Molecular Mechanisms of Metaplasia, Differentiation and Hyperplasia of Goblet Cell in Allergic Asthma

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Abstract

Bronchial allergic asthma (asthma) is an airway inflammation characterized by airflow obstruction of variable degrees with bronchial hyper-reactivity and is induced by a complex interaction of environmental and genetic factors. Asthma of the human can be divided into an immediate- and a late-phase reaction, and some of the patients develop a late-phase reaction after a symptom-free interval. Hallmarks of asthma are mucus overproduction by goblet cells associated with responses of helper T(Th)2 cells. Mucus hypersecretion from goblet cells themselves or by metaplasia and/or hyperplasia of goblet cells appears to be associated with disease severity in asthma, because mucus production by those cells in local bronchial–bronchiolar lesions causes airway mucus plugging. However, the mechanisms of mucus production are not fully understood. Molecular mechanisms of goblet cell metaplasia, differentiation and hyperplasia will be reviewed in this article. Also, the relationship between allergic inflammation in Th1/Th2 paradigm shift and thymic stromal lymphopoietin (TSLP) was included for the understanding of goblet cell responses. The clarification of mechanisms of mucin production in vivo may lead to the development of novel therapeutic strategies to suppress mucus production in asthma.

Keywords: Asthma; Goblet cell; Clara cell; Mucus; Inflammatory Th2; TSLP; Epithelium

Introduction

Bronchial allergic asthma (asthma) is a prevalent disease and is increasing in developed countries. Asthma is an inflammatory disorder characterized by airflow obstruction of variable degrees with bronchial hyper-reactivity. Asthma is caused by environmental factors and induced by a combination of genetic and environmental stimuli. Genetic studies have revealed that multiple loci are involved in the etiology of asthma [1]. Asthma of the human can be divided into an immediate- and a late-phase reaction, and some of the patients develop a late-phase reaction after a symptom-free interval [2]. Helper T(Th)2 cells play a fundamental role in both immediate and late reactions. Immediate reaction is induced by IgE-mediated increased mast cell response, whereas late response is characterized by recruitment of eosinophils, basophils, and Th2 cells producing cytokines such as IL-4, a switch factor for IgE synthesis, and IL-5, an eosinophil growth factor [3]. In addition, late reaction may be allergen dependent and IgE independent. Moreover, Tang et al. [3] reported that the chronic response in asthma is thought to include repeated epithelial shedding, mucus secretion, thickening of the basal lamina, airway wall fibrosis, smooth muscle hypertrophy, angiogenesis and hyperplasia of the submucosal gland. The degree of mucus production appears to be associated with disease severity in patients with asthma [4]. Hallmarks of asthma are goblet cell hyperplasia and mucus overproduction, leading to cough, shortness of breath and wheezing. In the most severe cases of asthma these effects can be dramatic, and pathologic evidence suggests that mucus plugging may be the cause of death in fatal asthma [5]. Goblet cells in the airway epithelium at autopsy were demonstrated to be 30-fold higher in patients with fatal asthma than those with asthma who died of other causes [5].

Thus, goblet cell responses in asthma play an important role in the pathogenesis of asthma (Figure 1). Recent cellular, molecular and animal-model studies have revealed several cellular events including inflammatory helper Th2 cytokines that are involved in the progression of asthma, leading to goblet cell responses. This article focusses on goblet cells and introduces molecular mechanisms of goblet cell metaplasia, differentiation and hyperplasia in asthma (Figures 2A and 2B).

Goblet Cell Function

In this section general information of goblet cell function will be explained briefly. In the normal human airway, mucus is produced and secreted by goblet cells. The submucosal glands in the larger airways also contribute to mucus production. Mucus is composed of water, ions, lipids, proteins, and complexed macromolecular glycoproteins called mucins, which impart viscoelastic and gel-forming properties to mucus [6]. More than 20 mucin genes have been identified, and 12 of these have been shown to be expressed in the respiratory epithelium [7]. There are two structurally and functionally distinct classes of mucin such as membrane-bound mucins and secreted mucins. Membrane mucins, which have transmembrane and cytosolic domains, are tethered to the plasma membrane, where they participate in functions such as cellular adhesion, pathogen binding, and signal transduction [8,9]. Proteolytic cleavage or synthesis of alternative splice variants lacking the transmembrane domain can result in release of membrane mucins into the mucus layer [10]. Secreted mucins are synthesized in epithelial cells and stored in intracellular secretory granules until stimulated for release by regulated exocytosis. The gel-forming mucins, MUC5AC and MUC5B, are the most prominent secreted mucins in the respiratory tract. When secreted, gel-forming mucins form a dense macromolecular matrix providing the adhesive and space-occupying properties of the mucus

