

Chitinases and Chitinase-like proteins in asthma

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ABSTRACT

Despite the lack of endogenous chitin synthesis, mammalian genomes encode two enzymatically active true chitinases (chitotriosidase and acidic mammalian chitinase) and a variable number of chitinase-like proteins (CLPs) that have no enzyme activity but bind chitin. Chitinases and CLPs are prominent components of type-2 immune response-mediated respiratory diseases. However, despite extensive research into their role in allergic airway disease, there is still no agreement on whether they are mere biomarkers of disease or actual disease drivers. Functions ascribed to chitinases and CLPs include, but are not limited to host defense against chitin-containing pathogens, directly promoting inflammation, and modulating tissue remodeling and fibrosis. Here, we discuss in detail the chitin-dependent and -independent roles of chitinases and CLPs in the context of allergic airway disease, and recent advances and emerging concepts in the field that might identify opportunities for new therapies.

41 **Abbreviations:**

42	AM	alveolar macrophage
43	AAM	alternatively activated macrophage
44	AMCase	acidic mammalian chitinase
45	BAL	bronchoalveolar lavage
46	CHIT1	chitotriosidase
47	CLP	chitinase-like protein
48	COPD	Chronic obstructive pulmonary disease
49	DAMP	damage-associated molecular pattern
50	DC	dendritic cell
51	GH18	family 18 of glycosyl hydrolases
52	GINAc	N-acetyl-glucosamine
53	HA	hyaluronan
54	HDM	house dust mite
55	IgE	immunoglobulin E
56	OVA	ovalbumin
57	PAMP	pathogen-associated molecular pattern
58	PRR	pattern recognition receptor
59	WT	wild-type

Introduction

Despite the lack of endogenous chitin synthesis, mammalian genomes encode two enzymatically active true chitinases, chitotriosidase (Chit1; encoded by *CHIT1* in humans/*Chit1* in mice) and acidic mammalian chitinase (AMCase; encoded by *CHIA* in humans/*Chia1* in mice) that bind and degrade chitin, in addition to a variable number of chitinase-like proteins (CLPs) that have no enzyme activity but bind chitin (Table 1) (1). Chitinases belong to the evolutionary conserved family 18 of glycosyl hydrolases (GH18), a gene family found in all six kingdoms, that catalyse the hydrolysis of chitin to simple sugars (2). They are produced for different purposes ranging from morphogenesis and nutrition to immune defence against chitin-containing pathogens (2-4).

Besides their potential host defense function in mammals (5), chitinases and CLPs have not only been associated with type-2 immune response-mediated respiratory diseases, but proposed as potential biomarkers and drivers of allergic asthma (4, 6). Allergic asthma is a Th2-mediated, chronic inflammatory disease of the lung that is triggered by the exposure to allergens and associated with aberrant expansion of Th2 lymphocytes, airway eosinophilia, elevated serum immunoglobulin E (IgE), and excessive airway mucus production. Non-allergic asthma also exists, some forms also associated with eosinophilia driven by innate type-2 lymphocytes, and others with more neutrophilic inflammation or little immune cell influx (7). While asthma prevalence has increased to epidemic proportions over the last few decades, the development of targeted therapies to treat this debilitating disease has not kept pace, and the economic burden of disease morbidity and control continues to escalate. Recent years have seen renewed interest in chitinases and CLPs as critical regulators of type-2 immunity not only in response to chitin-containing allergenic sources (i.e., house dust mites (HDM)) but also in chitin-independent processes driving type-2 immunity (3, 8). This is supported by the discovery of polymorphisms and major variations in the expression of the chitinase enzymes and CLPs in the human population which have been correlated with enhanced/decreased activity, reduced lung function and increased susceptibility to bronchial asthma (4, 9).

88 Here, we discuss the chitin-dependent and -independent roles of chitinases and
89 CLPs in the context of allergic airway disease and recent advances and emerging
90 concepts in the field that might identify opportunities for new therapies.

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Chitin

Chitin is an insoluble, linear polymer of N-acetyl-glucosamine units, which are linked by $\beta(1,4)$ -bonds. It is structurally similar to cellulose, except that one hydroxyl group of each subunit is replaced by an N-acetyl group. The resulting increased hydrogen bonding between separate polymers makes it one of the strongest and most resilient natural materials on earth. In contrast to α -chain carbohydrates such as the readily digestible glycogen and starch, these β -linked polymers are highly resistant to degradation. Chitin is the second most abundant polysaccharide on earth after cellulose, and is produced by fungi, nematodes, and arthropods (such as crustaceans, insects and house dust mites). Due to its physical properties, it provides excellent structural rigidity to these invertebrates, just like cellulose provides structural rigidity to plants.

Presumably, chitin is the evolutionary precursor of hyaluronan (HA), the major component of the extracellular matrix of many vertebrates. HA is composed of repeating disaccharides containing N-acetyl-glucosamine and glucuronic acid, which are connected exclusively by β -bonds, like chitin and cellulose. Despite having similar structures, chitin and HA considerably differ in physical properties. While chitin excludes water, HA attracts large volumes of water, allowing cells to travel through the extracellular matrix. In humans, chitin synthase genes have been lost, leading to the assumption that mammals do not produce chitin. However, this view has recently been challenged by the observation that short chitin oligomers, remnants of chitin synthesis, are produced during the initiation of HA synthesis in vertebrates (10). In addition, chitin-like polysaccharides in Alzheimer's disease brains have now been documented (11). *In vitro*, HA synthases make small chitin oligomers that potentially serve as endogenous primers for the initiation of HA synthesis (10). These short chitin oligomers might act as a hydrophobic needle that penetrates the cell plasma membrane, dragging the hydrated HA chain into the extracellular space to allow its unrestrained growth.

Chitin is considered non-self and as described above, it has a structure analogous to HA, which can be recognized as a damage-associated molecular pattern (DAMP) by TLR4 (12), suggesting that chitin might also be recognized by the innate immune system as a pathogen-associated molecular pattern (PAMP). Indeed, chitin has been shown to

activate the mammalian immune system. However, as there is no consensus yet on which pattern recognition receptors (PRR) sense chitin, chitin is considered an “orphan” PAMP. Several PRRs have been implicated in chitin signalling in immune cells (Figure 1), such as TLR2 (13-15), TLR9 and NOD2 (16), mannose receptor (14, 17), and dectin-1 (14, 18). However, direct receptor-chitin interactions have only been firmly demonstrated for TLR2 (15). That study also showed that six subunit long chitin chains were identified as the smallest, immunologically active motif. Chitin oligomers of 5 or less subunits are immunologically inactive, and even inhibit the binding of larger chitin particles to TLR2 in a dose dependent manner; potentially this short conformation fails to induce cross-linking of receptors.

The fact that small chitin oligomers are immunologically silent could be evolutionary beneficial, allowing the distinction between self and non-self, as endogenous chitin oligomers are produced during HA synthesis. In addition to TLR2, other direct chitin-receptor interactions have also been described. The transmembrane receptor FIBCD1 binds chitin (19), but is mainly expressed in the intestinal epithelium, limiting its relevance for chitin sensing in the lungs. In contrast, LYSMD3 is expressed on airway epithelial cells and also directly binds chitin (20). Chitin binding to LYSMD3 induced receptor signalling with consequent cytokine secretion, which could be reversed by knockdown or knockout of LYSMD3. However, further studies are required to determine its role *in vivo*. Finally, chitin has also been shown to directly activate the alternative complement pathway (21).

