Metabolism of the Extracellular Matrix in Bronchial Asthma (Review)

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Bronchial asthma is associated with upper airway (UA) disorders, primarily with allergic rhinitis, which, in turn, occurs in combination with other UA conditions, including hyperplasia of the nasal mucosa. Chronic rhinosinusitis, if confirmed, is a predictor of asthma severity.

The pathogenesis of these diseases includes the remodeling (restructuring) of the extracellular matrix and the adjacent UA structures, which is associated with further worsening of the diseases and their resistance to therapy. It is known that remodeling of the lower respiratory tract in bronchial asthma is characterized by epithelial desquamation, hyperplasia of goblet cells, thickening of the basement membrane, fibrosis of the subepithelium, hyperplasia of smooth muscles of the respiratory tract, and increased angiogenesis. At the same time, the UA remodeling in patients with asthma is still poorly understood; the data are still limited and often contradict each other. With isolated allergic rhinitis, the remodeling process is not very much pronounced and is limited, apparently, to a basement membrane thickening. In chronic rhinosinusitis, the UA remodeling manifests by epithelial hyperplasia and an increased sedimentation and degradation of the matrix along with the accumulation of plasma proteins.

Despite recent extensive studies, the cellular and molecular mechanisms involved in the respiratory tract remodeling remain largely undetermined, which necessitates further research into these processes. The review addresses several aspects of neuro-humoral control of the extracellular matrix metabolism and the associated remodeling of the upper and lower airway in patients with asthma.

Key words: bronchial asthma; allergic rhinitis; chronic rhinosinusitis; extracellular matrix; remodeling.

Introduction

Extracellular matrix (ECM) is a collective name for the extracellular tissue structures (basement membranes, interstitial matrix) that provide mechanical support for cells, perform signal conduction, participate in metabolism and transport, mediate the cell-cell contacts and cell locomotion. Components of the fluid connective tissues — blood plasma and lymphatic fluid — that mediate gas exchange and maintain homeostasis [1] are also attributed to the ECM.

Previously, ECM was considered an inert structure. Recent studies demonstrate though that ECM is a biologically active medium, crucial for the functioning of organs and systems, both in normal conditions and in disease [2]. Changes in ECM of the respiratory tract and/or lung parenchyma are considered key elements of bronchial asthma (BA) and the concomitant diseases of
reviews the upper airway (UA) [3–5]. It has been lately discussed whether changes in ECM are a consequence of these diseases or whether they are active elements of the pathogenesis that determines the clinical picture of the disease [6]. The most recent data suggest that changes in the composition and quality of ECM proteins may alter the functional characteristics of cells associated with the ECM [7].

Expanding the knowledge about the ECM in patients with BA and the associated UA disorders will contribute to a better understanding of respiratory chronic diseases and will help in a search for new approaches to their treatment [8].

**Structure and function of the extracellular matrix**

Extracellular matrix is a complex and dynamic structure that provides a mechanical framework for cells located inside it. ECM consists of a large number of macromolecules, mainly proteins and glycoproteins; their composition and structure are organ-specific [9].

There are two types of ECM: the first is represented by the basement membrane (BM), which lies directly beneath the epithelial and endothelial cells, and the second is the interstitial matrix of connective tissue surrounding the cells.

The BM functions as a supporting structure consisting of orderly arranged molecules, which adhere to endothelial and/or epithelial cells and protect them against biochemical and biophysical stresses [10]. The barrier function of the BM enforces the function of mucous membranes in the respiratory, gastrointestinal and urinary systems [9]. In addition, BM (or basal plates) represents a supporting structure for muscle and adipose cells, neurons and the Schwann cells of peripheral nerves [11]. These “leaf-like” layers appear at an early stage of embryogenesis; they separate tissues, function as macromolecular filters, and provide sites for cell adhesion [9]. By mediating the contacts between the epithelial membrane receptors and the BM proteins, BMs participate in cell arrangement and differentiation. In this respect, of particular importance are integrins — transmembrane heterodimeric cell receptors that interact with the ECM and transmit signals to the cells. On the basal side of the epithelial cells, hemidesmosomes — BM protein binding receptors — are expressed [9].