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cells have many mucin granules and are found throughout the glandular and surface airway epithelia and can be detected histochemically using alcian blue (AB) and periodic acid Schiff (PAS)'s stains. Submucosal glands, which contain a mixture of goblet cells and serous cells, are distributed throughout the cartilaginous airways in humans, but are limited to the laryngeal region of the trachea in mice [15]. In the small airways of humans and in all intrapulmonary bronchi-bronchioles in mice, there are very few goblet cells under basal conditions. In mice, for example, ciliated cells (~40%) and nonciliated cells (~60%), which are mostly Clara cells, account for the vast majority of airway epithelial cells [17]. In normal specific pathogen-free mice, there were mainly three cell types in the epithelium of the intrapulmonary bronchi-bronchioles such as ciliated, non-ciliated (Clara) and basal cells [18]. The epithelium was reduced in thickness in the more distal airways in human and mouse. Both the ciliated and Clara cells adhering to neighboring cells by desmosomes were seen equally throughout the bronchus and bronchiole. Clara cells with or without a few short microvilli have a lot of smooth endoplasmic reticulum (s.e.r.) and mitochondria, appearing dark with a few or no cristae. A few round, homogeneous fine electron-dense secretory granules (S.G.), which do not have a clear membrane bound, corresponding to few granules (a varying degree) stained with toluidin blue (TB) were also located most regularly in the apical portion of Clara cells. Non-ciliated cells without S.G., which seem to be Clara cells, are often observed. S.G. appeared to be in the process of extrusion into the lumen of the airway by meroerythroblast and apoptotic secretions in some Clara cells [18]. On the other hand, basal cells, which are small polyhedral cells on the basement membrane, were few and distributed throughout the bronchus and bronchiole and adhered to neighboring cells by desmosomes. The nucleus occupied a large portion of the cells with tonofilaments having a few organella.

### Goblet Cell Metaplasia from Clara and Ciliated Cells

In this section goblet cell metaplasia from mainly Clara and ciliated cells will be introduced. Epithelial cell hyperplasia and metaplasia are common consequences of airway inflammation and may be associated with protective as well as pathogenic outcomes. Metaplasia implies a change in cell phenotype, while hyperplasia indicates cell proliferation as the mechanism for the increase in goblet cell numbers. Clara cells have a capacity to differentiate into goblet cells [19]. In tracheal epithelium in a rat model that epidermal growth factor receptor (EGFR) activation may promote selective cell differentiation (not proliferation), from Clara cells to goblet cells [20, 21]. Nadel [21] hypothesized that the mechanism of EGFR expression and activation as follows (Figure 2A). Stimulation of airway epithelial cells with tumor necrosis factor (TNF)-a causes EGFR expression and EGFR ligandsbind to EGFR, resulting in EGFR tyrosine phosphorylation and a subsequent downstream cascade, which causes mucin gene and protein expression. Human airway epithelial cells stimulated with the EGFR ligands such as transforming growth factor-a (TGF-a) and EGF, develop a mucous phenotype [21, 22]. Intratracheal administration of these ligands after TNF-a-induced up-regulation of EGFR on airway epithelial cell in rats resulted in mucous metaplasia [22]. EGFR activation is required for inducing mucous metaplasia in animal models and for up-regulation of mucin expression in human airway epithelial cells in response to allergens, viruses, neutrophils, and cigarette smoke in human with asthma [22-25]. This chronic switch in epithelial behavior exhibits genetic susceptibility and depends on persistent activation of EGFR (also designated ErbB1 and HER1) signaling to PI3K that prevents apoptosis of ciliated cells, and IL-13 signaling that promotes transdifferentiation of ciliated to goblet cells.
of transcription factors that are important in lung development have been identified in the airway epithelium and have been shown to play key roles in mucous metaplasia. The forkhead box transcription factor FOXA2, regulates genes that are involved in lung maturation and epithelial cell differentiation. FOXA2 inhibits MUC5AC gene expression [37]. Allergen challenge or transgenic overexpression of IL-4 or IL-13 in mice resulted in pulmonary inflammation and goblet cell metaplasia, which was associated with significant inhibition of FOXA2 expression in the airway epithelium [38]. Thus, FOXA2 is required for maintenance of normal differentiation of the airway epithelium and inhibition of FOXA2 appears to be an important early step in the initiation of mucous metaplasia.