The immune responses induced by chitin are rather complex and depend on the chitin particle size (Figure 1). Initial studies focused on chitin sensing and uptake by macrophages (22). For macrophages to engulf chitin, chitin polymers need to have the appropriate size to be phagocytosed and to result in a proper cytokine response (14, 23). Phagocytosis-dependent immune responses only occur with small chitin, and have been suggested to be dependent on the mannose receptor (14, 16). In the absence of the mannose receptor, chitin can still be phagocytosed, however this phagocytosis does not induce a cytokine response. There is some controversy about the phagocytosis-induced cytokine response. While Da Silva et al. report that phagocytosis induces TNF α secretion (14), Wagener et al. observed inhibition of TNF α secretion and induction of IL-10 secretion

(16). Next to phagocytosis, chitin recognition by PRR might activate macrophages to secrete type-1 cytokines with consequent IFN- γ production by NK cells (13, 14, 17, 22). TLR2 and consequent NF- κ B pathway activation, and dectin-1 and Syk activation have been implicated in chitin sensing by macrophages (13, 14).

These early reports mostly analyzed type-1-biased immune responses initiated by macrophages. Importantly, the majority of these studies were done *in vitro*, and might not reflect the *in vivo* situation. Indeed, chitin elicits much more complex immune responses *in vivo*. In 2007, Reese et al. reported for the first time that chitin administered to mouse lungs induced the accumulation of IL-4-expressing innate immune cells in tissues, including basophils and eosinophils (24). They found that chitin induced alternatively activated macrophages (AAMs) that produce leukotriene B₄, leading to eosinophil recruitment. Macrophages fail to acquire an AAM phenotype upon *in vitro* stimulation with chitin and instead secrete TNF α (25), suggesting an additional factor is required for *in vivo* AAM polarization. Roy et al. found airway epithelium-derived CCL2 to be needed for driving AAM activation (25). Indeed, more recent studies have shifted focus from macrophages as chitin sensors to the airway epithelium. These studies show that inhalation of chitin induces different patterns of epithelial cytokine secretion leading to a mixed eosinophilic-neutrophilic infiltrate in the airways (26, 27). Van Dyken et al. demonstrated that chitin triggers IL-33, IL-25, and TSLP secretion by the epithelium, which non-redundantly activate IL-5/IL-13-secreting ILC2 and eosinophil recruitment (26). At the same time, chitin also induced the expression of IL-1 β , TNF α , and IL-23, which lead to IL-17A production by $\gamma\delta$ T cells, and consequent neutrophilia. Of note, chitin induced epithelial responses might be influenced by macrophages. Kim et al. demonstrated that chitin phagocytosis by macrophages resulted in the activation of caspase-7, which inactivates epithelial-derived IL-33 thereby resolving type-2 immune responses (23).

It is striking that most chitin-containing organisms are known to trigger eosinophil-rich type-2 immune responses in mammalian hosts, such as the anti-parasite response against gastro-intestinal nematodes and the allergic response against allergenic sources, such as HDM, cockroaches, and fungi. Moreover, asthma is a common occupational hazard among people working with chitinous substances, such as shellfish processors

(28, 29). These observations have led to the hypothesis that chitin might be implicated in the pathogenesis of allergic diseases. The induction of chitin-binding antibodies has been described in mice (30). Immunization of mice with certain bacteria, such as *Streptococcus pyogenes*, induced anti-N-acetyl-glucosamine (GlcNAc) antibodies which were shown to bind chitin. However, these antibodies were rather protective, instead of contributing to allergic airway disease, as chitin-induced airway inflammation was greatly reduced by administration of anti-GlcNAc antibodies, suggesting that chitin stimulates airway inflammation. The current view is that the development of asthma and allergic inflammation involves innate immune responses that promote allergic sensitization, and chitin might act as an adjuvant in these responses. In animal models of airway sensitization, chitin was proven to act as an adjuvant for adaptive allergic responses elicited by ovalbumin (OVA) and by *Aspergillus* antigens (21, 31, 32). Chitin induced allergen-specific Th2 cells and allergen-specific antibodies. Complement activation and IL-33-induced DC activation have been implicated in the adjuvant effect of chitin (21, 31). Yet, the exact molecular pathways remain to be elucidated. As B and T lymphocytes might express TLR2 and other PRRs, it is plausible that they might be directly activated by chitin. Except for one report showing chitin to be a polyclonal B cell activator (33), the fine molecular details remain to be elucidated.

In contrast to the studies described above, chitin has also been reported to downregulate allergic features in mouse models. Intranasal application of chitin before and during allergen challenges reduced inflammation in the lungs and systemic IgE levels in some models of intraperitoneal sensitization (34, 35). Two other studies administered chitin orally and found that this reduced allergic features in a model of peanut-induced anaphylaxis and in a model of intraperitoneal sensitization to ragweed allergen followed by intratracheal challenges (36, 37). These findings are seemingly at odds with studies demonstrating the adjuvant effect of chitin.

In conclusion, chitin can elicit a broad range of innate immune responses, leading to the initiation of/or influencing adaptive immune responses. Yet the underlying molecular pathways are not fully understood. Conflicting observations may result from different

- 211 routes and timing of sensitization and chitin administration, different chitin preparations
212 (degree of acetylation, size and endotoxin content), and experimental design.

Chitinases

Structurally, members of the chitinase family consist of a catalytic domain, a hinge region and a chitin-binding cleft (38). These enzymes catalyse the cleavage of chitin β -bonds, consequently resulting in the depolymerization and degradation of chitin (38). Chitin bearing organisms express various chitinases for the remodelling of their complex chitin structures (38). While chitin synthase genes have been lost during evolution, chitinase genes were highly conserved across species with multiple duplication events (39). In mammals, two active endochitinases are expressed: Chit1 and AMCase. These proteins resulted from an ancient gene duplication event (39). While AMCase is very acid-stable and has a pH optimum of 2, Chit1 is most active at a more neutral pH (40).

Considerable differences exist between mice and men in the expression of active chitinases across tissues. It is well established that AMCase mediates chitinase activity in the mouse lung and that its expression is increased by an IL-13/STAT-6-dependent pathway, resulting in increased chitinolytic activity during allergic inflammation (41, 42). Chit1 levels are very low to absent in the mouse lung. In contrast, in humans, chitinolytic activity in the airways is exclusively attributed to CHIT1 (43). Although AMCase is expressed in the human lung, it is an inactive splice variant which has no chitinolytic activity (43). Active AMCase is expressed in the stomach of humans (43). Because of expression in the barrier tissues of the airways and gastro-intestinal tract of mammals, it is plausible that chitinases confer protection against chitin-bearing organisms. However, chitinases seem to have more complex functions, and might either contribute to or inhibit inflammation depending on the context. Extensive research has been conducted on the potential value of chitinases as biomarkers and therapeutic targets of airway inflammation and remodelling, as discussed below.

Chitotriosidase

CHIT1 was the first active chitinase to be described in humans (44). It is produced, stored, and secreted mainly by phagocytes, such as macrophages and neutrophils (5). Two CHIT1 isoforms exist: a secreted 50 kDa protein and a 39 kDa protein which is found in the lysosomes (5). In the healthy human lung, chitotriosidase is expressed by alveolar macrophages (45), while Chit1 mRNA is often undetectable in the mouse lung at steady state (45-48).