The BM consist mainly of type IV, XV, XVIII collagens, laminins, glycoproteins, and proteoglycans, which separate the epithelium or endothelium from the surrounding stroma [9] (Table 1). Laminins provide attachment of epithelial cells to BM and, in conjunction with type IV collagen and other components of BM, ensure its stability. The laminin networks are non-covalent and, therefore, more active than the collagen networks [12]. Both networks are linked via nidogen, which plays a role in their structure stabilization [13–17].

The interstitial matrix is also tissue-specific and includes, as a rule, a large number of fibrous proteins (collagen, elastin), fibronectin and proteoglycans, which are interconnected with other molecules and assembled into complex fibriillary networks [1]. The interstitial component of the ECM is needed for structural stability of tissues, cell migration, their proliferation and adhesion; in addition, it takes part in the control over synthesis of inflammatory mediators and the regulation of water balance [9, 18].

| Table 1 | Major structural components of the extracellular matrix ([1, 9] with additions) |
|---------|--------------------------------------------------------------------------------|
| **Class** | **The most common types of collagen** | **Description** |
| Fibrillary collagens | I, II, III, V, XI, XXIV, XXVII | Localized largely in tissues with a fibrous ECM, such as skin, bones, tendons, and ligaments |
| Network-forming collagens | IV | Localized in the basement membranes (among others). Have a mesh structure associated with laminins and other proteins of the basement membrane |
| Short-chain collagens | VIII, X | Involved in regulation |
| Fibril-associated collagens with intermittent triple helix | IX, XII, XIV, XVI, XIX, XX, XXI, XXII | Molecular bridges linked to type I (XII, XVI, XIX, XXI) collagen fibrils and type II (IX, XVI, XIX) collagen fibrils |
| Membrane-associated (transmembrane) collagens with intermittent triple helix | XIII, XVII, XXIII, XXV/CLAC-P | Cell membrane-associated molecules with transmembrane and intracellular domains |
| Long-chain collagens | VII | Anchor fibrils. Collagen-containing anchor fibrils, mainly associated with basement membranes |
| Collagen shaped as filaments-beads | VI | Soft tissues, cartilages |
| Multiplexin | XV, XVIII | Localized mainly in endotheliocytes. Multiple interrupted triple helical domains containing chondroitin sulfate and glycosaminoglycan heparin sulfate |
By now, 28 basic types of collagen have been described; they differ from each other by the amino acid sequence, as well as by the mode of hydroxylation and glycosylation. The common feature of all these collagens is the existence of triple helix domains, which are considered part of the ECM. More than 90% of total collagen of higher organisms is represented by collagens of types I, II, III, and IV.

In addition to collagens, there are many other proteins containing triple helix domains [1]. Nevertheless, those are not classified as collagens, but as collagen-like proteins. Examples of such proteins are adiponectin, collectins, and ficolins, the terminal structure of acetylcholinesterase, etc. These proteins play both a structural and a regulatory role [19].

Cells populating the ECM interact with this macromolecular network through their surface receptors, such as integrins, proteoglycans (localized on the cell surface). Major proteoglycans of the extracellular matrix are shown in the table below.

| Classes                  | Groups (types)                  | Molecules                              |
|--------------------------|---------------------------------|----------------------------------------|
| Extracellular proteoglycans | Hyalectans                      | Aggrecan                               |
|                          |                                 | Versican                               |
|                          |                                 | Neurocan                               |
|                          |                                 | Breviscan                              |
|                          |                                 | Small leucine-enriched proteoglycans    |
|                          |                                 | Decorin                                |
|                          |                                 | Biglycan                               |
|                          |                                 | Fibromodulin                           |
|                          |                                 | Lumican                                |
| Pericellular proteoglycans | Perlecan                        | —                                      |
|                          | Agrin                           | —                                      |
|                          | Barnacan                        | —                                      |
| Proteoglycans embedded in the cell membrane | Syndecans                      | —                                      |
|                          | Glypicans                       | —                                      |
|                          | Chondroitin sulfate 4 proteoglycan | —                                   |
|                          | Beta-glycans                    | —                                      |
|                          | Phosphacans                     | —                                      |
| Intracellular proteoglycans | —                              | —                                      |

