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Severe asthma: entering an era of new concepts and emerging therapies

Highlights of the 4th International Severe Asthma Forum, Madrid, 2018

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To the Editor

In the past decades, new insights into molecular mechanisms of severe asthma have further unveiled its heterogeneous nature prompting the need for personalised and targeted approaches. These and several other hot topics related to severe asthma were addressed in the 4th International Severe Asthma Forum, organized by the Asthma, ENT, Immunology and Pediatric sections of the European Academy of Allergy and Clinical Immunology in collaboration with the Spanish Society for Allergy and Clinical Immunology.

Severe asthma is currently defined as asthma that requires treatment with high-dose inhaled corticosteroids combined with a second controller \pm systemic corticosteroids to maintain control or, asthma that remains “uncontrolled” despite this therapy (1). While severe asthma affects 5-10% of the entire asthma population, it accounts for >80% of the total healthcare costs related to asthma. Of note, severe asthma should be distinguished from difficult-to-treat asthma for which (re)evaluation of diagnosis, precipitating triggers, treatment adherence, as well as treatment of comorbidities is recommended. To improve treatment adherence, patient education on sustained use of controller medications in combination with a proper inhalation technique is mandatory. eHealth, comprising of a variety of tools and applications, including mobile devices (mHealth), can support patient awareness, improve adherence and support disease self-management (2).

Sensitizing agents at the workplace can cause occupational asthma. Irritants are often related to work-exacerbated asthma but at high concentrations may also elicit occupational asthma. A detailed occupational history including the insight into potential sensitizers at work is an integral, but often under-emphasized, part of the evaluation of patient with severe asthma. Patients with occupational asthma due to high-molecular-weight (HMW) agents commonly present with rhinitis, conjunctivitis, atopy and early asthmatic reactions following specific inhalation challenge, while asthma exacerbations are more frequently associated with occupational asthma induced by low-molecular-weight (LMW) agents. Interestingly, the inflammatory profile triggered by HMW and LMW agents is similar (3).

As for comorbidities, chronic rhinosinusitis with nasal polyps is a prevalent upper airway comorbidity in severe asthma, responding to corticosteroids and type 2 immune response (T2)-targeted therapies. Likewise, obesity, a key component of the metabolic syndrome, is linked to severe

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asthma. Asthma patients with metabolic syndrome experience worse respiratory symptoms resulting from both lung function impairment and increased airway inflammation. Obviously, proper advice on dietary intake, lifestyle adjustments and/or bariatric surgery should be part of severe asthma management.

Bronchiectasis is present in approximately 30% of patients with severe asthma. Although both diseases share overlapping features, such as symptoms, neutrophilic inflammation and airway hyperresponsiveness, common underlying pathways require further research.

Furthermore, a significantly higher prevalence of mental disorders, especially depression and anxiety, has been reported in patients with severe asthma. Patients with psychological comorbidities often present with low treatment adherence, smoking, inactivity and obesity. Hence, for severe asthma, psychological support is recommended as part of daily clinical practice.

Coinciding with a reduced response to standard treatment, severe asthma patients are often exposed to an overload of (combinations of nasal, cutaneous, inhaled and oral) corticosteroids causing adverse events which impose substantial health care costs. This clearly underscores the need for alternative treatment options enabling reduction of the overall corticosteroid dose.

Reflecting its heterogeneous nature, severe asthma comprises several different phenotypes, distinguishable by age of onset, allergy status, airway inflammatory pattern and response to treatment (4). Eosinophilic asthma is the most commonly studied inflammatory phenotype, driven by T2 cytokines and present in approximately 50% of asthmatics, with both allergic and non-allergic triggers initiating or aggravating the disease. In contrast, the relevance and driving factors in non-T2 asthma, including neutrophilic asthma, are less well-defined and often precipitated by various factors including pathogens, air pollutants, cigarette smoke, cold air or exercise (5). Finally, the pauci-granulocytic phenotype may reflect either well-treated airway inflammation or symptomatic severe asthma driven by episodic inflammatory or non-inflammatory events, e.g., airway smooth muscle (ASM) hypertrophy (6). More recent data showed that different inflammatory signatures may be present at different anatomic sites within the airways of individual severe asthma patients.

Recent insights into molecular pathways underlying the inflammatory phenotypes helped to define distinct endotypes. Presently, the sub-endotype(s) of T2 asthma are best defined. T2-driven mechanisms are involved in epithelial barrier dysfunction, airway eosinophilia, mucus hypersecretion and airway hyperresponsiveness. In these patients, biologicals targeting T2-pathways, *i.e.*, IgE, IL-4, IL-5, and IL-13, showed an overall 50-60% reduction in severe exacerbations

(7). Future studies will establish if biologicals to more upstream targets will safely deliver additional benefits, e.g. tezepelumab (anti-TSLP), which suppresses both IL4/13, IL5 pathways and also impacts non-T2 inflammatory events. However, targeted interventions so far did not achieve any disease-modifying effects and there is significant heterogeneity in treatment response due to the complexity of the T2 endotype. Additionally, recent health-economic evaluation showed that the cost of currently registered biologics for severe asthma exceeds the recommended maximal cost per quality-adjusted life-year by far.

Treatments targeting the following pathways: alarmins (e.g. TSLP, IL-25, IL-33), kinases (e.g. JAK, Pi3K), and other pro-inflammatory mediators (e.g. PGD2) are currently under clinical development while their long-term effectiveness remains to be seen. While bronchial thermoplasty may be considered for patients with severe asthma with predominant chronic airway obstruction due to ASM hyperplasia, non-T2 asthma and mixed endotypes still represent unmet needs.

Further understanding of the molecular pathways will help to elaborate accurate algorithms matching the right biological to the right patient. To this end, simple and reliable biomarkers are indispensable, which adequately reflect the underlying pathophysiology and the treatment response, as well as predict the long-term outcomes. In connection with the T2-signature, blood or sputum eosinophils and FeNO are presently the best validated biomarkers. While eosinophils can best be used to predict treatment response in eosinophilic (severe) asthma, FeNO proves useful in identifying T2 inflammation, partly unrelated to eosinophils and mainly IL13-driven.

More innovative biomarker techniques, *i.e.* breathomics, require specific expertise including metabolomic analysis of exhaled air through mass spectrometry or eNose technology. Other emerging approaches include transcriptomics, genomics and micro-RNA analysis. Recent data showed that nasal epithelium gene profiling reflects bronchial gene expression and hence may serve as a future biomarker to guide asthma treatment. Alternatively, further classifying biomarkers may arise from new imaging techniques and advanced physiology. Future pragmatic biomarker strategies, combining innovative methods for decoding the molecular, immunological, anatomical and physiological complexity, will further improve (sub)phenotyping, endotyping and personalized treatment.

Pulmonary rehabilitation is an alternative, effective but often overlooked treatment strategy (8). However, patients should be informed on potentially detrimental effects of exposure to irritants such as chlorine by-products or cold air in specific training environments.

In childhood, the increased asthma risk after severe viral bronchiolitis may be due to long-lasting epithelial and innate immune changes, resulting in diminished antiviral immunity and enhanced pro-inflammatory T2 responses. Most children with T2 asthma respond well to low-dose ICS ± additional controller, but there are currently no predicting biomarkers for step-up to LABA, LTRA or increased ICS dose. Rather than increasing pharmacotherapy in the non-responsive child, co-morbidities, social and environmental factors should be addressed, especially poor adherence. In addition to omalizumab, at least 3 additional biologicals (mepolizumab, benralizumab, dupilumab) should become available soon for childhood severe asthma (9). To enable personalised treatment approaches further research into response predictors is key.

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