

Asthma: personalized and precision medicine

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Purpose of review

In this review, we herein describe the progress in management of severe asthma, evolving from a 'blockbuster approach' to a more personalized approach targeted to the utilization of endotype-driven therapies.

Recent findings

Severe asthma characterization in phenotypes and endotypes, by means of specific biomarkers, have led to the dichotomization of the concepts of 'personalized medicine' and 'precision medicine', which are often used as synonyms, but actually have conceptual differences in meaning. The recent contribute of the omic sciences (i.e. proteomics, transcriptomics, metabolomics, genomics, ...) has brought this initially theoretic evolution into a more concrete level.

Summary

This step-by-step transition would bring to a better approach to severe asthmatic patients as the personalization of their therapeutic strategy would bring to a better patient selection, a more precise endotype-driven treatment, and hopefully to better results in terms of reduction of exacerbation rates, symptoms, pulmonary function and quality of life.

Keywords

biological therapies, biomarkers, personalized medicine, precision medicine, severe asthma

INTRODUCTION

Personalization of management and treatment of each patient has been the physician goal since Hippocrates. In the last century, progression of research and industry leaded to great achievements, such as the discovery of molecules able to treat a relevant proportion of patients with a certain disease. In that context, we lived the era of the 'blockbusters' drugs [1].

What was already evident at that time was the following: a need for generic drugs, whose intent is to cover more patients in the developed countries and to extend the treatment to underdeveloped countries; a decrease of the so-called 'blockbuster approach' while developing a 'phenotype driven treatment'; and finally, considering the burden of treatment with branded biologics, we also envisaged the possible forthcoming of biosimilars in asthma and allergic diseases.

Nowadays, mainly because of the new biologic therapies, 'Personalized Medicine' and 'Precision Medicine' are commonly used as terms to indicate a more individualized approach to single patient treatment [2].

We previously reported that the two terms are used interchangeably [2], mainly in this area, but we should revise that view, seeing not a complete overlapping between Personalized and Precision Medicine.

'Precision medicine' is mainly targeting the endotype of the patient, that is the mechanism(s) of the disease of the individual patient [3]. Support to this definition of target patient is coming from several 'omic' sciences, such as proteomics, transcriptomics, metabolomics, genomics, and so forth, ... detecting biomarkers able to identify the molecular mechanisms of the patient''s disease and

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KEY POINTS

- 'Personalized medicine' and 'precision medicine' are often used as synonyms but actually have slight meaning differences.
- Personalized-precision medicine would bring to better treatment strategies in dealing with severe asthmatic patients.
- Checking patient adherence to their therapies, the phenotypization and endotypization processes, the utilization of predictive biomarkers and the contribution of the omic sciences would be relevant in reaching this goal.

accordingly to lead to the choice of the correct and effective biologic drug [4].

Precision medicine is of key importance and it will lead to a new classification of the diseases on the basis of the molecular mechanisms [5].

'Personalized medicine' has a broader value and meaning. Of course, we can see precision medicine as a core moment of personalized medicine, but we should consider all the aspects related to the 'person' we have to treat [6]. In the context of omic sciences, FitzGerald recently proposed the term 'Humanomics', a very interesting concept, refocusing doctor's attention on patient's features [7,8[•]]. In this context, we herein report also 'Personomics', indicating all the aspects (clinical, emotional, psychological, functional, phenotypical, endotypical, etc...) [9] we should consider, evaluate and investigate about the 'person/patient' we have to treat. Thus, the term 'omic sciences' is not just restricted to system biology but it is also encompassing other areas of knowledge and research [10]. (Fig. 2).

Nowadays the most challenging task is to transfer into the current clinical practice the principle(s) of personalized medicine. Recently an interesting approach to this has been envisaged by Pritchard and colleagues and a suitable proposal of how to reach this goal has been formulated [8[•],10]. Interestingly, National Institute for Health and Care Excellence (NICE) reports that the knowledge of the personal experience about a disease (made by values, needs, concerns, beliefs, expectations, ...) represents a fundamental step to achieve the best possible experience of care [11].

THE PATIENT: A CHANGING ROLE

Recently Hood and co-workers promoted 'P4 Medicine', where in addition to preventive, predictive and personalized medicines, participatory medicine has also been included [12[•]]. This is giving a new and key role to the patient, whose 'participation' in context is not anymore limited to the obsolete "patient/doctor" interaction as such. P4 Medicine also means the active participation of the patients in the entire health process till an active involvement in clinical trials: an innovative approach.