Thyroid transcription factor 1 (TTF-1) plays a critical role in peripheral lung morphogenesis and is an important regulator of genes involved in many biological functions in the lung, including host defense, fluid balance, surfactant homeostasis, lung vasculogenesis, and epithelial cell differentiation [39]. ETS transcription factor (SPDEF), an epithelial specific transcription factor that is markedly induced by a phosphorylation mutant of TTF-1, was shown to be constitutively expressed in proximal airway epithelial cells and submucosal glands in the adult mouse lung [40]. Transgenic overexpression of SPDEF in the murine airway epithelium caused spontaneous mucous metaplasia. SPDEF appeared to interact with TTF-1 to have a synergistic effect on the promoters of a number of genes but had no direct effect on the MUC5AC gene promoter. SPDEF overexpression did, however, result in down-regulation of FOXA2 expression. SPDEF has been shown to be both necessary and sufficient for induction of a transcriptional program that results in goblet cell differentiation [41]. Expression of a member of the Clca gene family (i.e., calcium-activated chloride channel-3; mClca3) with the development of mucous cell metaplasia but not airway hyperreactivity [42] has been reported. The 5′-regulatory region of the mClca3 gene contains a putative binding site for the Stat6 transcription factor that mediates IL-13R signal transduction and that IL-13 stimulates mClca3 (and mClca5) gene expression in cultured airway epithelial cells [43]. The mouse system appears to be directly relevant to human airway disease because the human homologs for mouse mClca3 and mClca5, human calcium-activated chloride channel-1(hCLCA1) and hCLCA2, are both overexpressed in airway tissue of subjects with asthma [43,44]. mCLCA-3-gob(5)/human CLCA-1 acts as a downstream molecule of Th2 cytokines. IL-4/IL-9/IL-13 signals, playing an important role in mucus production and mucus transport, is induced by IL-13 and causes mucus production in airway epithelial cells [45].

Epidermal growth factor (EGF) and trefoil factor family (TFF) peptides support synergistically several processes of goblet cell metaplasia [46]. TFFs are well known particularly for their key role in mucusal reconstruction and TFFs also support mucosal differentiation processes. Particularly the latter process reflects the plasticity of the airway epithelium and causes intense airway remodeling. Goblet cells are derived, at least in part, from Clara cells, which trans-differentiate from a serous into a mucous phenotype. Using a murine asthma model it was shown that trans-differentiating Clara cells specifically express TFF1, which is stored in a specific subset of secretory granules [47]. These points to a role for TFF1 as an autocrine factor for the trans-differentiation of Clara cells toward goblet cells [47]. It is demonstrated that induction of TFF3 synthesis was shown with differentiation in vivo humanized tracheal xenograft and in vitro air-liquid interface culture models. Furthermore, exogenous TFF3 promoted differentiation of ciliated cells in an EGFR-dependent manner. Taken together, both studies imply that TFFs may play key roles for various differentiation processes of the airways [46].

