



Current perspectives on airway inflammation and remodelling in asthma and allergic rhinitis

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ABSTRACT

The development of AR and asthma requires an interaction between the environment, immune system and genetic susceptibility. While pollen-induced rhinitis is the most characteristic IgE-mediated allergic disease, in perennial allergic rhinitis the allergic triggers are more continuous, and lead to ongoing inflammation. Several cells and mediators orchestrate and maintain this inflammation. Although histamine is still one of the major mediators of the allergic reaction, many other mediators produced by different cell types are involved. Thus, the intricate interaction amongst these mediators, cytokines, chemokines, neuropeptides, adhesion molecules and various cells in the form of a complex network leads to the development of specific symptoms and the non specific hyperreactivity of allergic rhinitis. Asthma is characterized by variable degrees of chronic inflammation and structural alterations in the airways which include epithelial denudation, goblet cell metaplasia, subepithelial thickening, increased airway smooth muscle mass, bronchial gland enlargement, angiogenesis, and alterations in extracellular matrix components, involving large and small airways. Chronic inflammation is thought to initiate and perpetuate cycles of tissue injury and repair in asthma, although remodeling may also occur in parallel with inflammation. While AR and asthma share several similarities in the inflammatory cell and mediator profiles and responses, remodeling as seen in asthma is not characteristic of AR. In asthma, the relationships of airway inflammation, remodeling and lung function are becoming better understood. A variety of inflammatory cells and structural cells play a role in orchestrating the inflammation and structural changes in asthma. Increased airway responsiveness is a surrogate marker of inflammation and may reflect the development of structural changes in the airways. Such persistent increased bronchial responsiveness indicates remodeling which is partly resistant to therapy.

Keywords: Asthma, allergic rhinitis, airway remodeling, airway inflammation, cytokines.

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INTRODUCTION

Allergic diseases like allergic rhinitis (AR) and asthma are increasing to epidemic proportions worldwide and are major public health issues in both the developed and developing countries. Considering the focus of global bodies like the United Nations on non-communicable diseases and efforts to control these diseases, allergic diseases are amongst the earliest manifestations in early childhood of non-communicable diseases. While AR is an IgE-mediated chronic inflammation of the

nasal mucosa characterized by symptoms of sneezing, rhinorrhea, nasal itching and nasal obstruction, asthma is a chronic, inflammatory condition of the lower airways characterized by largely reversible airflow obstruction, airway hyperresponsiveness, and episodic respiratory symptoms, including wheezing, productive cough, and sensations of breathlessness and chest tightness.¹

Clinically, AR is a Th2-type inflammatory disease whereas asthma may be allergic or nonallergic,

distinguished by the presence or absence of IgE antibodies to common environmental allergens. About 80% of childhood and 50% of adult asthma is allergic (i.e., IgE associated). In the latter, the triggers are not so well defined, but are probably microbes or microbial components, viruses etc. However, both forms of asthma are characterized by T helper cell 2 (Th2) type inflammation, stimulation of inflammatory cells like mast cells, causing eosinophilia, leukocytosis, and enhanced B-cell IgE production, leading to airway hyperresponsiveness and tissue remodeling. The determinants for an individual to develop an asthmatic phenotype require both exposure to appropriate stimuli and a genetic predisposition.^{2,3}

Airway remodeling involves structural changes in the airway wall characterized by epithelial alterations (epithelial shedding or denudation as a result of toxic proteins released especially by eosinophils), goblet cell metaplasia (their numbers increase first), bronchial mucous gland enlargement (excess production of mucus), subepithelial fibrosis and extracellular matrix deposition, increased smooth muscle mass and increased vascularization.² Chronic inflammation causes tissue injury, which is partly repaired between inflammatory exacerbations. Airway remodeling is associated with poorly-reversible or non-reversible airway narrowing, more severe airflow limitation and airway hyperresponsiveness and may be involved in component of adult severe asthma.⁴ These changes involve both large and small airways, but the individual variation is large. Airway secretions also contribute to the air flow limitation in asthma, and increase in amount and viscosity of the secretions plays a crucial role especially in acute, life-threatening exacerbations.⁵ However, remodeling is not a characteristic feature of AR.