In order to realize this, biomedical advances alone are insufficient: all those psychological aspects that make 'unique' each person, should be integrated with the clinical, biological and genetic data [13]. Gorini and Pravettoni [14] proposed a more complex approach, highlighting the opportunity to add a fifth 'P' (psycho-cognitive) to the P4 Medicine: thus, reaching both the biological and the psychosocial dimensions.

Agusti *et al.* [15] proposed a novel strategy for the management of airway diseases aimed to overtake the limits of the traditional diagnostic classification that tends to make 'stereotypical' the patient. They suggested a bottom-up approach (from endotypes to disease phenotypes) instead of a classical top-bottom approach (from symptoms to mechanisms). In this perspective, the identification of 'treatable traits' (those traits that are identifiable and treatable according the current knowledge) in each patient permits to apply a targeted strategy of disease management. This personalized approach takes into account physical, psychological and behavioral features.

INHALER DEVICES AND ADHERENCE TO TREATMENT

A key aspect to be considered, whenever treating patients with asthma, is the adherence to treatment [16[•]]. Official recent data from the Italian Regulatory Authority reported adherence to treatment in obstructive pulmonary diseases, including asthma, as 13.8% of the expected (http://www.aifa.gov. it/content/luso-dei-farmaci-italia-rapporto-osmed-gennaio-settembre-2016; accessed 8th July 2017). On the other hand, adherence has been always indicated as a weak point in chronic disease management [17], with a dramatic economic impact on disease burden.

In asthma, a critical role is played by the inhaler devices, quite a few on the market [18] and it is a specific task of the specialist to know all the features on each inhaler and to choose the most appropriate one accordingly to the single patient characteristics. This key issue has been extensively described and discussed in the European Respiratory Society (ERS)–International Society for Aerosols in Medicine (ISAM) Task Force document published in 2011 [19]. Additionally, to the correct choice of the device, the prescriber should consider an

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FIGURE 1. Steps forward to the best patient's evaluation.

educational training to the patient about the device use. In fact, Virchow et al. [20] reported a very low capability of the patients (about 21%) to use properly inhalers just after reading the written instructions, whereas a specific educational intervention can rise the correct use up to 52%. Being this the case, educational intervention is a key element for a correct treatment of asthma patients [21]. A specific education is also crucial whenever a switch of device is needed [22], as we should avoid multiple inhalers whose prescription is confusing the patient, with the risk of defective adherence [23[•]]. The prescriber should also always properly educate the patient to new devices and check the inhalation patient's technique at each visit, as part of the correct asthma management [23[•],24].

Improvement of adherence has been the aim of several projects [25] involving different stakeholders and technologies. Recent useful approaches have been proposed such as using monitoring tools for inhalers that can be improve adherence [26].

These new technologies will certainly help both the patients and doctors to control adherence and they will also provide further interesting data about asthma management in general.

PRECISION MEDICINE AND PERSONALIZED MEDICINE IN ASTHMA

As already previously stated, the concepts of 'personalized medicine' and 'precision medicine' have been often considered as overlapping. We think that they are not interchangeable but strongly related and depending on each other, being 'Precision Medicine' essential to achieve the goal of personalizing the medical approach [Figs. 1 and 2]. According to the National Institution of Health (NIH) definition, Precision Medicine is 'an emerging approach for disease treatment and prevention that takes into account individual variability in



FIGURE 2. Steps forward to obtain the best patient's treatment.

genes, environment, and lifestyle for each person' (https://ghr.nlm.nih.gov/primer/precisionmedicine/ initiative; accessed 8 July 2017).

The precision medicine approach aims to find predictive parameters to more accurately choose which treatment or prevention strategy is more suitable for a particular disease in a specific group of patients. Therefore, precision medicine is the opposite of the so-called 'one-size-fits-all' approach, in which clinical decisions and therapeutical strategies are applied, without considering the possible individual [1]. Precision medicine implies the ability to classify individuals into subpopulations that differ in their susceptibility to a particular disease.

Two additional approaches are of great relevance in this context: 'choosing wisely' and 'slow medicine' [27], whose aims are the best for the patient while controlling sustainability of the process.