Specialized non-collagen proteins of the extracellular matrix:

| Protein                          | Locations                            | Notes                                                                 |
|----------------------------------|--------------------------------------|-----------------------------------------------------------------------|
| Elastin and elastin-associated proteins | Arteries, connective tissue, respiratory system, lungs, skin, bladder | Provide structural integrity along with reversible extensibility and deformability |
| Fibronectin                      | Blood, connective tissue             | —                                                                     |
| Laminins                         | Basement membrane                    | —                                                                     |
| Matrikines                       | Connective tissue                    | Disulfide-linked proteins involved in the formation of filaments      |
| Fibrinogen                       | Blood                                | Soluble glycoprotein, a component of the blood coagulation system     |
| Fibrillin                         | Connective tissue, cardiovascular tissue | The predominant component of microfibrils; creates a shell around elastin of the elastic fibers |
| Fibulins                           | Basement membranes, blood vessels, blood | Calcium binding glycoprotein                                      |
| Netrins                           | Basement membranes                   | Laminin-bound proteins that are involved in axon control and vascularization of tissues, including the lungs |
| Osteopontin                       | Various tissues                      | Bone mineralization, cell adhesion and cell attachment                |
| Tenascons                        | Connective tissue                    | Glycoproteins that are involved in inflammation and fibrosis          |
| Vitronectin                       | Blood                                | Adhesive glycoprotein associated with coagulation and wound healing   |
surface), the hyaluronan CD44 receptor and other molecules [20, 21]. This way the cells communicate with the ECM; these signals largely determine their function and behavior. Various growth factors, cytokines, and chemokines bound to specific ECM molecule are found within the ECM; eventually, these regulatory molecules are released and activated under specific physiological conditions. Changes in the composition and structure of individual ECM components affect the structural and functional parameters of the entire network, as well as the numerous cellular elements associated with ECM [22, 23]. Notably, all the ECM cells, including epithelial and endothelial, immune cells, fibroblasts, and smooth muscle cells participate in the synthesis and secretion of matrix macromolecules, thus maintaining the ECM structure and function. The term “matrikines” has been proposed for the bioactive ECM fragments that modulate various physiological processes. It has been shown that some matrikines — metabolites of ECM components — are involved in the mechanism of inflammation and immune homeostasis [24, 25].

There are reports on syndromes and diseases caused by abnormalities in the synthesis and metabolism of ECM components; those are currently considered as potential targets for pharmacological therapy [26].

**Structure of the extracellular matrix in the respiratory system**

Respiratory organs perform two vital physiological functions: passive gas exchange (alveolar respiration) and immune protection against exogenous antigens (by virtue of the epithelial barrier). The ECM of the respiratory system is saturated with elastin. The ECM structure often changes in response to environmental factors detrimental to the airway epithelium, including chronic exposure to inhaled allergens, cigarette smoke, pollutants, and infections [27]. Damaged epithelial cells can provoke a specific ECM remodeling, which, in turn, affects the epithelial cells themselves [28].

The respiratory system has a unique composition; the ECM maintains permeability of the airway, and gas exchange in the lung parenchyma. In the proximal airway segments, the ECM contains a large amount of collagens, laminins, and proteoglycans [29]. In the alveoli, type I epithelium prevails; its cell walls almost merge with endothelial cells of proximal capillaries, creating an ultrathin elastin-dominated ECM to ensure effective gas exchange [29].

The nose and paranasal sinuses provide the conditioning of the inhaled air, and the ECM and the epithelial UA lining protect the lower airways from antigens and pollutants [30]. The UA epithelium has a predominantly pseudo-layered columnar structure, and the supporting connective tissue is largely loose. In BA and comorbidities, abnormalities of the UA extracellular matrix manifest in different ways (from fibrosis to severe edema), which is due to extensive vascularization of the UA [31].

**Metabolism of the extracellular matrix**

Recent studies show that ECM is a highly active tissue that is constantly being reconstructed, thus responding to the needs of growth and repair [32].