Originally, Chit1 was believed to confer protection from chitin-bearing pathogens, such as fungi. *In vitro* exposure of chitin-containing fungi to recombinant Chit1 results in growth inhibition (5). The *in vitro* activity of recombinant Chit1 was also observed in *in vivo* models of systemic Candidiasis and systemic Aspergillosis (5). In contrast to these observations, *Chit1* knockout mice had prolonged survival due to decreased Th2 induction (49, 50). A common polymorphism in *CHIT1* exists in which a 24-bp duplication results in aberrant splicing, leading to an in-frame deletion making the expressed CHIT1 inactive (51). Homozygous individuals have no CHIT1 chitinolytic activity, while heterozygous individuals have reduced activity. This defect is frequently encountered in different ethnic groups, suggesting that CHIT1 chitinolytic activity is not (any longer) indispensable for the defence against chitin-containing pathogens. The divergent frequencies of the 24-bp duplication allele in different populations led to speculations that active CHIT1 might still have benefits in host defence (52). However, no strong or consistent support for this hypothesis exists. While some studies associated this 24-bp duplication polymorphism with increased susceptibility to *Plasmodium falciparum* and filariasis (53, 54), others could not confirm this (55). It might be hypothesized that active CHIT1 is no longer required in evolution because of the duplication event that resulted in a second active chitinase, *AMCase*. However, their different expression profiles (45) and the inactive splice variant of *AMCase* expressed in the human airway (43) do not support this hypothesis.

To date, CHIT1 is thought to have more functions than only host defence. Increased levels of CHIT1 are observed in a variety of granulomatous and fibrotic lung diseases associated with tissue remodelling, such as tuberculosis, sarcoidosis, idiopathic pulmonary fibrosis, scleroderma-associated interstitial lung diseases, and chronic

obstructive pulmonary disease (COPD) (56). *CHIT1* is expressed by a specific subset of recruited macrophages that also express other profibrotic genes, and co-express *CHIL3L1* (YKL-40) (57). This was also observed in yet unpublished single cell RNA-sequencing data from COVID-19 patients included in the SARPAC trial (58). The elevation of CHIT1 in these patients reflects chronically activated tissue macrophages and has been suggested to also contribute to tissue inflammation and remodelling (2). One study using mice genetically deficient in *Chit1* reported that Chit1 was instrumental in bleomycin- and IL-13-induced pulmonary fibrosis by augmenting TGF- β and MAPK signalling in mice (59). However, the role of Chit1 in inflammatory and remodelling responses remains to be elucidated, as only few functional studies are available.

To date, no strong evidence supports CHIT1 being a major player in asthma pathogenesis. Two independent studies showed that CHIT1 concentrations in the bronchoalveolar lavage (BAL) fluid of asthmatics did not differ from those observed in healthy controls (43, 60). Rather, a smoking habit and COPD were observed to result in increased chitotriosidase activity in the BAL fluid. Similarly, another study found serum CHIT1 levels were only very modestly increased in asthmatics compared to healthy controls, while much more pronounced increases were observed in COPD patients (61). Conversely, CHIT1 levels were observed to be locally increased in the lungs after segmental allergen challenge during bronchoscopy in patients with atopic asthma (62). However, this study did not include healthy controls, meaning the transient increase in CHIT1 expression might have been due to phagocyte activation, possibly also observed in healthy controls. Most genetic association studies failed to find an association between *CHIT1* polymorphisms and asthma/atopy (52, 63-65). Only one study reported that a single polymorphism in *CHIT1* was associated with atopy in Korean children (66). While not all reports found *Chit1* expression in the lungs of mice, those that did detect *Chit1* expression found it was not altered upon induction of allergic airway inflammation (42, 46), making Chit1 unlikely to play an exacerbating role in allergy and asthma in mice. In contrast, Hong and colleagues reported CHIT1 concentrations were higher in the sputum of asthmatic children compared to age-matched controls, and that the Chit1 concentration was increased in the lungs from OVA sensitized and challenged mice (67). Using *Chit1* knockout mice, they claimed that Chit1 inhibits allergic airway disease through induction

of Foxp3⁺ regulatory T cells via regulation of TGF- β signalling (ref.). These findings need confirmation, as no other reports on the functional role of chitotriosidase in asthma are available to date.

AMCase

AMCase is expressed in the stomach and to a lesser extent in the lungs (45). While being inactive in the human lung, AMCase is the major chitinase contributing to airway chitinolytic activity in mice (67). Airway epithelial cells are the main source of AMCase in mouse airways, while AMCase was found to be mainly produced by macrophages in human lung (43). Also in mice, AMCase was reported to be expressed by alveolar macrophages (23, 42), but this appears to be under the control of specific stimuli as not all studies could confirm this (48). Kim et al. found both epithelial cells and macrophages to contribute to biologically significant amounts of AMCase (23), while Van Dyken et al. found the epithelium to be the main producer of AMCase (48).

Its high expression in the gastro-intestinal tract, together with its acid stability and low pH optimum, suggest that AMCase might be implicated in the processing of chitin in the gut. However, it seems that humans and rodents do not use chitin as a dietary source. Instead, chitinase expression in the gut is likely to confer protection against chitin-bearing parasites. Vannella et al. showed that mice genetically lacking AMCase had decreased clearance of gastro-intestinal parasites due to profound defects in type-2 immunity (46). The authors hypothesized that the chitinolytic activity of AMCase in the stomach aids in disrupting, releasing, and processing parasite-associated chitinous antigens that are critical for the initiation of protective immune responses in the gut.

In analogy with its function in the gastro-intestinal tract, it is plausible that AMCase also protects the airways from chitin-containing organisms and chitin-induced inflammation. Yet, its function in the airways seems to be much more complex than in the gut, possibly depending on both its chitinolytic activity and chitinolytic-independent effects. As many sources of aeroallergens, such as fungi, HDM and cockroaches, contain chitin, AMCase might also influence the development of atopic asthma. Zhu et al. were the first

to report AMCase is increased in the lungs of asthmatics and claimed a crucial role for this chitinase in asthma based on their mouse studies (42). Since then, extensive research on AMCase and allergy pathogenesis was conducted.

In accord with increased AMCase levels in the lungs of asthmatics (42), AMCase expression is also increased during other manifestations of allergic inflammation, such as allergic rhinitis, allergic rhinoconjunctivitis, and nasal polyposis (68-70), as well as in the lungs of allergen-challenged mice (23, 42, 47). Increased airway chitinolytic activity upon allergen challenges in mice is dependent on IL-13 and STAT6 signalling (41, 42). Some polymorphisms in the *CHIA* gene are associated with pediatric asthma (71), while polymorphisms associated with increased chitinase activity *in vitro* were protective against asthma (72). However, other genetic association studies failed to find associations between *CHIA* polymorphisms and asthma (65).