In order to identify subpopulation of patients ('phenotypes' [28^{••}]), the precision-medicine approach looks into the underlying mechanisms of different forms of each disease ('endotypes' [3]) by the use of surrogate measures that acts as biomarkers [29].

The ability to provide precision medicine to patients in routine clinical settings depends on the availability of easily assessable biomarkers, possibly with point-of-care technology [30^{••}], together with the knowledge bases on how correctly translate biomarkers result into a personalized clinical-therapeutical decision. Therefore, we should be aware that a personalized and precise approach, such as in severe asthma, will be managed by few well equipped reference centers in which high-level trained staff works [31,32].

The precision-medicine approach became a priority for clinicians and researchers when, in 2015, also the United States President Barack Obama

stated his intention to fund a 'Precision Medicine Initiative' 'to enable a new era of medicine through research, technology, and policies that empower patients, researchers, and providers to work together toward development of individualized treatments' (https://obamawhitehouse.archives.gov/node/ 333101; accessed 8 July 2017). Such an initiative has been confirmed early this year [33] by the NIH research investment in the next decade.

After the first biologic agent (omalizumab, 37) 10 years ago, recently new biologics have been approved for severe asthma: mepolizumab (a monoclonal antibody against IL-5) and reslizumab (another anti-IL monoclonal antibody). In the next few years, some more will come; benralizumab, an IL-5 receptor antagonist; dupilumab, an IL-4 receptor alpha antagonist, blocking both the IL-4 and IL-13 inflammatory pathways; tezepelumab an antithymic stromal lymphopoietin (TSLP) antibody, we are now at the beginning of a new era in the management of severe asthmatic patients: a 'Precision Medicine Era'. In fact, severe asthma patients necessarily must be clinically, functionally, inflammatorily and molecularly phenotyped in order to personalize the therapeutic approach.

FROM PHENOTYPES TO ENDOTYPES: A ROUNDTRIP

Asthma was previously viewed as a single disease characterized by chronic airway inflammation, bronchial hyperreactivity, airway obstruction and airway remodeling. More recently, this simplistic concept, ignoring the heterogeneity of the disease, became obsolete: nowadays, asthma is considered a multidimensional disease with different clinical, inflammatory, pathologic and physiologic involvements [28^{••},29,30^{••},31,32–35].

Mainly using cluster analysis-based approaches, several asthma phenotypes have been described so far [36^{••}]. The most common phenotypical classification of asthma highlights the type of inflammatory airway involvement, separated in four subgroups according to sputum eosinophilia or neutrophilia: an eosinophilic asthma generally (earlyonset) well responsive to inhaled corticosteroids, a noneosinophilic but neutrophilic asthma generally with more severe disease and less sensitive to inhaled corticosteroids, a mixed eosinophilic-neutrophilic form with features of both the previously described forms and a paucygranular sputum pattern asthma whose existence is still highly debated [37,38]. Subsequently, a more complex way of dividing asthmatic patients in subgroups was developed according to the presence (Th2-high) or not (Th2low) of assessable biomarkers of Th2-mediated airway inflammation [39,40]. The Th2-high phenotype includes mainly the classical allergic one and the late-onset, nonallergic but highly eosinophilic asthma (including the subgroup associated with acetylsalicylic-acid sensitivity); on the other hand, the Th2-low phenotypes are more diversified and less well defined. The existence of other phenotypes has been reported: asthma associated with obesity; a very late-onset asthma without eosinophilic inflammation mainly in women; cigarette smoker asthmatic patients with neutrophilic sputum and a paucygranular asthma phenotype with predominant airway hyperreactivity [41].

These phenotypic, cluster-analysis-based classifications helped clinicians and researchers to better understand that asthma is a multifaceted syndrome.

Additional consideration should be paid to comorbidities, which may negatively influence asthma control, severity and response to therapy (including response to biologics).

Comorbidity could be a real hallmark as it happens to chronic rhinosinusitis with nasal polyps (CRSwNP), which is present in the great majority of late-onset, highly eosinophilic, nonallergic asthma [42]. The presence of CRSwNP, in fact, being associated with more severe, steroid-dependant asthma [43], was seen to increase lower airway inflammation [44], and negatively correlate with asthma control [45]. Another example of relevant comorbidity associated with a specific phenotype of asthma is obesity: there is evidence that overweight and obese asthmatic patients tends to have a less eosinophilic and more neutrophilic airway inflammation with the consequence of a reduced corticosteroid sensitivity, and therefore, a more severe asthma [46]. Weight loss has been associated with restored asthma control and improved asthmarelated quality of life [47].