In the body, ECM undergoes continuous degradation mediated by special enzymes, accompanied by simultaneous synthesis and restructuring of matrix components. Changes in the ECM affect the surrounding cells so to regulate their proliferation, migration, or differentiation [33]. Normally, ECM homeostasis is characterized by an optimal balance between the formation, secretion, alteration, and degradation of the matrix [34]. The local degradation of ECM is a prerequisite for cell migration and proliferation.

The restructuring of the ECM is called “remodeling”. This term refers to structural and geometric changes in the ECM and has both a physiological and a pathogenetic meaning. In a healthy person, remodeling occurs from the birth through the maturity as an adaptive response to the body growth. Abnormal remodeling of ECM may reflect a disease, for example, chronic obstructive pulmonary diseases [33].

The proteolytic degradation of ECM components is mediated by enzymes, including matrix metalloproteinases (MMPs); disintegrins and metalloproteinases (ADAMs — a disintegrin-like and metalloproteinase domain); ADAMs with thrombospondin motifs (ADAMTS — ADAMs with thrombospondin type I motifs); enzymes of the meprin family; serine/threonine proteases (elastase, matriptase, urokinase activator and tissue plasminogen); cysteine proteases; aspartate proteases [35–37] and other enzymes [38].

The enzymes of the ADAMS group mainly cleave the transmembrane protein ectodomains associated with the cell membrane. This is accompanied by a release of cytokines, growth factors, receptor and adhesion activation [33, 39].

The meprins are proteases that cleave ECM proteins, such as collagen IV, nidogen and fibronectin [40]. Meprins can also contribute to the formation of true collagen molecules by cleaving procollagen I, which is then assembled into collagen fibrils [41]. Indirectly, meprins control ECM remodeling by activating other metalloproteinases [33].

Excessive degradation of ECM leads to tissue destruction. At the same time, excessive production of ECM components and their precipitation, which occur in response to severe tissue damage, can cause fibrosis in the absence of adequate degradation of ECM components [14].

**Remodeling of the extracellular matrix in bronchial asthma**

Airway remodeling in BA can be defined as an advancing pathological reorganization of their cellular and molecular structure. The onset and progression of
the structural changes remain the subject of discussion; nevertheless, the negative impact of the airway remodeling on the respiratory system is now generally recognized [42].

In BA, the airway remodeling likely results from the bronchial tree inflammation caused, among other factors, by exposure of sensitized patients to allergens [43]. The airway remodeling in asthma can be caused by a mechanical stress or triggered by earlier life events [44, 45].

Nevertheless, the linear model suggesting that an environmental sensitization leads to allergic inflammation (mediated by Th2 cells) and subsequent airway remodeling, has been questioned for asthma in pediatric patients. Bronchial biopsies in children with asthma show pronounced remodeling at an early stage of the disease, thus implicating the allergic component of asthma rather than chronic airway inflammation [46, 47]. Elliot et al. demonstrated that the mechanisms of airway remodeling in BA could be associated both with the inflammation and/or with non-inflammation processes [48]. There is a debate about whether the inflammation precedes the remodeling and whether it can be associated with primary anomalies (e.g., atopy) of the epithelial recovery [49], or the remodeling does not depend on inflammation and can occur before clinical manifestations of BA [50].

Remodeling of the airways in asthma includes large-scale structural changes, which lead to a narrowing of the airways, increasing their resistance and retaining mucus in the bronchial lumen. Remodeling of the airways in asthma also includes thickening of their walls, hyperplasia and hypertrophy of the bronchial smooth muscles (BSM), edema, subepithelial fibrosis, quantitative and qualitative changes in the ECM, remodeling of the ECM subepithelial space, loss of the barrier function of the BM, accumulation of immune cells and fibroblasts, angiogenesis, epithelial metaplasia, loss of cilia, and mucous hypersecretion [51]. The deposition of ECM in the subepithelial area may play an important role in modulating the structure and function of the epithelial cells and fibroblasts, since these deposits are in close contact with the epithelial cells and, moreover, they all form a functional epithelial-mesenchymal-trophic unit (EMTU).