Several mouse models have been used to investigate the functional roles of AMCase in allergies, yielding highly conflicting results. Zhu et al. observed that the inhibition of AMCase ameliorated pulmonary inflammation and airway hyperreactivity in a chitin-independent model of OVA-induced allergic airway disease and IL-13 overexpressing mice (42). While not affecting IL-13 levels, AMCase inhibition diminished the expression of chemokines normally induced by IL-13, suggesting that AMCase contributes to the downstream chemokine responses to IL-13. Later studies using different chitinase inhibitors and AMCase-silencing RNA also observed that blocking AMCase ameliorates features of allergic airway disease in the OVA-induced asthma model (73-75). In addition, in a model of HDM-driven allergic asthma, targeting AMCase with a specific inhibitor also reduced lung inflammation and serum IgE concentrations (76). One could question whether the systemic administration of chitinase inhibitors is ideal to study airway chitinase biology, as this strategy might have broader systemic effects. Moreover, chitinase inhibitors have targets other than AMCase. They might block the other active chitinase Chit1, and have the potential to bind a range of CLPs, possibly interfering with their function as well. As the above-described studies were mainly performed with non-chitinous allergens and adjuvants, AMCase is suggested to exert its effects through a mechanism that is independent of its chitinolytic activity, including the protection of airway

epithelial cells from apoptosis (77). Yet, it should be noted that two separately generated AMCase transgenic mice exhibit normal lung function and do not develop lung disease in the steady state, meaning that high concentrations of AMCase in the airways are not sufficient to trigger or potentiate allergic inflammation. However, this does not rule out the possibility that AMCase contributes to inflammation once the appropriate stimuli are present.

In contrast, in the presence of chitin, AMCase does have a protective role in airway inflammation through its capability to degrade chitin (23, 24, 48, 78, 79). This was first proposed by Reese et al., who observed that chitin-induced airway eosinophilia was ameliorated in mice overexpressing AMCase (24). Conversely, AMCase-deficient mice exhibit premature morbidity and mortality, concomitant with accumulation of environmentally derived chitin polymers in the airways and increased pulmonary inflammation (48). The inflammatory infiltrate observed in these aged AMCase-deficient mice resembled the profile of lung cellular infiltrates induced after acute inhalation of purified chitin in wild-type (WT) mice (26). The inability to degrade chitin resulted in the development of pulmonary fibrosis, which could be reversed by restoring chitinase activity in the airways. AMCase was also protective against allergic airway inflammation driven by chitinous aeroallergens. Constitutive overexpression of AMCase protected mice from airway eosinophilia in response to fungal challenges (78, 79). Conversely, treatment with the chitinase inhibitor allosamidin prolonged the duration of tissue eosinophilia (79). In addition, mice expressing enzymatically inactive AMCase showed enhanced type-2 immune responses upon intratracheal sensitisation and challenges with HDM extract (23). The authors of this study showed that the inability to cleave chitin prevented macrophages from phagocytosing chitin; uncleaved chitin remains outside the cells, continuing to cause tissue damage and IL-33 release. These inactive AMCase expressing mice showed no difference from WT mice in airway inflammation in the chitin-independent OVA-alum model, confirming that the observed effects of AMCase are mediated through its chitinolytic activity, and suggesting that AMCase does not influence allergic airway disease via a chitinolytic-independent mechanism, in contrast to the earlier reports. In contrast with the above-described studies, mice congenitally lacking AMCase demonstrated little to no role for the enzyme in establishing type-2 immunity in models of

HDM- or OVA-induced allergy in the lung, despite using several asthma models, with both chitinous and non-chitinous allergens (46, 47). This might possibly be due to chitin contents being too low in the HDM extracts used.

The seemingly contrasting observations in mouse models on the role of AMCase might have several reasons. First, AMCase might have distinct functions depending on the context. It clearly is protective against chitin-induced airway inflammation, while in the absence of chitin, AMCase only exerts functions independent of its chitinolytic activity. However, it remains to be elucidated if these functions exist and what they are. Of note, it might be questioned how relevant models using purified chitin or filtered, crushed allergen extracts, are for studying the function of AMCase in airway inflammation. Usually, chitin is not inhaled as a pure polymer, but rather complexed with other sugars and protein antigens. When intact particulate antigen is inhaled, chitinases might act to create additional free chitin ends in particles, promoting chitin recognition by TLR2 (15). However, additional chitinase activity might generate small chitin fragments, which were shown to inhibit TLR2 signalling (15). Moreover, it can be hypothesized that chitinases might promote the release of allergens associated with chitin, promoting the induction of more profound adaptive type-2 immune response.

A second reason for the controversy on AMCase is the use of different strategies to block its function. Thirdly, the hypothesis has been raised that when AMCase is absent, Chit1, the other active chitinase, might compensate for its loss and maintain airway chitinolytic activity. This hypothesis was introduced with the observation that AMCase-deficient mice still had low levels of chitinolytic activity in the serum, which might be attributed to Chit1 (47). This idea has been challenged by some studies that could not find Chit1 expression in the lungs of either WT or AMCase-deficient mice (47, 48). However, Vannella et al., who did not observe different asthmatic features in WT and AMCase deficient mice, did observe expression of Chit1 in the lungs (46). The controversial observation of Chit1 expression in mouse lungs might be due to different activation status of lung macrophages, possibly influenced by housing conditions and microbiome.

The potential significance of targeting AMCase in human asthma remains uncertain, since AMCase was shown to minimally contribute to chitinolytic activity in the

human lung (43). Inactive AMCase retains its ability to bind chitin, and as such, might still influence inflammation and tissue remodeling in analogy with CLPs, as described below. Hence, future research should try to unravel the possible functions of AMCase that are independent of its chitinolytic activity.

Chitinase-like proteins

Mammalian CLPs have a chitin binding domain, yet lack enzymatic activity: YKL-39 (*CHI3L2*) and YKL-40 (*CHI3L1*) in humans and Ym1 (*Chil3*), Ym2 (*Chil4*) and BRP-39 (*Chil1/Chi3l1*) in the mouse (Table 1). In CLPs, the substitution of an essential glutamic acid to leucine, isoleucine or glutamine within the highly conserved enzyme site accounts for the lack of chitinolytic activity. In both species, CLPs have been implicated in an enormous variety of pathologies, suggesting broad generalized functions (8). Among human and murine CLPs, YKL-40 retained the property of binding chitin with high affinity (6), while Ym1 does not bind chitin but chitin-like saccharides such as glucosamine oligosaccharides, heparin and heparan sulfate (80). The fact that some CLPs retain chitin-binding ability suggests they contribute to the recognition and immune signaling of chitin-associated PAMPs, or might control the substrate specificity of real chitinolytic enzymes by competitive or synergistic binding. However, CLPs have been evolving at a remarkably rapid rate, potentially associated with the acquisition of novel, chitin-independent regulatory functions. It is therefore no surprise that the role of CLPs in host immunity, especially their contribution to type-2 immunity, has received great attention in recent years. CLPs are expressed in immune and structural cells in steady state, while type 2-polarized immune responses associated with repair and regeneration boost their expression. Dysregulated CLP expression is often associated with inflammatory conditions including allergic asthma and related obstructive lung diseases, as will be discussed in detail below. In addition, at least in mice, decreased lung CLP transcripts are a genetic determinant of lung function associated with lower basal pulmonary capacity (81).