**Figure 2:** Nadel [21] reviewed that stimulation of airway epithelial cells with TNF-α causes EGFR expression. Curran and Cohn [33] proposed the hypothesis that goblet cell differentiation requires two signals (A). The hypothesis: Signal 1 activates EGFR on the epithelial cells (Clara and ciliated cells), leading to inhibition of epithelial cell apoptosis. Epithelial cells that survive have the potential to become goblet cells if provided with Signal 2. IL-13 binding to its receptor. Upon IL-13R and Stat6 activation, epithelial cells show an increase in SPDEF and reduced FoxA2 expression. GABA_R responsiveness is required for goblet cell metaplasia, which is induced by IL-13. After expression of MUC5AC, CLCA and TIFF, cells lose the features of epithelial cells from which they arose and become goblet cells that produce mucins. Also reversible changes from established goblet cell metaplasia to ciliated cells may occur. Hypothetical differentiation and hyperplasia depending on tentative working model for the self-renewal and differentiation of basal stem cells in mouse and human airways proposed by Rock et al. [82] are shown in (B) with some modification. A subset of BCs in the normal airway epithelium expresses both KRT5 and KRT14. BCs self-renew over the long term and generate ciliated, secretory cells and goblet cells. On the other hand, Clara cells expressing SCGB1A1, may be one of stem cells, can proliferate and give rise to goblet, Clara and ciliated cells via progenitor cells. Also Sox2 is required for proliferation and differentiation of Clara and basal cells. Consequently, metaplasia from Clara cell to goblet cells and ciliated cells may occur after differentiation.
Mucous metaplasia does not occur in all immune responses, particularly those rich in IFN-γ by mutual inhibitory effects between Th1 and Th2 cells (Figure 3). Th1 cell–induced airway inflammation did not stimulate mucous metaplasia [55]. IFN-γ inhibits mucus cell transition, even in the presence of high levels of Th2 cytokines (Cohn et al. [49], unpublished data). Thus, IFN-γ may inhibit airway epithelial mucus by blocking steps required for IL-13 to act, such as inhibiting EGFR actions to promote cell survival. Alternatively, IFN-γ may act by inhibiting downstream effects of IL-13. The molecular mechanisms by which IFN-γ inhibits mucus are still being defined.

In cultured human airway epithelial cells, IFN-γ was shown to inhibit IL-4–induced STAT6 phosphorylation and to reduce the expression of IL-4 target genes [56]. Furthermore, IFN-γ also induced expression of suppressor of cytokine signaling-1 and -13RA2, inhibitors of IL-4/IL-13 signaling, as well as increasing the decay of IL-4 target gene mRNA. Thus, IFN-γ may inhibit Th2-induced responses in epithelial cells through a number of pathways. IFN-γ also promotes resolution of allergen-induced mucus metaplasia by stimulating airway epithelial apoptosis via Bax- and caspase-dependent mechanisms [57]. IFN-γ-induced cell death involves STAT1-dependent translocation of Bax to the endoplasmic reticulum [58]. Serha et al. reported [59] that peristin, a 90-kDa member of the fasciclin-containing family and functions as part of the extracellular matrix, is expressed in a variety of tissues and expression is increased in airway epithelial cells from asthmatic patients as follows. Sensitization and challenge of peristin-deficient mice with OVA resulted in increased peripheral Th2 responses. In the lungs, peristin deficiency resulted in increased airway resistance and significantly enhanced mucus production by goblet cells concomitant with increased expression of gob5 and Mcu5ac compared with wild type littermates. Peristin also inhibited the expression of gob5 involved in the regulation of mucus production, in primary murine airway epithelial cells, suggesting that peristin may be part of a negative-feedback loop regulating allergic inflammation.