Although much more is known in terms of the inflammatory mechanisms underlying AR and asthma, there is still a gap in our understanding of the cellular and molecular pathways involved in remodeling. Both inflammation and remodeling occur in the tracheobronchial tree of patients with asthma. Persistent, eosinophilic inflammation seems to be a prerequisite for the development of remodeling.⁶ Increased vascularity and expression of vascular endothelial growth factor (VEGF) are features of the asthmatic airway, but little is known of their contribution of airway remodeling. Siddiqui et al.⁷ showed that vascular remodeling is a feature of asthma, and is inversely correlated with postbronchodilator Forced Expiratory Volume in One Second (FEV₁) indicating a role in airflow obstruction.

Until recently, airway remodeling was considered to be a secondary phenomenon, developing later in the disease process as a consequence of persistent inflammation. The presence of airway inflammation and remodeling in children⁸ with asthma indicates that the

process of remodeling begins early in the disease process of asthma and synchronously with ongoing and repeated airway inflammation rather than as consequence of airway inflammation. Structural changes of the airways are already present and maximal in severely asthmatic school children,⁹ indicating that changes begin early in life between the ages of 1 and 3 years; and in good association with tissue eosinophilia and reticular basement membrane thickening.¹⁰ This gives a window of opportunity for early intervention that can possibly modify the natural history of asthma.¹¹

CHRONIC INFLAMMATION IN ALLERGIC RHINITIS AND ASTHMA

Effector cells of inflammation

The upper and lower airways have a similar mucosal structure, and exhibit similar inflammatory reactions to irritants and allergens. Increasing evidence indicates that there is marked similarity in the degree of cell infiltration or cytokine expression during allergic inflammation between the nasal mucosa and bronchial mucosa.^{12,13} However, there are minimal structural changes of the nasal mucosa in AR unlike that in asthma.

In allergic asthma and AR, inhaled allergens penetrate the mucociliary lining and enter the airway epithelium either via the tight junctions that surround the apical zone of epithelial cells or by direct uptake by the cells per se. Allergens are presented to T cells that then react by forming a pool of Th2 cells (IL-4, IL-5 and IL-13 secreting T cells) that drive the production of IgE via interaction with B cells. Subsequently the IgE binds to the high affinity IgE receptor on mast cells, and cross-linking of allergen specific mast cell surface IgE by allergens causes the release of mediators such as histamine and leukotrienes. They increase vascular permeability and initiate a cascade of recruitment of more inflammatory cells and further release of pro-inflammatory mediators.

Mast cells and basophils

Mast cells are critical in mediating the early phase of inflammatory response in AR and asthma. However, more recently it is known that mast cells are not only effector cells of the immediate phase response, but also play a role in ongoing allergic inflammation. They store and produce Th2 type cytokines, and chemokines, induce IgE synthesis in B cells,^{14,15} express the Cysteinyl Leukotriene 1 Receptor (CysLT₁),¹⁶ Glucocorticoid Receptor (GR),¹⁷ and SAF-2,¹⁸ and upregulate the production of cytokines/ chemokines in epithelial cells and fibroblasts.¹⁹

Accumulation of mast cells into the nasal and bronchial epithelium, a feature of AR and atopic

asthma, is also reported in idiopathic rhinitis.²⁰ RANTES (Regulated on Activation Normal T-Cell Expressed and Secreted chemokine) and TGF-beta (Transforming Growth Factor beta) are implicated in the intraepithelial migration of mast cells.^{19,21} Increase in mast cells and basophils occurs within 1 hr of nasal allergen challenge (NAC).²² Patients with AR with or without asthma have similar number of intraepithelial mast cells,²³ and segmental bronchial challenge (SBC) induces an increase in basophils and eosinophils which predominate in nasal and bronchoalveolar lavage (BAL).²⁴⁻²⁷