YKL-40/BRP-39

In humans, several CLPs have been identified. Among these, YKL-40 is the most prominently expressed during lung inflammation and tissue remodelling (11), with increased expression associated with many pathologies including allergic asthma (8). In fact, YKL-40 is increased in the serum and lungs of patients with asthma, correlated with disease severity and asthma exacerbation rate and inversely correlated with lung function (82-84). In these settings, YKL-40 can be produced by a multitude of cells of both hematopoietic and non-hematopoietic origin as reviewed in (85). YKL-40 was shown to contribute to the differentiation of monocytes to activated macrophages in inflamed tissues (86), facilitating the late stages of human macrophage maturation (87). Enhanced expression of YKL-40 protein is also found in bronchiolar epithelial cells and alveolar macrophages adjacent to fibrotic lesions (88) and YKL-40 levels in bronchial smooth muscle cells show a clear correlation with the thickness of bronchial epithelial basement membrane in asthmatic patients (89). Experimental studies then showed enhanced proliferation and migration of bronchial smooth muscle cells in response to YKL-40 (90). This might be due to the binding of YKL-40 to type I collagen and subsequent formation of collagen fibrils and stimulation of pro-fibrotic and inflammatory mediators (91, 92).

Thus, YKL-40 seems to be closely related to the pathogenesis of asthma, especially airway remodeling. It is hence not surprising that variation in *CHI3L1*, the gene that encodes the YKL-40 protein, is closely linked to asthma susceptibility and reduced lung function, and that YKL-40 is a significant biomarker for asthma severity (4, 93). A correlation between *CHI3L1* SNPs and asthma is evident as *CHI3L1* polymorphisms have been associated with increased serum YKL-40 levels, reduced pulmonary function, bronchial hyperresponsiveness and airway remodeling (93-97). One of the most recent meta-analyses investigating a total of 13 articles showed that the serum concentration of YKL-40 in asthmatic patients was significantly higher than in healthy subjects and YKL-40 tracked with asthma severity (98). However, the correlation between YKL-40 and Th2-related inflammation in asthma is still controversial (98) as asthma is a highly heterogeneous disease presenting with different endotypes and inflammatory signatures that may be broadly viewed as type-2-high or type-2-low (7, 99). Initial studies suggested that YKL-40 may enhance allergen sensitization and IL-4/IL-13 Th2 cytokine responses (100). More recent work rather points to a connection of elevated YKL-40 levels in non-

eosinophilic and paucigranulocytic asthma as compared to type-2 high, eosinophilic asthma. This is likely through elevated YKL-40 expression by various cell types including neutrophils and macrophages (101). These findings are in line with other reports of significant positive relationships between YKL-40 levels, blood neutrophil numbers, asthma severity and reduced lung function (102-104), strengthening the association of serum YKL-40 with non-eosinophilic, type-2-low asthma.

However, while this seems to be true for adult-onset asthma, data on elevated YKL-40 levels and asthma severity in childhood asthma are conflicting. While Konradsen et al. found that compared with healthy controls, serum YKL-40 levels were higher in children with severe, therapy-resistant asthma and were associated with the *CHI3L1* promoter single nucleotide polymorphism (105), another study found no association of elevated YKL-40 levels with asthma severity, lung function or type-2 inflammation in asthmatic children (106).

In mice, BRP-39 is often referred to as the 'prototypical' CLP, because it is the genetic orthologue of YKL-40 in humans (8). Elias et al. generated and characterized BRP-39-deficient mice (*Chi3l1*^{-/-}), YKL-40 transgenic mice, and *Chi3l1*^{-/-} mice that simultaneously produced transgenic human YKL-40 only in their respiratory epithelium to overcome any contributing effects of the mouse ortholog BRP-39 (100). BRP-39 was significantly increased after OVA-induced allergic airway inflammation, predominantly in type 2 alveolar cells and alveolar macrophages. Others found BRP-39 levels not significantly altered following allergic inflammation, claiming that Ym1/2 proteins mimic the biological effects of YKL-40 more closely (8, 107, 108).

Loss of BRP-39 resulted in substantial reductions in all the hallmarks of allergic asthma, along with increased apoptosis of CD4 T cells and eosinophils. In addition, BRP-39 potently stimulated AAM polarization and the production of the Th2 chemokines CCL17 and CCL22 from alveolar macrophages. Overexpression of YKL-40 in BRP-39-deficient mice rescued type-2 responses to levels comparable to those seen in WT animals (100, 109). Mechanistically, BRP-39 was driven by IL-13 and as such seems to be a critical downstream target of IL-13 effector responses (109). Following up on this, the authors could demonstrate that both YKL-40 and BRP-39 bind to the interleukin 13 receptor alpha

2 (IL-13R α 2), in concert with IL-13, although this finding still needs validation by other groups (109). Since then, numerous YKL-40/BRP-39 binding partners have been identified including Prostaglandin D2 receptor, receptor for advanced glycation end product and type I collagen (8).

Similarly diminished type-2 inflammation in *Chil3l1*^{-/-} mice could be observed after HDM exposure (110) and in a model of fungal-associated allergic airway inflammation (111). However, despite significantly lower type-2 responses to fungal exposure, mice showed significantly increased airway hyperresponsiveness, indicating that BRP-39 protects against airway hyperresponsiveness during fungal asthma despite contributing to type-2 inflammation (111). The authors also found higher serum YKL-40 concentration in fungal-sensitized asthmatics (111), which is, as already discussed, often associated with more severe disease, reduced lung function and features of type-2 low asthma. Along these lines, BRP-39 has been shown to support IL-17A production by $\gamma\delta$ T cells and both total numbers of $\gamma\delta$ T cells and IL-17A-producing $\gamma\delta$ T cells were significantly lower in *Chil3l1*^{-/-} mice than in wild-type mice (108). However, levels of IFN- γ and IL-17A were unaffected by the absence of BRP-39 during fungal asthma as were the levels of neutrophils in the lung (111).

In summary, the studies demonstrate that correlation of asthma endotypes and *CHI3L1* SNPs is warranted when looking at the contribution of YKL-40 to allergic asthma (98). Whether YKL-40 participates in the pathogenesis of asthma and/or is a biomarker of asthma severity remains an open question, given the divergent evidence at present (83).

Ym1/Ym2

Ym1 and Ym2 proteins, encoded by the genes *Chil3* and *Chil4*, respectively, on mouse chromosome 3, are rodent-specific CLPs with no true human orthologs (4). They are highly homologous, sharing up to 91% amino acid sequence identity, indicating a relatively recent gene duplication event (8). Nevertheless, Ym1 and Ym2 show obvious tissue expression specificity (112, 113) and have fairly distinct cellular sources (107).

Under steady state conditions, Ym1 is the most prominently expressed CLP in the mouse lung, while Ym2 is barely detectable. Allergens (107, REF, 108, 114), particulate matter (115), helminthic parasites (108), and cancer (116) significantly increase expression of Ym2, which then becomes the predominant lung CLP. While Ym2 is largely expressed by airway epithelial cells (107) under these conditions, Ym1 is mainly restricted to the myeloid compartment (117-120). This is a very recent finding because due to the lack of specific tools, Ym1 and Ym2 could not be clearly distinguished and the literature is confounded by observations that can be attributed to both Ym1 and Ym2.

A distinct feature of Ym proteins is that they are commonly found to form crystals during 'type-2-high' responses including allergic airway inflammation (114, 121). Recent findings suggest that a hyperactivated IL-33/ILC2 axis might drive Ym1/Ym2 crystallopathies in mice (122), like Charcot-Leyden crystallopathy in humans during hypereosinophilic inflammation (123). In this context, Ym proteins are best known as IL-4/IL-13-inducible (114, 117, 124) hallmark genes of AAMs, which is mediated by STAT6 (118) and further facilitated by PPAR γ signaling (125). The dependence on IL-4/IL-13 might also partly explain the occurrence of Ym crystals in viable motheaten mice that harbor a mutant Src homology protein tyrosine phosphatase that, amongst others, negatively regulates IL-4R α signaling, leading to uncontrolled Ym protein expression in these mice (121). However, Ym1/ Ym2 protein expression is far more than just a mere marker of type-2-high responses although their detailed functions during such so far remain poorly understood.