Morphologically increased S.G., identical to Clara cell secretory (specific) protein (CCSP), are seen in the late allergic asthmatic model mice [18]. Clara cells having increased S.G. (CCSP) without mucus suggest that the significance of increased CCSP in allergic status may be the result of protection against injurious agents in environmental air primarily [32]. Thereafter, those cells with mucus may become goblet cells and there are two structural discharge of mucin such as secretion of mucins of membrane-bound mucins and apocrine fashion [18]. In addition, the loss of cristae and mitochondrial swelling has been found in an asthmatic mouse model [60]. Moreover increased CCSP will be induced by IL-13 [20] and IL-13 was shown to induce MUC5AC expression via a mechanism requiring a secondary TGF-β2/SMAD-mediated signal in airway epithelial cell cultures [61]. These suggest
that CCSP may be related with goblet metaplasia. Thus, allergen-induced goblet cell metaplasia from CCSP-positive Clara cells contains mucous granules in mice [18,62,66]. In fact, significant levels of CCSP-driven recombination can be found in ciliated and nonciliated airway epithelial cells [65]. CCSP and β-tubulin co-localization with MUC5AC in mice and in humans suggest that ciliated cells and Clara cells may each demonstrate sufficient plasticity to contribute to goblet cell metaplasia [66]. However, the relationship between increased S.G. and mucous production in Clara cells is not always identical [18]. Thus further study is needed to clarify this point. Alternatively hypoxia-inducible factor-1 (HIF-1) is a transcriptional activator that mediates gene expression in response to cellular oxygen concentrations [64]. HIF-1 is known to contain a conserved STAT6 motif in its promoter, and binding of HIF-1 to the MUC5AC promoter is induced by IL-13 and EGF stimulation including that HIF-1α is activated in antigen-induced airway disease through PI3K-δ signaling [65]. Goblet-cell metaplasia may occur in the intrapulmonary airway in a mouse model of asthma. A few fine granules, which correspond to PAS-positive granules stained with TB in semi-thin sections, are observed in the apical site of the non-ciliated epithelium (Clara cells) in normal mice. Thus most goblet-cell metaplasia in the bronchus and bronchiol of inflamed mucosa may be derived from Clara cells. In re-population of airway after injury, a morphological derivative of the Clara cell, referred to as a type-A cell, lack the differentiated ultrastructural features, such as s.6.c.r and S.G., that are characteristic of mature Clara cells [41] may be immature Clara cells (designate here as iClara cell) instead of type-A Clara cell. In asthma animal model, iClara cells and/or goblet cells may not be derived from neuroendocrine cells, which are only located in terminal bronchioles, since goblet-cell metaplasia was observed in bronchus rather than terminal bronchiole [18,66-68]. On the other hand, hyperplastic goblet cells at distal sites, at least in part, are considered to be metaplasia from Clara cells in human asthma [69]. In addition, it has been reported that IL-9 in airway injury repair resulted in goblet-cell hyperplasia in murine models of asthma [17]. Ciliated cells can be converted to goblet cells in primary cell culture from Guinea pig tracheal epithelium [26,70] as described above and that metaplasia is the dominant mechanism mediating the expansion in goblet cell numbers in human asthma [70].

**Goblet Cell Differentiation and Hyperplasia from Stem Cell**

In this section, goblet cell hyperplasia after differentiation (regeneration) from proliferated progenitor cells including basal cells (BCs) will be introduced. The increase in hypertrophic goblet cell numbers in the respiratory epithelium during airway inflammation has been described both as mucous metaplasia and as goblet cell hyperplasia. In human airways, a detailed analysis of the epithelial transition to a mucus-secreting phenotype has not been undertaken. In a murine model of asthma, there was a dramatic increase in goblet cell number after sensitization and allergen challenge, while the number of epithelial cells per unit surface area of the basal lamina remained constant and Clara cells were undergoing metaplasia to mucous cells [62] as mentioned already. Thus, these seem to be two types of increased goblet cells such as the metaplasia rather than hyperplasia and so called goblet cell hyperplasia explained below.

In the normal human airway epithelium, a tight balance between BC self-renewal and generation of physiologically appropriate proportions of secretory and ciliated cells is required [71]. This may be mediated by the behaviors of stem cells and potential multipotent intermediate progenitors. Alternatively the changes might result from altered paracrine signaling within the dynamic and complex BCs niche. In either case, hyperproliferating BCs secrete cytokines, such as IL-1, that promote airway wall fibrosis [71]. It is believed that Clara cells’ change their phenotype to either express mucins or chitinases, depending on their position in the airways [38], and there seems to be direct, reversible conversion of Clara cells into goblet cells, a change that is correlated with expression of SPDEF and downregulation of another transcription factor, FOXA2 [41], where SGB1A1+ Clara cells (Clara cells expressing secretoglobin, family 1A, member 1; also known as uteroglobin) can self-renew over the long term and they may be bronchoalveolar stem cells (BASCs) in the trachea, bronchiol and alveoli, which are maintained by distinct populations of epithelial progenitor cells [72]. Mucous metaplasia and the expression of SPDEF are observed in human small airways, in which BCs are abundant [72]. It will be important to model this condition in proximal mouse airways in which BCs are found. It is likely that cytokines such as IL-13, IL-6, IL-1β and TNFa directly or indirectly influence the fate of BCs and their immediate progeny [72,73,15]. In support of this idea, treating human air-liquid interface airway epithelial cultures (probably initiated mainly with BCs) with exogenous Notch ligand or IL-13 resulted in a drastic increase in the number of mucus-producing goblet cells [73]. Upstream events include a new immune axis for growth factor and cytokine production and downstream events that include ciliated epithelial cell survival and transdifferentiation to mucous cells as well as expression of CLCA genes [15].