Airway mast cells stimulated via the FcεRI are an important source of Th2 cytokines, proinflammatory cytokines like TNF-alpha (Tumor Necrosis Factor alpha, preformed in mast cells) and chemokines like Thymus and Activation-Regulated Chemokine (TARC), as well as the IL-7 like cytokine Thymic Stromal Lymphopoietin TSLP.^{28,29} FcεRI-mediated TSLP production from mast cells is further enhanced in an autocrine manner by IL-4.³⁰ Moreover, mast cells can interact with structural cells like epithelial cells to enhance the production of cytokines and chemokines from epithelial cells. TNF-alpha in concert with IL-4 and IL-13 enhances the production of TARC, TSLP and eotaxin from epithelial cells.¹⁹ These result in increased infiltration of Th2 cells, eosinophils and differentiation of dendritic cells. Moreover, tryptase and chymase from mast cells can upregulate RANTES and GM-CSF (Granulocyte-Macrophage Colony-Stimulating Factor) production from epithelial cells.¹⁹ TNF-alpha also promotes antigen- and Th17 cell-dependant neutrophilia after allergenic stimulation and induces dendritic cell migration.^{31,32} Murine mast cells have been shown to induce CD4+T-cell migration in vitro, but down-regulate FcεR1 expression in T reg cells, while activated Treg cells suppress mast cell FcεR1 expression.

Mast cells can induce IgE synthesis in B cells and local IgE synthesis has been demonstrated in the nasal mucosa of patients with AR.^{33,34} Both IL-4 and IgE can upregulate FcεRI expression in mast cells suggesting a crucial role for the mast cell-IgE-IgE receptor cascade in ongoing inflammation in AR and asthma.³⁴ A role for mast cells in antigen presentation is also suggested.³⁵ While classically mast cell activation occurs following the allergen crosslinking of FcεR1-bound, allergen-specific IgE, mast cells may also be activated through multiple other mechanisms like via the complement receptors and Toll-Like Receptors (TLRs) even in the absence of FcεR1 stimulation.³⁶⁻⁴⁰

Basophils are increased in the nasal secretions and bronchoalveolar lavage of patients with AR and asthma. Like mast cells, basophils can also initiate allergic inflammation through the binding of allergen-specific IgE antibodies to the FcεR1 on basophil cell surface.⁴¹ Basophils also drive Th2 cell differentiation of activated

naive CD4+T cells via production of IL-4 and direct cell-cell contact.⁴² Basophils in AR and atopic asthmatics also express high levels of the FcεRI.⁴³ Down regulation of FcεRI in basophils and mast cells comprises one of the mechanisms of anti-IgE monoclonal antibody (Omalizumab).

Eosinophils

Eosinophils increase in both the early and late phase of AR⁴⁴ and correlate with nasal flow, IL-4, IL-5, IL-8 and IFN-gamma, spirometric values, methacholine test positivity, percentage of predicted FEV₁ and BHR.⁴⁵⁻⁵⁴ Eosinophils are increased in symptomatic atopics and increase in nasal as well as bronchial epithelium and lamina propria after NAC^{22,55-57}. Eosinophils are a major source of Macrophage Migration Inhibitory Factor (MIF)⁵⁸ and Nerve Growth factor (NGF),⁵⁹ express 5-Lipoxygenase (5-LO), Five-Lipoxygenase Activating Protein (FLAP), Leukotriene C₄ Synthase (LTC₄S), and CysLT₁ and CysLT₂ receptors,⁶⁰ are involved in the loss of epithelial integrity⁶¹ and are also seen in the esophageal mucosa of symptomatic patients with respiratory allergy.⁶²

IL-5 has a key role modulating eosinophil differentiation and survival. Targeting IL-5 as a therapeutic strategy for allergic asthma with anti-IL-5 therapy has demonstrated marked reduction of peripheral blood eosinophils, but only partial abrogation of the pulmonary eosinophilic response and minimal impact on clinical outcomes.^{63,64} More recently, in selected patients with refractory, eosinophilic asthma anti-IL-5 monoclonal antibody (mepolizumab) reduced both peripheral blood and sputum eosinophil levels, asthma exacerbations, and resulted in a reduction in oral corticosteroid dose.^{65,66}