In BA, the characteristic feature of the airway remodeling and structural changes in the ECM is the BM thickening described by Huber and Koessler as early as in 1922 [52]. At present, it is known that the BM thickening in patients with asthma is accompanied by the accumulation of collagen types I, III, IV, and fibronectin [5]. For example, in healthy adults, the thickness of BM reticular plate is 5–6 µm, while in patients with chronic BA it is 9 µm [27].

In asthma, ECM structural abnormalities are observed in all parts of the respiratory system (central respiratory tract, bronchial smooth muscles, distal parenchyma, and blood vessels). Manifestations of ECM remodeling depend on the severity of asthma, the degree of disease control, the patient's age, and the pharmacotherapy [27].

Structural changes in ECM in asthma can be accompanied by a loss of some elastin fibers, which, coupled with a gradual decrease in lung function, can lead to a loss of the airway elasticity and, consequently, to the formation of air traps in the lungs [53].

Poon et al. [54] suggested that the angiogenesis and neovascularization of the mucous membranes are key factors in the airway remodeling in severe asthma. It was also demonstrated that the deposition of proteoglycans or collagen in the airway walls can contribute to bronchial hyperreactivity [55, 56].

An important role in the initiation of ECM rearrangement is played by the epithelium [57] (Table 2). For example, in asthma, repeated damage to epithelial cells by exogenous antigens and pollutants (in combination with immune factors) contributes to recurrent epithelium desquamation. This leads to permanent activation of the EMTU to maintain reparative processes that may be accompanied by prolonged and progressive remodeling of the airways [58, 59].

Another effector of the airway remodeling in asthma is the BSM, which, (in patients with BA) actively secretes ECM components, cytokines and growth factors. BSM itself is linked to the ECM structure and thereby able to increase its contractile

| Table 2 Participation of various cell types in the pathological airway remodeling in bronchial asthma ([60] with additions) |
| Cell types | Role in the airway remodeling |
|---|---|
| **Epithelial cells** | Desquamation of epithelial cells | Affects mucus secretion |
| | Subepithelial fibrosis | Hyperplasia of the mucous glands |
| | Stimulating smooth muscle cell proliferation in the bronchi through the release of growth factors | Recruitment of pro-inflammatory cells |
| | Accelerates sediment deposition in the extracellular matrix | Stimulates angiogenesis and neo-angiogenesis |
| **Bronchial smooth muscle cells** | Bronchial smooth muscle hypertrophy that narrows the airway lumen | Migration of bronchial smooth muscle cells and their invasion into the epithelium |
| | Transformation of the bronchial smooth muscles into the “synthetic phenotype”: results in excessive secretion of transforming growth factor beta (TGF-β), chemokines, and extracellular matrix components | Interaction with immune and other cells mediated by cell adhesion molecules |
| **Fibroblasts** | Differentiation into myofibroblasts, secretion of extracellular matrix components | Accumulating in the subepithelial layer |
potential under ECM remodeling caused by airway inflammation [61, 62]. A BSM mass gain in combination with a BM thickening results in a narrowing of the airway lumen [42, 51].

It was found [63] that bronchoconstriction alone, without an inflammation, could induce airway remodeling in patients with BA. In this regard, it is an assumed that the mechanical forces developing in the airway walls during bronchoconstriction were those that triggered the remodeling [64].

It has been experimentally demonstrated [65] that the inflammation correlates with the increase in BSM contractility; the effect may be mediated by pro-inflammatory cytokines such as tumor necrosis factor. This process also involves an increase in the number of contractile units (active actin + myosin complexes). As a feedback, signals from the altered BSM can induce both the remodeling and the inflammation [66, 67]. It is likely that both the inflammation and the remodeling can develop rapidly if caused by a viral infection, exposure to allergens, or even by excessive BSM contraction [63, 68]. All this underscores the need for additional research into the airway remodeling [51].