Rock et al. [15] reviewed that BCs are a population of multipotent stem cells that drives both homeostasis of the normal epithelium and its orderly regeneration after injury. Disruption of the normal balance between BC proliferation and differentiation can lead, at two extremes, to BC hyperplasia or epithelial hypoplasia. Changes in the lineage choice of BCs or their undifferentiated daughters might contribute to the mucous cell hyperplasia, metaplasia or squamous metaplasia seen in many respiratory disorders. BCs are important in the initiation and progression of airway disease [15]. Among cytoskeletal proteins, cytokeratins 5 and 14 (KRT5 and KRT14, respectively) are typically expressed in BCs. There is differential expression of these proteins in airway BCs. At steady state, most mouse tracheal BCs express KRT5, whereas only a subset expresses KRT14 [82]. Mouse BCs generated lineage-labeled basal, ciliated and Clara cells [15]. However, during repair after naphthalene-induced injury, KRT14 expression is upregulated in the BC population [74-75]. KRT14 also has a more restricted expression pattern than KRT5 in the human airway, but is expressed in TRP63+ BCs in regions of squamous metaplasia [15]. Thus they pointed out that the functional significance of differential cytokeratin expression at steady state after injury and pathological changes of remodeling [15].

In the uppermost mouse trachea, BCs form a mostly continuous monolayer, but they are found in clusters or as individual cells more distally [76]. In the larger human airways, BCs also form a continuous monolayer but similarly become distributed into clusters and individual cells in the small terminal bronchioles [76]. In both the mouse and human, BCs form desmosomal contacts with neighboring columnar cells and are anchored to the basal lamina via hemi-desmosomes and other highly expressed adhesion molecules. The subjacent lamina propria contains mesenchymal cells, such as fibroblasts and immune cells (e.g. macrophages) [15] and dendritic cells (DCs) [77]. In addition, the mouse trachea and human intrapulmonary airways are richly endowed with blood vessels and nerves, which enter between the cartilage rings and encircle the airway [15,78]. Systemic blood vessels of the bronchial circulation supply many generations of
intrapulmonary airways in humans, whereas in the mouse, these vessels do not penetrate beyond the main stem bronchi, even after pathological stimulation of angiogenesis [79]. However, despite notable differences between mouse and human airways, BCs of the pseudostratified airway epithelia in both species are molecularly and histologically very similar. By definition, a stem cell can self-renew and generate differentiated progeny of one or more cell types. Relevant studies have been reviewed that BCs gave rise to both ciliated and Clara cells. By contrast, lineage-labeled Clara cells expressing SCGB1A1+ cells in the trachea proliferate and generate ciliated cells but a population was replaced over time by unlabeled progenitor cells, presumably BCs [12]. This suggests that, in airway epithelia containing BCs, the Clara cells do not function as stem cells over the long term under normal conditions. On the other hand, Randell [80] reviewed that both BCs and nonciliated columnar epithelial cells of the rodent trachea proliferate in the steady state and in response to injury [81,82]. BCs can regenerate a fully differentiated mucociliary epithelium when seeded in denuded tracheas (host tracheas that have been stripped of their epithelium) and transplanted subcutaneously in nude rats [83,84]. Alternatively, some luminal cells might have the capacity to regenerate BCs under these conditions. Preliminary evidence that mouse SCGB1A1+ cells can generate both self-renew and generate ciliated cells after epithelial injury in in vivo lineage-tracing studies [72]. Individual sorted TRP63-NGFR+KRT5+BCs from the mouse trachea self-renew give rise to clonal ‘tracheospheres’ containing both ciliated and secretory cells, and NGFR lumenal cells do not form tracheospheres. Among human tracheo-bronchial epithelial cell types, TRP63-NGFR+KRT5+BCs demonstrating their capacity to self-renew generate ciliated and secretory cells [15]. Also, human TRP63+KRT5+BCs are isolated from normal human airways [85]. These cells, estimated to constitute 0.01% of the total airway epithelium, proliferate and generate ciliated and secretory cells. This progenitor might still proliferate (in which case it would be considered a transit-amplifying cell) and give rise to either ciliated or secretory cells.