Eosinophil chemoattractants include eotaxin, macrophage/monocyte chemotactic protein 4 (MCP4), RANTES, and CysLT among others.^{67,78} They act on distinct cell surface receptors (e.g., CC chemokine receptor 3 [CCR3] and CysLT₁) present on the eosinophil, but not exclusively so. Challenge with leukotriene E₄ (LTE₄) or LTD₄ results in increased nasal secretion and nasal obstruction in AR and greatly increased numbers of eosinophils in the bronchial wall.^{67,68} CysLT₁ receptor is expressed on a variety of nasal and bronchial inflammatory cells including eosinophil, neutrophils, mast cells, macrophages, B lymphocytes, and plasma cells.⁶⁹ These numbers of inflammatory cells expressing the CysLT₁ receptor are also increased in AR and asthma and there is a further, significant increase among patients experiencing a severe exacerbation of asthma leading to hospitalization.⁷⁰

Activated eosinophils release highly toxic granules, the evolutionary function of which has probably been killing the potentially dangerous invader. Especially in

asthma, eosinophil derivatives damage the surface epithelial cells, loosening their attachments and resulting in shedding of cells into the airway lumen, where they admix with eosinophils, neutrophils and mucus.

T lymphocytes

In perennial AR (PAR) patients, CD3+, CD25+ (activated) and CD45RA+ (naïve) T lymphocytes are increased in the nasal mucosa. In PAR, memory T cells and in idiopathic rhinitis CD8+ T cells correlate with mucosal mast cells.⁷¹ While CD86 is expressed on CD19, CD1a, CD14 and CD3 T cells in PAR, CD80, CD28, and CD152 are expressed after NAC.⁷² Moreover, CCR4+ CD4 cells are increased in AR.⁷³ Besides Th2 cytokines and chemokines, T cells in AR express IL-16,⁷⁴ CXCR1⁷⁵ and CX(3)CR(1).⁷⁶ Mucosal gamma delta T cells in PAR and asthma are increased, induce IgE synthesis in B cells and induce proliferation of T cells.⁷⁷ CD23+ B cells increase in PAR, but do not correlate with the mucosal Th2 cells.⁷³

Although conventionally both AR and allergic asthma have been considered to be due to a disruption in the normal Th1/Th2 balance, more recently, there is new evidence on the emerging roles of Th17 cells which are a distinct subpopulation of CD4+ T cells that produce IL-17A, IL-17F, IL-22, TNF-alpha, and IL-21.⁷⁸ Th17 cells were found in the nasal mucosa of AR patients⁷⁹ and in bronchial biopsies of asthmatics.⁸⁰ IL-17 induces the release of proinflammatory cytokines / chemokines from a variety of cell types,⁷⁹ is linked to the development of airway neutrophilia, and its presence in the asthmatic airway correlates with increased severity of disease. Treg cells play roles in the determination of self tolerance and the regulation of immune responses. Th17 and Treg cells have opposing actions, and T reg cells secrete IL-10 and TGF-beta and are increased in patients after immunotherapy.

Neutrophils

Although neutrophils are predominantly increased in non-allergic infective rhinitis and chronic rhinosinusitis, the increased expression of activation markers on neutrophils and Myeloperoxidase (MPO) levels in AR and increase in neutrophils in bronchoalveolar lavage after nasal allergen challenge suggests roles for neutrophils in AR.^{25,81} In acute, severe exacerbations of asthma, there are increased eosinophils and neutrophils within the airway, and the increase in neutrophils is proportionately greater.⁸² Inhaled corticosteroids reduce airway eosinophils but increase airway neutrophils and neutrophil chemoattractant IL-8, with loss of asthma control.⁸³

Epithelial cells

Conventionally, epithelial cells placed at the interface between the external environment and the host have been considered to play a role as a defense barrier against environmental agents. However, over the past several years the roles of epithelial cells as effector cells have become more evident, directly via the action of inflammatory mediators as well as via cell-cell interaction with immune cells. Moreover, its pivotal position in orchestrating airway remodeling and fibroblast proliferation is also crucial.⁸⁴

