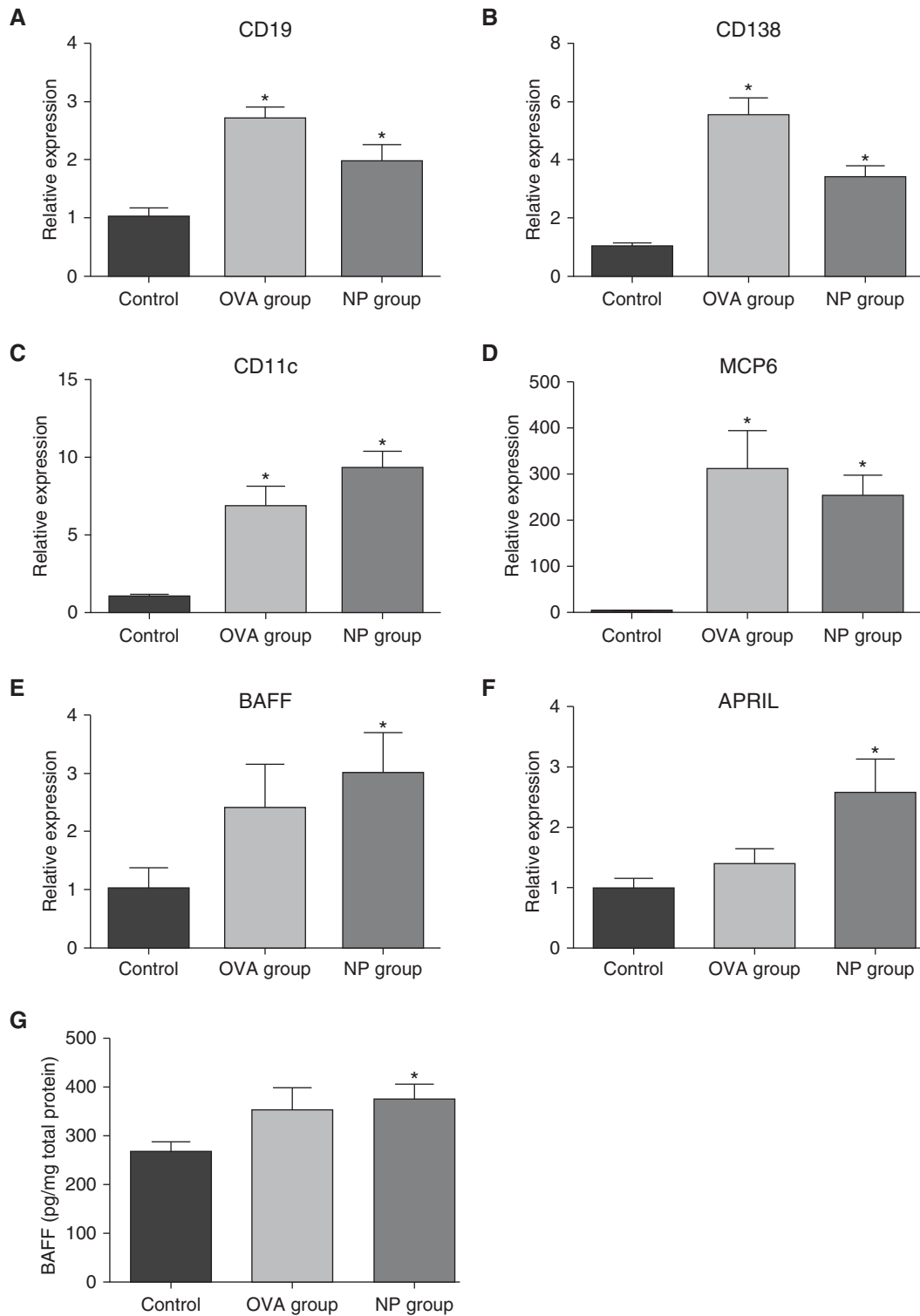


Figure 3. Expression of inflammatory and B cell markers in the NP group. Total RNA was extracted from nasal tissues, and expression of CD19 (B cell marker [A]), CD138 (plasma cell marker [B]), CD11c (dendritic cell marker [C]), and mast cell protease-6 (MCP6) (mast cell marker [D]) was analyzed using real-time PCR. In more analyses of B cell activation, mRNA expression of B cell-activating factor (BAFF) (E) and a proliferation-inducing ligand (APRIL) (F) was found to be significantly increased in mouse NP tissue. BAFF protein concentration in nasal lavage fluid (NLF) was also significantly higher in the NP group than in the control (G). Data presented are means \pm SEM. * P < 0.05 versus control.