Tompkins et al. reported [86] that Sox2, which is a member of the SRY-high mobility box transcription factor family, is expressed in epithelial cells of the foregut, including pharynx, esophagus, trachea, bronchi, and bronchioles, but is excluded from the peripheral and alveolar regions of the lung physically, interacted with Smad3 and inhibited TGF-β1/Smad3-mediated transcriptional activity in vitro, a pathway that negatively regulates proliferation. Also Sox2 is required for proliferation and differentiation of Clara and basal cells that serve as the progenitor cells from which Clara, ciliated, and goblet cells are derived. The existence of such an early progenitor population is still hypothetical, but these cells might correspond to some of the ‘indeterminate’ or ‘intermediate’ epithelial cells that have been described in human airways [87].

Environmental Factors in Driving Goblet Cell Responses

In this section, interaction among DCs, respiratory epithelium and environmental factors being responsible for goblet cell responses in the development of inflammatory Th2 cells (Figure 3B) in airway allergy will be introduced [88] other than regulatory Th2 cells, which are already mentioned. DCs are the professional antigen presenting cells (APC) that have the capacity to present antigen to naïve-T (Th0) cells and T-effector (Te) cells and also DCs have an important role in the immunological outcome of the disease [89]. Circulating immature DCs were found to represent a small leukocyte population that produces large amounts of IFN-α in the presence of certain viruses [90]. Two major types of DCs have been characterized in humans: the CD11c+CD123+ myeloid dendritic cells (mDCs), for which there are several subtypes, and the CD11c+ CD123+ (IL-3Ra-chain) lymphoid cells known as the plasmacytoid DCs (pDCs) [91,92]. mDCs with CD11c+ DCs, a very potent antigen-presenting cell (APC) type, for their capacity to induce allergen-dependent activation of Th2 memory cells [93]. Tuma et al. [94] demonstrated the presence of mDCs in nasalmucosa with and without allergies. Jahnset al. [95] reported that pDCs, most likely has an important role both in innate defense against pathogens and as APCs in the adaptive immune system. This cell type exists in bone marrow, blood, organized lymphoid tissues and at effector sites with direct antigens (allergens) exposure such as the mucosae. In addition, pDCs are identified by their high expression of CD123, together with CD45RA and are present in low and variable numbers in normal nasal mucosa (including probably bronchial mucosa) but increased dramatically during the nasal (and also probably bronchial) allergic reaction [95].

DCs play a critical role in directing the types of T-cell responses, including Th1, Th2, and Th17. The ability of DCs to induce either Th1 or Th2 responses appears to be dictated by the type of signals that the DCs received at an immature stage [96-98]. In addition, bronchial epithelial cells, fibroblasts, smooth muscle cells, and mast cells have the potential to produce TSLP in human [96,99]. Wang et al. [100] reviewed the interaction between ligands and epithelial cells as follows, since exposure of the airway inflammatory stimuli through activation of epithelial Toll-like receptor (TLRs) to provide an important link between innate immunity and allergic disease. Additionally, airway epithelial cells can act as inflammatory promoters capable of directing DCs towards a Th2 response. TSLP production can be induced in airway epithelial cells by ligands that activate TLR2, TLR3, TLR8 and TLR9 to release TSLP protein, and TLR8 and TLR9 to stimulate TSLP gene transcription through NF-κB activation [100]. Ligands (TLR-LR) capable of activating these TLRs include bacterial lipoteichoic acid and peptidoglycans from bacteria (TLR2), single stranded (TLR8) and double-stranded (TLR3) viral RNA, and CpG DNA motifs in both viruses and bacteria (TLR9). TLR activation can also be initiated by other factors such as viruses, dustsand chemicals [100]. TSLP activated immature mDCs rapidly express TSLP receptor (TSLPR) and DCs undergo maturation in response to TSLP [96]. Unlike OX40L and TLR ligands, TSLP does not stimulate mDCs to produce the Th1-polarizing cytokines such as IL-12 or the proinflammatory cytokines (e.g. TNF-α, IL-1β, and IL-6) [96]. TSLP causes mDCs to produce large amounts of the chemokines such as IL-8 and Eotaxin-2, which attract neutrophils and eosinophils, as well as thymus and activation regulated chemokine (TARC) and macrophage-derived chemokine (MDC), which attract Th2 cells [96,101]. Liu [96] suggests that the inability of TSLP to induce the production of Th1-polarizing cytokines of mDCs is one of the most important features of TSLP-activated DCs, although the molecular mechanisms underlying TSLP’s ability to promote mDC maturation without inducing the production of Th1-polarizing cytokines are unknown. OX40L on DCs represents the original Th2-polarizing signal from TSLP-DCs, and IL-4 represents a critical autocrine stabilizer and enhancer of the developing Th2 cells. Thus, OX40L and IL-4 work synergistically and sequentially in driving Th2 responses in T cells [101,102]. Also, TSLP costimulates mast cells to produce inflammatory and Th2 cytokines. Also, TSLP is highly expressed by skin keratinocytes and airway epithelial cells during allergic inflammation. Although the triggering mechanisms of TSLP production by allergen or viral and bacterial infection are not well known, it appears to involve STAT5 activation, independent of the classic NF-κB and MyD88 signaling pathways. In the presence of IL-12, OX40L signaling instead promotes the development of Th1 cells. Liu [96] proposed the subdivision of Th2