Airway epithelial cells are an important source of a variety of inflammatory mediators including multifunctional cytokines and chemokines like IL-1, IL-6, IL-8, TNF-alpha, GM-CSF, RANTES, Eotaxin, TARC leading to their crucial role in the migration and activation of immune cells like eosinophils, basophils and Th2 cells.^{85,86} More recently, TSLP derived from epithelial cells is increased in the nasal mucosa of AR patients and in the asthmatic airway.⁸⁷ TSLP can activate dendritic cells, promote Th2 responses, and activate mast cells.³⁰ Epithelial cells also express co-stimulatory molecules like CD86 and HLA-DR, CD86 and the FcεRI can present antigen to T cells.^{88,89} Particulate matter like diesel exhaust particles can induce the release of proinflammatory mediators and enhance the expression of co-stimulatory molecules on epithelial cells. The expression of IL-17F in the the airways of asthmatics correlates with disease severity and induces several asthma-related molecules such as CCL20 that can attract Th17 cells into the airway thus amplifying airway inflammation. A recent study demonstrated that bronchial epithelial cells express IL-17F in response to IL-33 via ST2-ERK1/2-MSK1 signaling pathway.⁹⁰ Moreover, IL-17F is involved in airway remodeling and steroid resistance. Hence, IL-17F may be a valuable therapeutic target for development of novel strategies. Recently the importance of serum periostin measurements and the role of anti-IL-13 as a treatment in those with high periostin levels and anti-IL-5 in those who are steroid sensitive with high eosinophils demonstrated that there are phenotypes and endotypes of asthma.

REMODELING IN ASTHMA

Remodeling is defined as a change in structure that is inappropriate to maintain normal airway function.^{91,92} Some features of remodeling are evident, even in newly diagnosed or mild asthmatics and is characterized by epithelial fragility and reticular basement membrane thickening. With increasing severity of asthma, the changes are more pronounced and clear: increases of airway smooth muscle mass, vascularity, numbers

of fibroblasts, and interstitial collagen, as well as mucous gland hypertrophy.⁹³ These changes appear to be greatest in the larger, more proximal airways. Thickening of the reticular basement membrane occurs early in asthma, even before diagnosis, and is detected in children with mild asthma.⁹⁴ In school children between the ages of 6 and 16 with severe asthma there is already maximally thickening but with no relation to the age or symptom duration.⁹ These changes appear in preschool wheezy children by the age of 29 months.⁷ Bourdin et al.⁹⁵ showed recently that reticular basement thickness is a hallmark of severe asthma, but not of mild asthma or Chronic Obstructive Pulmonary Disease (COPD).

Airway smooth muscle surrounds the airways as two opposing helices, i.e., a geodesic pattern and as muscle shortens, it constricts and also tends to shorten the airway against an elastic load. Airway smooth muscle mass is increased in the asthmatic airway.⁹⁶ Airway smooth muscle cells can secrete mediators that may promote mast cell chemotaxis, proliferation, and survival, while cell-cell interaction between airway smooth muscle cells and mast cells enhances activated complement-induced mast cell degranulation.⁹⁷⁻⁹⁹ Human lung mast cells migrate toward Th2 cytokine-stimulated airway smooth muscle cells from asthmatics, while supernatants obtained from airway smooth muscle cell cultures of non-asthmatics inhibit this chemotaxis.¹⁰⁰ Large numbers of mast cells have been found located within bronchial smooth muscle of asthmatics and mast cell mediators, such as tryptase and cytokines, can modulate airway smooth muscle cell function. Mast cells also express MMP9 and contribute to multiple features of chronic asthma and they play an important role in tissue remodeling.¹⁹

Bronchial epithelial cell experiments have demonstrated a role for TLR signaling in the activation of epidermal growth factor receptor, suggesting a role for TLRs in potentiating remodeling.¹⁰¹ In the asthmatic airway, there are increased numbers of subepithelial myofibroblasts, and allergen challenge in people with asthma leads to increased accumulation of myofibroblasts in the airway mucosa.^{84,102} Histamine can induce the transition from fibroblasts to myofibroblasts (as measured by alpha-smooth muscle actin expression), and induce connective-tissue-growth-factor expression in fibroblasts, suggesting the ability to participate in the process of remodeling.^{102,103} Fibroblastic infiltration of the lung may be secondary to the recruitment of circulating bone marrow-derived progenitors of fibrocytes to the airway and to the proliferation and expansion of resident fibroblasts, or possibly, epithelial cells may undergo phenotypic change to effector fibroblasts through a process termed epithelial-