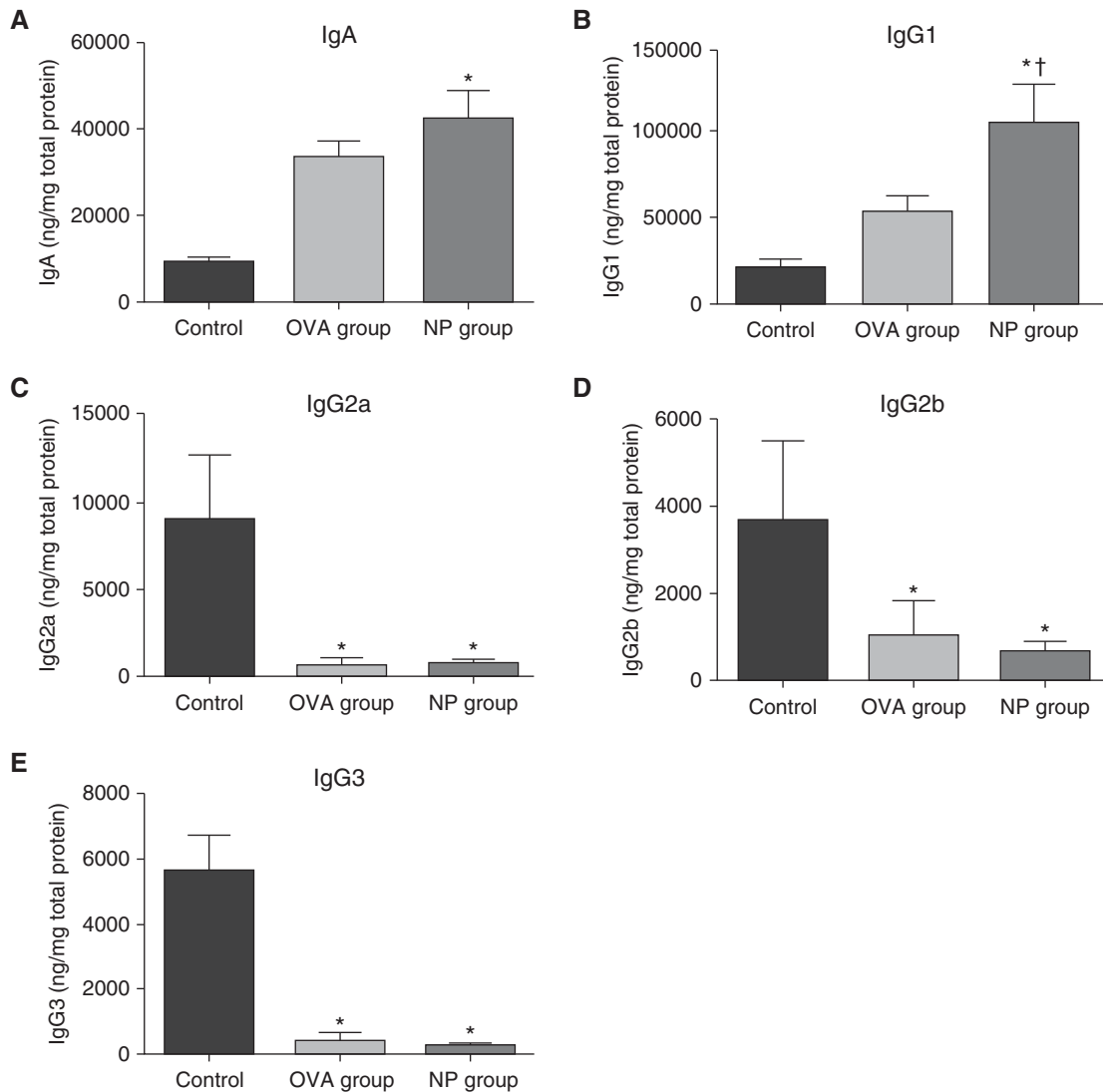


Figure 4. Antibody levels in NLF. Total Ig isotype assay by the Mouse Ig Isotyping Multiplex kit demonstrated that NLF of the NP group contained significantly increased levels of IgA (A) and IgG₁ (B) compared with those seen in control. IgG₁ level was also significantly higher in the NP group than that in the OVA group. The levels of IgG_{2a} (C), IgG_{2b} (D), and IgG₃ (E) were significantly lower in the NP group than those in control. Data presented are means ± SEM. **P* < 0.05 versus control; †*P* < 0.05 versus OVA group.

absence of the IgE item in the multiplex kit and the insufficient amount of NLF, the original report of the protocol of the mouse NP model (1) already described increased levels of IgE in NLF. Furthermore, a number of studies have confirmed a significantly higher local concentration of IgE in human NP than control tissues (17–21).

This mouse NP model has some limitations. First, as it is based on allergic inflammation with eosinophilic infiltration, which is biased to Th2 responses, it might better reflect mechanisms of formation of Western polyyps, but not Asian polyyps,

which are often not eosinophilic (22). Second, the morphology of mouse NPs is somewhat different from that of human NPs, which often become quite large and expand to fill the nasal cavity. This could be due to a reduced distribution of respiratory epithelium in the nasal airways of the mouse compared with the human nasal airways, where the NP-like structures that we have observed originated. Third, the necessity of prolonged challenges to make the model is a practical obstacle for investigators, although our protocol has been simplified by the modification of the previous protocol. Finally, this mouse

model is driven by antigen, and requires the enterotoxin adjuvant to develop NPs, and is thus antigen dependent. Human nasal polyposis has not been definitively shown to be driven by antigens, although, as mentioned previously here, studies from many laboratories support a role of local Igs of various isotypes in CRS pathogenesis.

In summary, this study demonstrates novel findings of enhanced expression of B cell and plasma cell markers, and elevated levels of IgA and IgG₁ as a consequence of B cell activation in a new murine model of NPs. Up-regulation of BAFF and APRIL

may play an important role in the B cell activation that occurs during NP generation. Based upon this study and previous work in human NPs, testing of an anti-BAFF human monoclonal antibody

(belimumab) may be warranted in the treatment of NPs. This animal model of CRS and NPs can be used to explore pathogenic mechanisms of the disease, as well as to test potential therapeutic

targets for the pharmacologic interventions. ■

Author disclosures are available with the text of this article at www.atsjournals.org.

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