While Ym1 was initially described as a chemotactic factor for eosinophils (known as eosinophil chemotactic factor or ECF-L) (126), this chemotactic activity could later not be confirmed in *in vitro* and *in vivo* settings (80, 114). There is, however, consensus that Ym protein expression is highly increased in the mouse lung and BAL fluid during experimental models of allergic asthma (112, 114, 124). In response to IL-13, dendritic cells (DC) upregulate Ym1/2 expression to enable Th2 cytokine production by CD4⁺ T cells; Ym1/2 supplementation of IL-13-deficient DCs restored the ability to stimulate IL-5/IL-13 secretion in DC/CD4 T cell cocultures by inhibiting the production of 12-hydroxyeicosatetraenoic acid by 12/15(S)-lipoxygenase (124). In turn, blocking Ym1/2 signaling during the induction of allergic airways disease with an anti-Ym1/2 antibody or

targeted delivery of siRNA attenuated IL-5 and IL-13 production in the mediastinal lymph nodes (124) and protected from allergen-induced allergic airway inflammation (127, 128), underscoring the relevance of Ym1/Ym2 in the enhancement of type 2 responses. However, there was no experimental distinction between Ym1 and Ym2 proteins and it was not clear whether both or one of the two was triggering type-2 cytokines.

In addition, a growing body of literature has accumulated around the role of Ym1 in IL-17-mediated neutrophil recruitment in allergic lung inflammation (108, 129). Plasmid-directed overexpression of Ym1 protein increased neutrophilia in the lungs, while eosinophilia was significantly decreased, dependent on IL-17 production from $\gamma\delta$ T cells. By blocking Ym proteins, the number and proportion of neutrophils in the lung were decreased, along with reduced expression of IL-17A and IL-17A target genes in OVA-challenged mice (108). In mice infected with *N. brasiliensis*, blocking Ym1 in the early innate immune stage could reduce the amount of Th2 cytokines in mice; while during the adaptive type-2 response, blocking Ym1 significantly enhanced type-2 cytokine production from both ILCs and CD4⁺ T; indicating that Ym1 is important for limiting the magnitude of type-2 responses. More importantly, Ym1 could directly contribute to lung repair via enhanced RELM α production (129).

Recent publications underscore the role of Ym1 in tissue repair while also reinforcing its function in type-2 immune responses and AAM polarization. Mice harboring a natural polymorphism in the *Chil3* promoter, which results in partial Ym1-deficiency, show reduced OVA-induced allergic inflammation, and demonstrate an enhanced AAM phenotype, indicating that Ym1 may control or limit the alternative activation of macrophages (112). Inhibition of Ym1⁺ AAMs by cynaropicrin, a galectin-3 pathway inhibitor, dampened eosinophilic lung inflammation in a model of HDM-induced allergic airway inflammation, while simultaneously inducing neutrophil influx. This resulted in worsened airway hyperresponsiveness to methacholine but significantly lower collagen deposition (130). In addition, tissue injury leads to the recruitment of Ym1⁺Ly6C^{hi} monocytes from the bone marrow, which are associated with the resolution of inflammation and tissue healing, and exhibit an immunoregulatory phenotype (120). One of the postulated functions of Ym1 is binding to components of the extracellular matrix, facilitating matrix deposition and tissue repair (80, 131). Differences in deposition of

specific collagen subtypes have been noticed between mouse strains (107), and Ym1 expression differs considerably between distinct inbred and wild-derived mouse strains due to natural polymorphism in their respective promotor regions (112). Hence, Ym1 polymorphisms might be correlated with differential susceptibility to type-2 inflammation-related tissue remodeling (112).

In summary, Ym proteins are generally recognized as fundamental players in allergic airway disease in mice, contributing to type-2 responses, AAM polarization and tissue repair. The effects ascribed to Ym proteins are, however, multifaceted, and even contradictory. One of many reasons is the lack of specific tools to clearly distinguish Ym1 and Ym2 proteins in the lung; as such, the literature is confounded by observations that could be attributed to either Ym1 or Ym2. However, they clearly differ in their spatial (myeloid vs. epithelial cells) and temporal (constitutive vs induced) expression patterns, and hence can be assumed to perform distinct roles in allergic airway inflammation. In addition, bi-phasic upregulation of Ym1 has been observed, which might play diametrically opposite roles in the early vs. late phase of inflammation and explain type-2 polarization vs. Th17 responses but also the different cellular sources of Ym1. The field urgently awaits genetic tools to delete either *Chil3* or *Chil4*, but creation of such tools is hampered by the fact that the chromosome 3 locus has a gene duplication.

Evolutionary biology of the chitinase and CLP family

It is intriguing that the genes coding for chitin synthase have been lost in mammals, while several genes encoding chitinases have been preserved. Not only have chitinase genes persisted, but multiple gene duplication events and loss-of-function mutations have led to emergence and diversification of CLPs with species- and tissue-specific expression patterns (9, 11, 132, 133). Especially CLPs have evolved at a speed that points towards positive selection rather than natural genetic drift (8, 9, 11, 132, 133), indicating distinct evolutionary benefits. One possible explanation is offered by the ‘innovation, amplification, and divergence’ model of gene evolution (133, 134). This model infers that active chitinases were no longer needed, which enabled (or facilitated) mutations in critical regions such as the catalytic active site in duplicated genes, giving rise to new, non-enzymatic proteins. However, these “dead enzymes” were still selectively maintained for

their beneficial and novel regulatory functions (133, 134). Specific sites within CLP proteins are under positive selective pressure that drives CLP functional diversity across species - yet, the sugar-binding barrel structure is highly conserved (8). As many authors have remarked, it is tempting to look at CLPs as examples of recent adaptations and gene evolution 'happening right now' (11, 133). In addition, despite the shared chromosomal location and high homology, individual genes have evolved independently and can have distinct functions within and across species (2, 8). This also raises concerns as to whether chitinase and CLP research using mouse models of allergic asthma is relevant to human pathology.

So the question remains: Why are chitinases and CLPs evolutionary hot spots? One possibility is that they might still be taking part in the defence against chitin-containing pathogens. However, in higher organisms, their chitin-independent and regulatory functions seem to be dominant. This is best illustrated by the human *CHIT1* gene, for which the 24-bp duplication polymorphism leaves around 5% of the Caucasian population without an active enzyme, with no obvious effects for the host's defense system (51, 52). Nevertheless, the polymorphism is strongly associated with an increased rate of lung function decline in COPD, pointing to a protective role in pulmonary tissue homeostasis (133). Similarly, AMCase shows regulatory and Th2-promoting functions that are completely independent of its enzymatic activity (77). In case of CLPs, regulatory functions have been discussed in the context of type-2 related tissue inflammation and remodeling through binding to extracellular matrix components. While Ym1 has specific binding capacity for GlcNAc oligomers and heparin/heparan sulfate proteoglycans, YKL-40 interacts with type I collagen. Whether CLPs could perform their regulatory functions also through protein-protein interactions, i.e, by binding to specific surface receptors on immune cells or by altering the substrate specificity of the true chitinases, remains to be established. The mining of multi-omics data may help modelling interactomes to identify new interaction partners and biological processes associated with the non-chitin binding, regulatory function of chitinases and CLPs.