Fibroblasts and myofibroblasts contribute to the process of remodeling by excessively secreting ECM components. In addition to increasing the airway wall thickness, ECM components can modulate cell proliferation and migration. There is evidence [69] that increased expression of hyaluronic synthase 2 in myofibroblasts and smooth muscle cells leads to an increase in airway fibrosis. Thus, it is quite possible that the accumulation of hyaluronan participates in the airway wall thickening in asthma via this pathophysiological mechanism.

It is known that airway remodeling reduces the efficacy of bronchodilators [5, 32–34]. A correlation was established between the degree of airway remodeling and the severity of asthma, but the clinical implications of this relation are not fully understood [35, 70]. The insufficient knowledge impedes the development of pharmacotherapy targeting the asthma manifestations associated with the airway remodeling. Thus, it is important to assess the current therapeutic strategies in asthma for their abilities to counteract the pathological airway remodeling [51].

Airway remodeling can also be observed in clinical situations that are not accompanied by pronounced BA symptoms; for example, in allergic rhinitis or in elite athletes [71].

Possible adverse effects of airway remodeling in asthma include a reduction in the lung function, an irreversible airway obstruction, a decrease in the airway distensibility and the reaction to bronchodilators, a persistent hyper-reactivity, a decrease in the BSM ability to relax, a decrease in the lung elasticity, and an increase in the BSM contractility caused by its hypertrophy [72–74]. There are also positive consequences of the airway remodeling in asthma: for example, the airways become better protected against a bronchospasm due to the increased rigidity of their walls [75].

**Metabolism of the extracellular matrix in asthma patients of various ages**

The updated evidence suggests that the most of the airway remodeling in BA occurs before the onset of clinical symptoms or at an early age [76–78]. Thus, according to biopsy data, in preschool children with wheezing, the BSM and the reticular layer of the BM are thickened [79]. Owens et al. [80] showed that a decrease in the lung function in infancy was a predictor of asthma in young adults, indicating the likelihood of a very early (possibly intrauterine) structural change in the airways. It is assumed that the remodeling of BSM and the reticular layer of BM is possible at early stages of asthma and may be independent of a concomitant inflammation [76]. Age-dependent changes in the lung elasticity are also important determinants of the lung function; however, these changes are yet to be characterized in detail [3, 81, 82].

**Neurohumoral regulation of metabolism of the extracellular matrix in asthma and related diseases of the upper airway**

ECM metabolism and remodeling processes are controlled by neuroendocrine signals; studies on this mechanism are currently under way. Particular attention is paid to the effect of glucocorticoids on the process of remodeling because these agents are commonly used in the anti-inflammatory treatment of asthma and associated UA diseases.

It is well documented that inhaled glucocorticoids help restore the integrity of the epithelium, prevent vascular changes and the deposition of ECM components of in the airways. The glucocorticoids can affect the ECM through various mechanisms, including the effect on the genes encoding for ECM components [83, 84].

In recent years, close attention has been paid to the relationship between the nutritional status of patients and the course of asthma and associated UA diseases. We previously [85–87] showed that the severity of asthma is associated with the physical body development and nutritional status of patients. In this regard, it is also of great interest to assess the effect on the ECM of hormones regulating the energy exchange and adipose tissue metabolism. Williams et al. [81] found that the synthesis of metalloproteinases by fibroblasts is controlled by leptin. The stimulation of fibroblast by leptin can change the metabolism of ECM components, including fibrillar collagens. Zhang et al. [88] showed that leptin reduced the expression of matrix metalloproteinases 2 and 9 that catalyze the degradation of collagen, and simultaneously increased the production of the inhibitor of metalloproteinases 1 and collagens I and IV. These studies indicate that in
the presence of leptin, the collagen synthesis dominates over its degradation.

In the experiment, adiponectin showed the exact opposite effect (increased expression of matrix metalloproteinases 2 and 9, and a decrease in TIMP-1 protein and collagens), but its effect manifested only in the presence of leptin. Notably, the effect of leptin on the ECM is not only a direct interaction with the synthesis of structural components, but it is also mediated by the nervous system. Leptin dilates the airways by reacting with the M3 receptors on the smooth muscles, regardless of inflammation in the respiratory tract, and thus modulates the autonomic control of the airways [89].