mesenchymal transition. Airway epithelial cells derived from asthmatics demonstrated increased susceptibility to TGF β -induced epithelial-mesenchymal transition than those derived from normal subjects.¹⁰⁴ In addition to stimulation of epithelial cells and extracellular matrix synthesis, TGF β can elicit other responses in bronchial fibroblasts, including stimulating their proliferation and synthesis of a range of growth factors. It also has effects on the asthma susceptibility gene, a disintegrin and metalloprotease (ADAM)³³, which has been implicated as an asthma remodeling gene.

Measuring airway function (e.g. FEV₁) can give indirect information on the long-term airway inflammation and structural changes, but cannot detect the early inflammatory processes. Such inflammatory changes can exist even in patients with normal lung function but with symptoms indicative of asthma.¹⁰⁵ On the other hand, symptomatic infants, who have reversible airflow obstruction may not have bronchial mucosal eosinophilia or remodeling.¹⁸

How early can remodeling start

Airway biopsy studies in children suggest that pathologic changes such as epithelial loss, basement membrane thickening, and angiogenesis occur early in the asthmatic airway. In children with difficult asthma (mean age of 13, age range of 6-16) recruited to investigate whether the thickening of reticular basement membrane could occur in childhood asthma⁹ have shown that the thickening of reticular basement membrane was seen in the airway. In another study, Barbato et al. have examined the biopsy specimens of airway in children that is nine children with asthma (age of 4-12 yr), six children with atopy without asthma (age of 4-12 yr), and eight control children without asthma or atopy to elucidate whether airway inflammation and remodeling could occur even in mild childhood asthma,⁹² demonstrated that airway eosinophilia and basement membrane thickness were present in children with mild asthma, and even in children with atopy without asthma. This indicates that 1) airway inflammation shown by airway eosinophilia occurs even in mild as well as in difficult asthma, 2) airway inflammation occurs in the airway before developing episodic wheezing, although asthma symptom is difficult to establish in children, 3) the presence of both airway inflammation and remodeling indicate that remodeling process begins early in the disease process of asthma and occurs synchronously in ongoing and repeated airway inflammation rather than as subsequent event of airway inflammation.

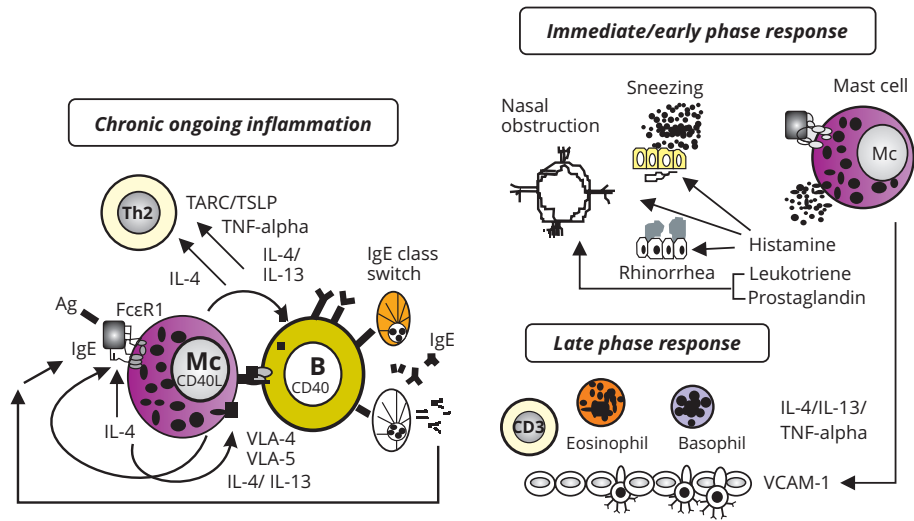


Figure 1 - Inflammatory mechanisms in allergic rhinitis (Modified from Pawankar et al. Allergic Rhinitis pathomechanisms)