Conclusions and future perspective

Despite extensive research into the role of chitinases and CLPs in allergic airway disease, there is still no agreement on whether they are mere biomarkers of disease or actual disease drivers. However, their ubiquitous presence and evolutionary conservation suggest they have functions that are far more complex and diverse than previously realized.

Functions ascribed to chitinases and CLPs include but are not limited to: (1) a role in host response against chitin-containing pathogens, by either degrading or binding to chitin, which triggers an innate immune response; (2) directly promoting chemotaxis or production of pro-inflammatory cytokines; (3) modulating tissue remodeling and fibrosis, possibly through interactions with components of the host extracellular matrix. These functions are likely beneficial to combat chitin-bearing parasites, a process in which these molecules are required for expulsion and wound healing. However, in pulmonary diseases, these same properties may cause pathology (117).

In the future, it will be crucial to identify yet unknown receptors and/or ligands of chitinases and CLPs to better understand their type-2 immunity-triggering properties. One limitation of mouse models in human CLP research is assigning orthology and/or functional homology to human genes and proteins. Although BRP-39 is the predicted genetic ortholog of YKL-40, Ym1/Ym2 proteins, which are often disregarded as being restricted to the mouse system, are found among the most upregulated genes after allergen exposure, similar to YKL-40 in humans. As such, considering CLPs in their entirety and understanding their functional redundancies could help to understand their potential and value for human CLP research in allergic airway diseases. Lastly, as there have been quite opposing findings between different chitinase and CLP studies reported in the literature, a number of considerations with major potential impact on research outcomes should be taken into account: (1) differences in allergen models and routes of exposure; (2) genetic background of the mouse strains as well as genetic strategies to create knockout strains; (3) differences in the housing and husbandry conditions (i.e., diet, commensals).

In conclusion, chitin is a size-dependent, pathogen-associated molecular pattern that is recognized by the human immune system through TLR2 and possibly other

receptors. Acute chitin challenges in the lungs result in the activation of complex pathways inducing a mixed neutrophilic-eosinophilic infiltrate, which can be reduced by active chitinases. However, chitinases might have functions beyond their chitinolytic activity. This is suggested by the evolution driving CLP duplication and studies that find roles for active chitinases in the absence of chitin.

Table and Figures

True chitinases					
Gene	Protein	Human	Mouse	Tissue/cellular expression	Reference
<i>CHIA</i>	AMCase	Chr1	Chr3	Macrophages, lung epithelial cells	(42, 45, 48)
<i>CHIT1</i>	CHIT1	Chr1	Chr1	macrophages, neutrophils, microglia	(5, 43, 45)
Chitinase-like proteins (CLPs) = chitolectins = chilectins					
Gene	Protein	Human	Mouse		
<i>CHIL1/Chil1</i>	YKL40/BRP39 (CHI3L1, HC-gp39, GP39, cartilageglycoprotein1)	Chr1	Chr1	chondrocytes, sinovial cells, macrophages, fibroblasts, neutrophils, epithelial cells, osteoclasts, astrocytes	(85)
<i>CHIL2</i>	YKL39 (CHI3L2, chondrocyte protein 39)	Chr1		cartilage chondrocytes	(135, 136)
<i>Chil3</i>	Ym1 (Chi3l3, ECF-L)		Chr3	macrophages, dendritic cells, monocytes, neutrophils	(107, 112, 113, 117-120)
<i>Chil4</i>	Ym2 (CHI3L4)		Chr3	bronchial epithelial cells, gut	(107, 112, 113)
<i>Chil5/Chil6/Gm6522</i>	Ym3/Ym4		Chr3	pseudogene	(39)

<i>OVGP1</i>	OVGP1 (oviductin,mucin 9)	Chr1	Chr3	oviductal epithelial cells	(137, 138)
<i>SI-CLP</i>	SI-CLP (CHID1)	Chr11	Chr7	monocytes, macrophages, neutrophils	(139-141)

Table 1: Summary of chitinases and CLPs genes found in humans and mice

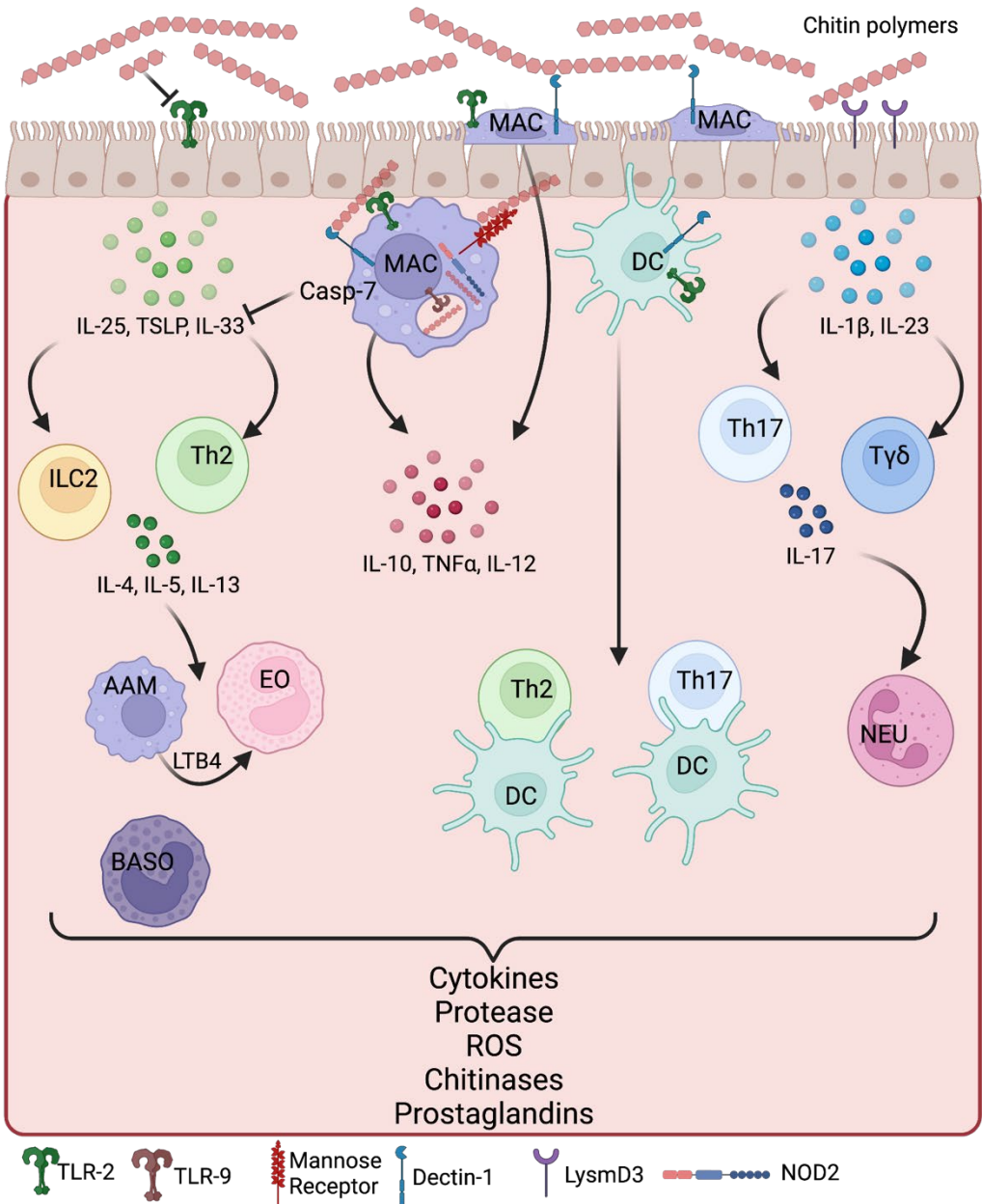


Figure 1: Overview of chitin induced immune responses in the airways.