It has been demonstrated that a thyroid function insufficiency leads to excessive accumulation of hyaluronic acid in the ECM and also the development of myxedema [89]. The effect of the T3 hormone, like that of hydrocortisone, involves an inhibition of the hyaluronan synthesis; however, these two hormones may act through different pathways: 1) decreasing the synthesis of hyaluronan; 2) enhancing its catabolism, mediated by hyaluronidase. The effect of T3 is probably carried out through a decrease in the production of hyaluronate synthase 2. It should be noted that T3 does not affect the synthesis of collagen or chondroitin sulfate [90].

The surface receptor for iodothyronines is represented by αV/β3 integrin, which is a heterodimer that interacts with a large number of ECM proteins. Thyroid hormones, by binding to αV/β3, modify the expression of the vascular endothelial growth factor, promote angiogenesis, cell proliferation and cell migration [91].

As mentioned earlier, BSM cells are actively involved in ECM remodeling. The BSM involvement is thought to be controlled by the paracrine system. This mechanism where the BSM contractility is regulated by distant organs can help understand the pathogenesis and heterogeneity of asthma [59, 92]. The effects of hormones on the BSM are multidirectional. Thus, the activation of the BSM leptin receptor inhibits the proliferation and migration of BSM due to stimulation of prostaglandin E2 secretion [93]. A negative correlation has been found between the BSM mass and the serum level of vitamin D in children with severe asthma resistant to therapy [94].

Insulin stimulates BSM hyper-contractility, possibly due to an increase in the free insulin-like growth factor, which is associated with BSM proliferation [95].

Sex steroids have a dual effect on ECM and BSM in patients with BA. Thus, high doses of estrogen increase the contractility of BSM ex vivo and reduce the lung function in vivo, while the estrogen replacement therapy reduces the airway reactivity and improves the lung function [96, 97].

Androgens are able to relax contracted BSM and potentiate the relaxing effect of β2-agonists, which suggests their overall bronchodilation effect [98, 99]. On the other hand, in the same preparations of BSM and at the same concentrations, androgens potentiate the contractile response to the spasmogen [99].

Inter-hormonal interactions and their impact on the ECM are also of considerable interest. In experiments of Ishida-Takahashi et al. [100], glucocorticoids inhibited the effects of leptin-mediated by the transcription factor pSTAT3. STAT3 is a signaling protein and a transcription activator from the STAT protein family, which is encoded by the human STAT3 gene. STAT3 is one of the intermediary proteins participating in the cell’s response to signals via interleukin receptors and growth factors.

Neuro-regulation of ECM in patients with BA is the best studied in terms of cholinergic receptors. Stimulation of muscarinic receptors leads to proliferation and remodeling of BSM, proliferation of human lung fibroblasts, synthesis of collagen and matrix metalloproteinases 1 and 2 [101–104].

The muscarinic effects on the ECM are expressed by enhancing the contractile potential of human BSM cells [104]. A few recent reports described the effects of the M3-choline blocker tiotropium bromide, a prolonged action drug [105]. Tiotropium bromide inhibits the remodeling of the bronchial wall, mitigates the wall thickening, the mucous gland hypertrophy, and the smooth muscle hyper-reactivity, and also reduces the level of Th2 cytokines and pulmonary eosinophilia caused by an allergen [106]. Part of this pharmacological activity of tiotropium is the inhibition of TGF-β-induced expression of metalloproteinases in human lung fibroblasts [107]. Tiotropium also inhibits the increased peribronchial collagen deposition in a model of chronic swine flu [108].

Remodeling of the upper airway in patients with bronchial asthma

Between 80 and 100% of patients with atopic asthma were also diagnosed with concomitant allergic rhinitis, which necessitated a closer look at their UA [109–113]. Recent studies demonstrate that the inflammation processes in the mucous membrane of the upper and lower airways in patients with asthma are identical and likely associated with the Th2-dependent mechanism [114, 115].

Assuming the remodeling is caused by inflammation, there must be detectable structural changes in the UA of patients with asthma due to the persistent inflammation caused by allergic rhinitis. However, no clear evidence of any rhinitis-induced remodeling has been provided in the literature [5].