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Page 6 of 10

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cells into inflammatory Th2 cells that produce high levels of TNF-a but little IL-10, and conventional Th2 cells that produce little TNF-abut high levels of IL-10. If that is the case, these mechanisms may operate in allergic rhinitis, although it is not well understood in nasal epithelial cells.

Future Direction

The mechanisms of goblet cell metaplasia, differentiation and hyperplasia are clarified by mainly in vitro study [103]. However, in vitro cultures do not always reflect physiologic conditions 

in vivo

[104,105]. The pathological changes in asthma affect the epithelium morphology and function. The constitute of bronchial mucosa such as ciliated epithelial cells, goblet cells, basal cells, basement membrane,fibroblast,smooth muscle, capillary, DC and bronchial gland (niche) may contribute to the development of asthma (Figure 4). The interaction of epithelial cells and underlying mesenchymal including inflammatory cell responses may affect or determine the fate of epithelial cells. Tang [106] pointed out that BCs proliferation and differentiation in vivo is still not known. Thus, it will be important to understand the difference between in vitro and in vivo, and how they are coordinated and how microenvironmental signals are integrated in metaplasia, hyperplasia and differentiation of goblet cells to direct homeostasis, repair and pathological remodeling of airways. Moreover, we have to remind that the data obtained from model mousenot simply applicable for human with asthma, since there are some species differences such as composition of epithelial cells and their function.

TSLP play a key role in goblet cell responses as the result of the development of asthma. TSLP is a cytokine produced by the skin and airway epithelium stimulated with environmental factors including TLRL that is capable of directing DCs towards an inflammatory Th2 response, thereby providing an essential link between epithelial cell activation and allergic-type inflammation. In addition, TSLP can interact directly with mast cells to initiate Th2 cytokine production to also provide a non-T cell route to mediate its pro-allergic effects. Holgate [107] pointed out that induction of TSLP production occurs through the activation of epithelial TLRL to provide an important new link between innate immunity and allergic disease. Moreover, it has been demonstrated that TSLP diverted airway tolerance against ovalbumin to Th2 sensitization and inhibited the generation of OVA-specific Foxp3 regulatory T cells [108]. Therefore, TSLP and OX40 ligand may represent important targets for intervention of the initiation of allergic inflammatory responses [107].

The asthmatic epithelium is known to be abnormally susceptible to apoptosis induced by oxidants, possibly due to a lack of endogenous protective factors [109,110]. Moreover, there is some evidence indicating the possible involvement of mitochondrial DNA defects in the etiology of asthma [110-112]. The recruitment of Th2 cells to the airway and the activation of those cells may produce ROS leading to mitochondrial damage and dysfunction by activated inflammatory cells [60,113,114]. Several oxidative stress studies revealed that endogenous antioxidant enzymes were decreased in the peripheral tissues of asthmatic adult patients relative to the control subjects, indicating that mitochondrial dysfunction and oxidative stress is present in asthmatic adults [115]. However the relation of those changes and mucus production in vitro and in vivo [105] is totally unknown. Thus, additional work is needed to fully answer this question for better therapy in the control of mucus hypersecretion.

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