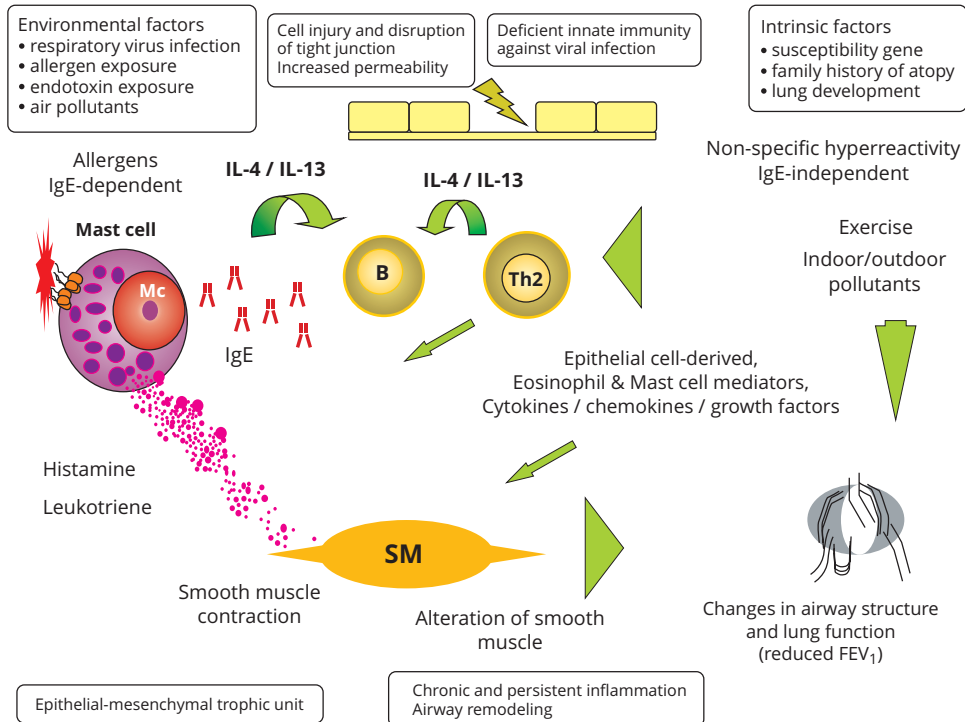


Figure 2 - Pathomechanisms of asthma

Summary

With the increasing evidence on the links between allergic rhinitis and asthma from epidemiologic, immunologic and clinical studies, early intervention and downregulation of inflammation are key to better control of both AR and asthma. New modes of immunomodulatory therapies and biologics that target specific phenotypes like IgE, IL-5, IL-13 and the IL4 receptor alpha chain have shown some efficacy in phenotyped patients. For those without evidence of Th2 inflammation, no specific therapies have been identified. Anti-IgE - and anti-IL-5, anti-IL-13, anti-IL4 R antagonists, and therapies that target TSLP, IL-33, IL-17 as well as those that induce tolerance hold more promising outcomes, but their effect on halting airway remodeling in severe asthma is not known. Clinical cluster analysis from the Severe Asthma Research Program (SARP) identified 5 asthma subphenotypes that represent the severity spectrum of early-onset allergic asthma, late-onset severe asthma, and severe asthma with chronic obstructive pulmonary disease characteristics. Analysis of induced sputum from a subset of SARP subjects showed 4 sputum inflammatory cellular patterns. Subjects with concurrent increases in eosinophil ($\geq 2\%$) and neutrophil ($\geq 40\%$) percentages had characteristics of very severe asthma. This multivariate approach identified 4 asthma subphenotypes representing the severity spectrum from mild-to-moderate allergic asthma with minimal or eosinophil-predominant sputum inflammation to moderate-to-severe asthma with neutrophil-predominant or mixed granulocytic inflammation.¹⁰⁶

Given the immunological similarities that exists between the patterns of inflammation in asthma and allergic rhinitis as well as chronic rhinosinusitis and asthma and the impact of AR on asthma, treatment should focus to have a global approach to treat both the upper and lower airways for better outcomes.

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