Depending on its size, chitin can stimulate epithelial and immune cells in the airways by triggering pattern recognition receptors (PRR). TLR-2, TLR-9, mannose receptor, dectin-1, LysMD3, and NOD2 have been implicated in chitin sensing. Activation of epithelial cells by chitin leads to the production of type-2 cytokines (such as IL-25, TSLP, and IL-33), but also production of IL-1 β and IL-23. These epithelial cytokines can stimulate adaptive and innate type-2 cells (Th2 cells, ILC2, basophils, eosinophils, and AAM) and type-17 cells (Th17 cells, $\gamma\delta$ T cells, and neutrophils), which produce many effector cytokines and mediators that contribute to inflammation, damage, and tissue remodelling. Moreover, the epithelial cytokine response can stimulate DCs to activate adaptive immune responses. DCs express PRR implicated in chitin sensing, but direct effects of chitin on DCs have not been investigated. Chitin can directly stimulate macrophages to secrete a variety of cytokines such as IL-10, TNF α , and IL-12. In addition, appropriately sized chitin can be phagocytosed by macrophages, also leading to cytokine secretion. Chitin phagocytosis by macrophages leads to the induction of casp-7, which can inactivate IL-33, leading to inhibition of type 2 immune responses.

Abbreviations: AAM: alternatively activated macrophage; BASO: basophil; Casp: caspase; DC: dendritic cell; EO: eosinophil; IL: interleukin; ILC: innate lymphoid cell; LTB4: leukotriene-B4; MAC: macrophage; NEU: neutrophil; ROS: reactive oxygen species; T gd: $\gamma\delta$ T cell; Th: T helper cell; TNF α : tumor necrosis factor α ; TSLP: thymic stromal lymphopoietin.

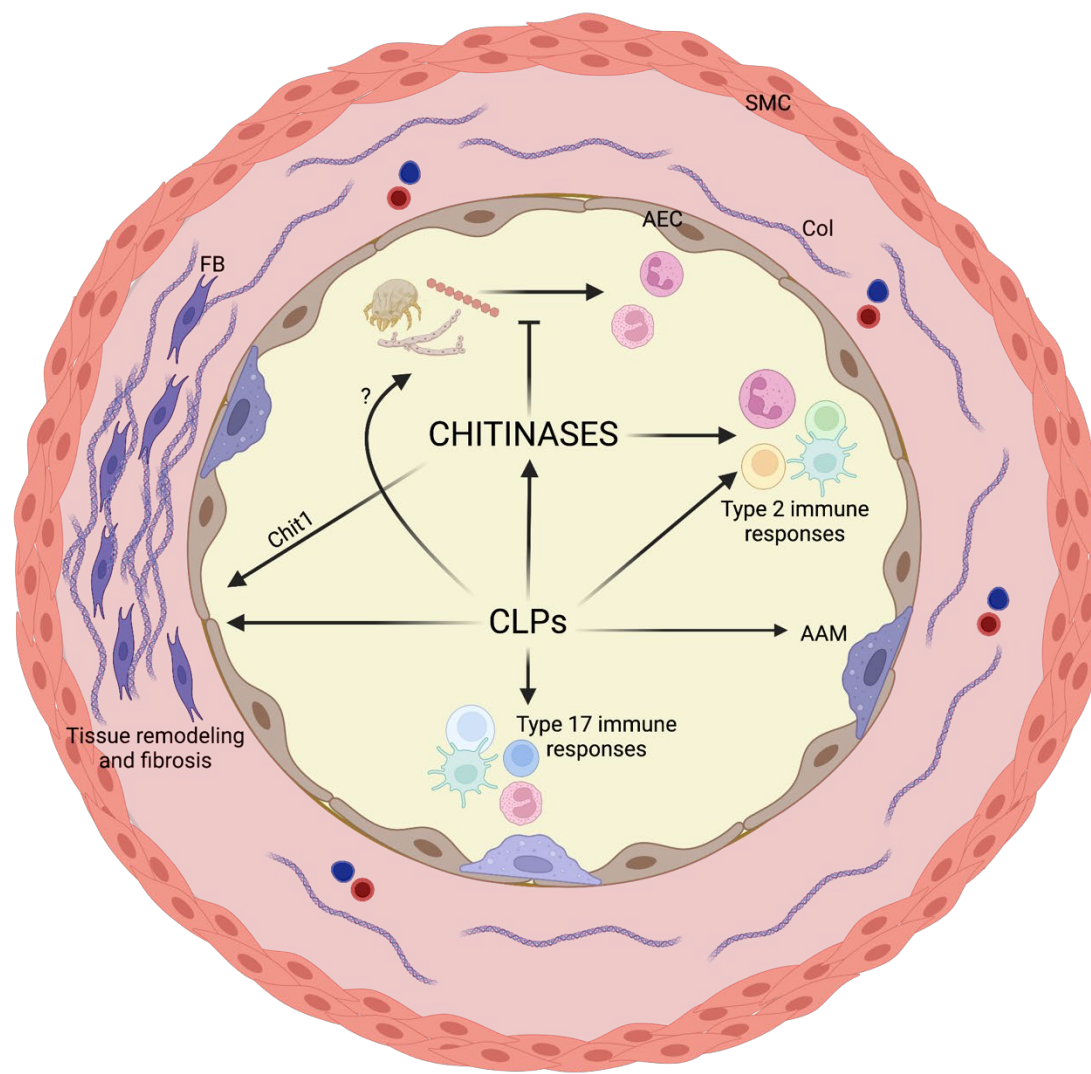


Figure 2: Overview of functions of chitinases and CLPs in the airways.

When environmental chitin, i.e. the one contained in house dust mites and fungi, is inhaled, it is degraded by active chitinases (AMCase and Chit1), thereby limiting chitin-induced airway inflammation. At the same time, AMCase might also contribute to type-2 immune responses in the airways. In addition, Chit1 secreted by activated macrophages might contribute to tissue remodeling and fibrosis through yet unknown mechanisms. CLPs do not cleave chitin but might control the substrate specificity of real chitinolytic enzymes by competitive or synergistic binding. In addition, CLPs might be implicated in chitin sensing by binding to chitin polymers. Moreover, CLPs contribute to type-2 immune responses and alternative activation of macrophages but have also been implicated in Th17 skewing. Like Chit1, CLPs are implicated in tissue remodeling and repair by

740 interaction with extracellular matrix components and consequently facilitating matrix
741 deposition and tissue repair.

742 *Abbreviations:* AAM: alternatively activated macrophage; AEC: alveolar epithelial cell;
743 Chit1: chitotriosidase; CLP: chitinase-like protein; Col: collagen; FB: fibroblast; SMC:
744 smooth muscle cell.

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