Phenotypes describe characteristics of asthmatic patients but do not provide insights of the disease causes. Therefore, the term 'endotype' was proposed to indicate a subtype of a condition defined by a specific biologic mechanism [3]. Each endotype may account for one or more phenotype, and vice versa. The study of underlying mechanisms of asthma gave the opportunity to the researchers to identify more precise therapeutic targets [48]. In particular, some inflammatory pathways were identified as relevant in different severe asthma endotypes: a proportion of patients are characterized to have a 'classical' allergic asthma induced by the exposure to relevant (mainly perennial) allergens, resulting in an increased concentration of cytokines as IL-4, IL-5 and IL-13 and IgE production [49]; another relevant proportion of severe asthmatic patients, often associated with CRSwNP, blood and serum

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eosinophilia, and generally without evidence of atopy, is characterized by an intense production of IL-5 and IL-13, which is responsible for the high levels of blood and tissue eosinophils [42]. This endotype seems to be because of the activation of a subset of inflammatory cells, with similar characteristics to Th2 but different in their trigger, called innate lymphoid cells of type 2 (ILC2) [49]. The hypothesis is that a still unknown agent (i.e. virus, bacterium, fungi, pollutants, cigarette smoking, occupational agents, ...) directly damage airway epithelium, inducing the activation of epithelial cells with the production of epithelial cytokines such as TSLP, IL-25 and IL-33, which act as potent activators of ILC2 [50]. 3) A smaller proportion of patients with severe asthma are characterized by a noneosinophilic, mainly neutrophilic airway inflammation, higher frequency of chronic-recurrent airway infections, higher prevalence of smokers or ex-smokers, and higher BMI [51]. In these patients, the TNF α [52] and IL-17 [53] inflammatory pathways are the most studied and more probably etiopathogenetically correlated underlying processes.

BIOMARKERS TO IDENTIFY ENDOTYPES AND PHENOTYPES

The definition of different endotypes and phenotypes implies the need of finding reliable and possibly easily assessable biomarkers that may guide the clinician in order to identify the right therapeutic target for each single endotype, and therefore, for each single patient.

An ideal biomarker should be suitable to identify the disease or a specific endotype/phenotype, and it should appear or disappear over the course of disease progression, and thus, be useful in determining the prognosis of a disease within an individual. Moreover, the ideal biomarker should change as a biomarker-driven therapy is started, adjusted or discontinued, and should be easily obtained with minimum discomfort or risk to the patient (https://www. fda.gov/ohrms/dockets/ac/01/briefing/3798b1_04_ holt/tsld005.htm; accessed 8th July 2017).

In severe asthma, many possible biomarkers have been investigated but only few of them, so far, had at least one of the above-mentioned characteristics and can be easily used in clinical practice.

Briefly, the current most reliable biomarkers are: blood eosinophil count, sputum eosinophils and neutrophils, serum total IgE and exhaled nitric oxide (FE_{NO}) [29]. Among the other molecules that need more studies in order to be validated as biomarkers of severe asthma endotypes-phenotypes, serum periostin and dipeptidil-peptidase 4 (DPP4) seems those with the most promising profile [54,55], for a certain asthma endotype.

The severe allergic asthma phenotype is generally characterized by no or mild blood eosinophilia, high levels of FE_{NO} , and high levels of serum total IgE [49]. This endotype differs from the nonallergic, late-onset, eosinophilic refractory one because the latter is generally characterized by important blood and sputum eosinophilia despite high dose of inhaled and oral or just oral corticosteroid treatment; these patients generally have also high FE_{NO} but serum total IgE can be both normal or elevated but probably with a lower etiopathogenetical importance [42]. Among the eosinophilic refractory asthma patients, some probably have a high IL-5 expression whereas others may have a predominant IL-13-mediated and IL-14-mediated inflammation. Serum periostin has been proposed to identify patients responsive to anti-IL-13 monoclonal antibody treatment [56], but other subsequent studies reduced the enthusiasm on this biomarker [57]. More recently, DPP4 has been proposed to be able to distinguish those endotypes with a predominant IL-13-mediated and IL-4 mediated inflammation from those in which IL-5 is the main key player [55]; further studies are needed to confirm these first evidence.

