Peripheral Eosinophil Counts Correlate With Nasal Eosinophil Counts in Patients With Rhinitis

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Eosinophilic inflammation affecting the nose indicates a T(H)2 immune response, which is typical in allergic rhinitis and in nonallergic rhinitis with eosinophils (NARES), as well as in eosinophilic asthma [1]. Nasal cytology is a convenient method that is very useful in clinical practice, mainly in the diagnostic and prognostic work-up of patients with rhinitis [2]. In addition, it has been reported that the nasal eosinophil count correlates better with symptom severity and IgE level [3,4].

In their cross-sectional study of adults with moderate-severe asthma, Amorim et al [5] demonstrated a convincing association between nasal and sputum eosinophilia and a link between the former and bronchodilator response, ie, postsalbutamol FEV₁. These results agree with those of recent studies that showed close similarities in tissue inflammatory changes in asthma and rhinitis, further supporting the concept that the upper and the lower airways should be considered a single entity influenced by common physiologic processes, namely, the one-airway hypothesis [6]. Therefore, the evaluation of upper airway inflammation may provide additional insight into lower airway involvement and suggests that evaluation of nasal eosinophilia could be a surrogate for sputum analysis in these patients. In other words, nasal eosinophils may mirror bronchial eosinophils, thus enabling the nose to be considered the window of the bronchi.

Another pathway for indirect evaluation of bronchial eosinophils is through blood eosinophils. Peripheral eosinophils have been reported to be a reliable surrogate biomarker for phenotyping type 2 asthma [7].

Therefore, we tested the hypothesis of whether peripheral eosinophil count is correlated with nasal eosinophil counts. To verify this possibility, we compared nasal eosinophils with blood eosinophils in a group of patients with rhinitis in a real-world setting. The study sample comprised 41 consecutive patients (23 males, 18 females; mean age, 38.7 years) attending a rhinology clinic who were enrolled on 2 consecutive days. All patients underwent a through otorhinolaryngologic examination (including endoscopy, nasal scraping, and nasal...
lavage) and had inflammatory rhinitis (15 allergic rhinitis, 10 NARES, and 11 nonallergic noneosinophilic rhinitis). Diagnosis of allergic rhinitis was based on the agreement between a positive result in the skin prick test and the occurrence of nasal symptoms after exposure to the sensitizing allergen. The diagnosis of NARES was based on nasal eosinophilia (≥5% of cell count) and a negative allergy test result. The diagnosis of nonallergic noneosinophilic rhinitis was based on the presence of inflammatory cell infiltrate in the absence of allergy and a diagnosis of NARES.

Nasal scraping for traditional cytology was performed according to validated criteria [2]. Nasal lavage was performed by slowly instilling 10 mL of sterile isotonic saline into each nostril using a 10-mL syringe while the patient inclined his/her head and closed the soft palate. The solution was retained for approximately 10 seconds in the nasal cavities without swallowing. It was then expelled by flexing the head forward, exhaling lightly, and rinsing lavage fluid into a sterile plastic beaker. Patients were strictly instructed to collect only secretions from the nose in the sterile beaker; secretions from the mouth had to be spat into the toilet. Immediately after collection, nasal lavage fluid was cytocentrifuged and the cell pellet was suspended in flow cytometry buffer (PBS, 0.09% sodium azide, 1% heat-inactivated FBS) and stained with monoclonal antibodies to CD294 and CD203c (CD294 positivity and CD203c negativity identify eosinophils) for 20 minutes at room temperature in the dark. Cells were washed with flow cytometry buffer, resuspended in 0.5% paraformaldehyde, and stored at 4°C in the dark. Samples were acquired within 24 hours on a flow cytometer (Cytomics FC 500, Beckman Coulter Diagnostics). Isotype-matched single color controls were used to control for nonspecific staining and to set analysis gates.

As shown in the Figure, A, the relationship between peripheral eosinophils and nasal eosinophils recovered by nasal scraping (mean numbers of 10 fields; original magnification, ×1000) and identified by microscopy, B, Relationship between peripheral eosinophils and CD294-CD203c– cells recovered by nasal lavage and identified by flow cytometry.

Figure. A, Relationship between peripheral eosinophils and nasal eosinophils recovered by nasal scraping (mean numbers of 10 fields; original magnification, ×1000) and identified by microscopy. B, Relationship between peripheral eosinophils and CD294-CD203c– cells recovered by nasal lavage and identified by flow cytometry.
In conclusion, peripheral eosinophil count correlates well with nasal eosinophil count and could reasonably be considered a biomarker for suspecting type 2 inflammation in the nose.

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**Conflicts of Interest**

The authors declare that they have no conflicts of interest.

**References**