Both in seasonal and year-round allergic rhinitis, biopsy specimens of the nasal mucosa (taken within first hours after exposure to significant allergens) showed an increased thickness of BM as compared to that in healthy subjects. However, 24 h after the exposure, despite the influx of eosinophils, no further BM thickening occurred [116, 117]. Apparently, remodeling of UA is not a key feature of the immunopathogenesis of allergic rhinitis in patients with asthma.
According to our earlier studies, isolated allergic rhinitis occurs only in part of children with asthma [112, 113]. In most of these patients, we found a combination of rhinitis and other UA disorders, including abnormal changes in the intranasal structures and hypertrophy of the pharyngeal tonsil. In some adolescent patients, there were hypertrophic changes in the nasal mucosa, which might indicate a possible debut of polyposis/rhininosinusitis [110, 111, 118].

It is known that chronic rhinosinusitis (CRS) and BA are closely related, especially in adult patients [119, 120]. CRS is a broad term for a group of heterogeneous inflammatory diseases—it is classified into CRS with or without nasal polyps [117, 121]. Up to 14% of patients with CRS-associated polyps and patients with aspirin-induced respiratory diseases were also diagnosed with severe asthma [120, 122].

Unlike allergic rhinitis, tissue remodeling in CRS is a hallmark of this condition. In CRS with polyps, the epithelial cells of the nasal mucosa degrade; under this condition, the excess mucus accumulation can be explained by hyperplasia of the mucous glands and hypersecretion of mucin [123, 124]. The BM thickening reflects the severity and duration of the disease and the concomitant asthma, which, apparently, does not depend on the degree of eosinophilic inflammation [125, 126].

The degradation of ECM is a key pathological change in CRS with polyps; in addition, there are changes in the tissue architecture and growth, including the occurrence of pseudocysts (Figures 1, 2).

By now, the available descriptive studies have not yet provided a detailed understanding of ECM remodeling in patients with CRS [5]. The mucosa ECM remodeling is not a characteristic feature of allergic rhinitis, except for some thickening of BM. In CRS with polyps and in asthma, the epithelial cells look damaged and the goblet cells show hyperplasia. In both cases, EMTU activation is noted. Perhaps, the original epithelial vulnerability plays a role in this mechanism, leading to submucosal mesenchymal activation. As for the submucosa, the remodeling processes in CRS with polyps and in allergic BA include hypertrophy of the mucous glands and excessive generation of ECM components. However, there are significant differences: pseudocysts in the nasal and paranasal mucous membranes are observed only in patients with CRS with polyps, but not in the mucous membrane of the lower airways in asthma [5].

**Conclusion**

According to the current concepts, the extracellular matrix is a biologically active medium that impacts the course and outcome of diseases. The restructuring (or remodeling) of the extracellular matrix is a pathogenetic mechanism that underlies a number of clinical conditions, including bronchial asthma and concomitant diseases of the upper airway.

Airway remodeling is characteristic of bronchial asthma; it starts at early stages of the disease and may reflect previous life events and/or persistent airway inflammation. The restructuring of the extracellular matrix in patients with asthma is an active and very complex process involving multicomponent changes in the lower airway, such as epithelial desquamation, hyperplasia of the goblet cells, thickening of the basement membrane, subepithelial fibrosis, hyperplasia of smooth muscles of the bronchi, and increased angiogenesis.

At the same time, in patients with asthma combined with isolated allergic rhinitis, the remodeling is limited just to thickening of the basement membranes despite the severe Th2-dependent mucosa inflammation.

However, some patients demonstrate hyperplasia of the nasal mucosa, which apparently leads to the development of chronic rhinosinusitis with or without polyps. These processes are characterized by gross
changes in the extracellular matrix, which are yet to be specified.

With regard to the pathological processes caused by the extracellular matrix remodeling, an efficient therapeutic approach is yet to be proposed. Further research into the tissue remodeling in the upper and lower airways in patients with bronchial asthma and concomitant upper airway diseases is needed to elucidate their pathogeneses and develop therapeutic strategies for the treatment of allergic diseases and clinical consequences of airway remodeling.

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