Interestingly, blood eosinophil count, FE_{NO} and serum periostin seems to be able to predict sputum eosinophilia, with best results for blood eosinophils and serum periostin [57]. The combined evaluation of blood eosinophils and lymphocytes ratio and eosinophils and neutrophils ratio in a mathematical model (the so-called ELEN index [58]) seems to be more precise than blood eosinophil count alone to identify sputum eosinophilia.

This evident overlap between the biomarkers implies the need of using panels of biomarkers instead of single ones [59]: this approach will probably increase the precision in identifying endotypes for a more precise medicine approach to severe asthma management.

As mentioned before, two of the desired characteristics of a biomarker are the easiness and noninvasiveness of assessment; this is also part of the definition of point-of-care testing (POCT). Among the available severe asthma biomarkers, FE_{NO} is the one with all POCT characteristics, being performed in a noninvasive, very easy and rapid way, usually by using a portable device; these features give the opportunity to clinicians to assess the measure directly in their offices during the time of a visit [60[•]]. We recently published the validation of a blood eosinophil count assessed by a portable, POCT and noninvasive device (the system is able to determine this information by analyzing a single drop of

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blood collected with a finger prick) in a severe asthmatic setting [30^{••}]. Similar devices but for the assessment of serum total/free IgE [61] and periostin [62] are being developed and need a validation in a severe asthma setting. We believe that these POCT devices, if applied to the assessment of severe asthma biomarkers, could really accelerate the path leading to a precision medicine approach and clinical management of severe asthma.

BIOMARKER-TARGETED THERAPIES

Biomarkers, in addition to their role in defining phenotypes and endotypes may also have a predictive value for defining responders to each biological treatment.

As far as biomarkers useful to choose the right endotype-based treatment, serum IgE, blood eosinophils were the most studied ones.

Serum total IgE is in fact used for verification that the patient with severe allergic asthma is a suitable candidate for anti-IgE therapy (omalizumab) with total serum IgE levels between 30 and 1500 IU/ml [63].

As far as the available or closely available anticytokine strategies (i.e. anti-IL5, anti-IL4/IL-13, ...) the biomarker chosen to define the possibility to prescribe such treatments is peripheral blood count. Several different cutoffs in blood eosinophil measurements were chosen as minimum values for prescribing biologicals acting in the eosinophilic refractory forms of severe asthma: they range from a eosinophil count of 150/ml for dupilumab [64] (more recent data did not confirm eosinophils as predictive biomarker of response to dupilumab [64]) and mepolizumab (together with at least one historical report of more than 300/ml for the latter) [65] to 300/ml for benralizumab [66] to 400/ml for reslizumab trials [67].

Whenever considering predictive biomarkers of response to biological treatments, a proteomic approach showed that galectin-3 tissue levels were directly correlated with a good response, in terms of reduction of airway remodeling to omalizumab [68,69].

Post hoc analysis of clinical trials, showed that patients with high-serum periostin levels were those with the highest response to lebrikizumab (an anti-IL13 agent) in terms of lung function improvement [56]. Furthermore, as mentioned previously about POCT, high FE_{NO} levels, high-serum periostin and high-blood eosinophil levels positively correlate with improvement in terms of reduction in exacerbations in patients treated with omalizumab [70].



FIGURE 3. Severe Asthma Center Networking would ameliorate the exchange of data and information on severe asthmatic patients.

CONCLUSION

We should definitely put aside the concept of the 'one-size-fits-all' traditional approach [6].

In the last 10 years, only omalizumab was available, later followed by mepolizumab and hopefully by the other monoclonal antibodies previously described.

We will move from the availability of only one monoclonal antibody to a situation in which we will have to choose one monoclonal antibody among many: this implies the need of more selective biomarkers (or panels of them) in order to identify the right biological for each single patient, in a more personalized and precise medicine approach to the disease treatment.

The era of new 'Magic Bullits' to target molecular mechanisms of severe asthma [34] has come and it implies new expertises and professionalities, also because severe asthma precision medicine has a cost [71] and in this context, the leading role of the Severe Asthma Reference Centers is crucial [72[•]]. The potential of networking among the severe asthma centers is of great relevance [73^{••},74], as collecting data, big data, will provide real life and research evidences capable of driving the process to the best practice (Fig. 3).

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Conflicts of interest

G.W.C. has been member of advisory board, speaker, scientific meeting for GSK, Teva, Sanofi, Roche,

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