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CHAPTER 8

GENERAL  
DISCUSSION AND  
IMPLICATIONS  
OF THE  
RESEARCH

*This thesis consists of two parts; section 1 contains chapters on biomarker applicability; section 2 describes the early clinical development of a new anti-asthmatic drug.*

## SECTION I

### BIOMARKER DEVELOPMENT AND EVALUATION

**CHAPTER 2 & 3** Inhalation of allergens has been effectively and safely applied as a tool to evaluate new drug treatments for asthma, including inhaled steroids, anti-leukotriene therapy, and biologics including anti-IgE and anti-TNF- $\alpha$  therapies [1]. The main application of these models in early drug development has been to estimate treatment changes in pulmonary function assessed by FEV<sub>1</sub>, bronchial hyper-responsiveness (BHR) assessed by methacholine challenge, and changes in inflammatory state as assessed by cell counts after inhaled allergens.

*Chapter 2* investigated an extension of the allergen challenge model with the ability to quantify changes of a broad range of specific TH<sub>2</sub> cytokines in post allergen challenge collected sputum.

**THE VALUE OF THIS STUDY FOR CURRENTLY DEVELOPED ASTHMATIC DRUGS** ✱ The research pipeline for asthma contains a number of potentially new anti-inflammatory compounds and therapies with a need for PD airway biomarkers [2], *Table 1*. The study in chapter 2 describes the combined use of ultracentrifugation with multiplex quantification methods of sputum biomarkers following allergen challenge, reversed by fluticasone. This approach allows for non-invasive identification of pharmacodynamic targets for anti-asthma therapies. In the study, the most notable effects, in terms of reproducibility and the reversal effect of fluticasone, were observed for IL-5, IL-13, eotaxin-3 and TARC.

Interleukin 5 is a key inflammatory cell mediator in the pathogenesis of asthma as it promotes the proliferation, differentiation, recruitment and survival of eosinophils [15]. High eosinophil counts in sputum are associated with poor asthma control, can predict future exacerbations, and can help direct medication changes. These findings suggest that IL-5 may be an attractive target in allergic asthma. Mepolizumab is a monoclonal antibody treatment directed against IL-5, and Mepolizumab treatment has resulted in a decrease in asthma exacerbations in patients with severe eosinophilic asthma [12]. Benralizumab, a monoclonal antibody that targets  $\alpha$  chain of the IL-5 receptor, has shown promising results in phase 2 trials [16].

Interleukin 4 and IL-13 are functionally and structurally related. Both induce B cells to produce IgE. Lebrikizumab is an IL-13 monoclonal antibody. Improvements

were seen in patients with high levels of serum periostin, a protein produced by bronchial epithelial cells. This suggests that lebrikizumab may be useful for a specific group of patients with allergic asthma [13]. The compound is now in phase 3 trials. Dupilumab inhibits both IL-4 and IL-13 signaling and shows promising results in early clinical studies [14].

Eotaxin-3 and TARC are both chemokines for which no specific compound has been designed so far.

The allergen challenge model provides useful information on the ability of potential controller therapies to block the asthmatic response. The model also offers high negative predicting values for effectiveness. Indeed, many - but certainly not all - novel compounds with anti-inflammatory characteristics have been studied using the allergen challenge to assess the efficacy of the compound in allergic asthma with FEV<sub>1</sub> as the main outcome parameter [14]. Few novel compounds however, have used the allergen challenge focusing on a panel of TH<sub>2</sub> driven cytokines rather than individual TH<sub>2</sub> cytokines, resulting in a more comprehensive insight in the pharmacodynamic effects of investigated compounds [17]. This is curious, since expectations are that this approach will result in a more robust predictive assessment of anti-inflammatory asthma medication in the early phase clinical drug development. The study described in chapter 2 contributes to this approach by determining the reproducibility and reversal effects of fluticasone of a wide range of cytokines and chemokines after allergen challenge in allergic asthmatic patients.

*Chapter 3* presents the results from microarray analysis of sputum samples obtained from the study described in chapter 2. Little is known about microarray assessment of gene expression in induced sputum obtained after allergen challenge in allergic asthmatics. It was found that microarray technology yields reproducible levels of gene expression in induced sputum following inhaled allergen challenge. Moreover, a refined allergen induced gene signature was effectively reversed by fluticasone and individual signatures correlated with lung function and eosinophil measurements. It appears that this approach has the potential to further elucidate the complex interactions occurring during an inflammatory condition.

**LIMITATIONS TO THE ALLERGEN CHALLENGE** ✱ Asthma is a heterogeneous disorder consisting of several ill-defined endotypes. Nevertheless, adult asthmatic patients are likely to fall within 5 clusters, [18-20], *Table 2*.

The allergen challenge model triggers a TH<sub>2</sub> pathway inflammation. Initially, mast cell fixed IgE leads to the immediate phase of the asthma attack. In the late phase, which is in essence an progressive inflammatory reaction, TH<sub>2</sub> play a prominent role [21]. TH<sub>2</sub> inflammation however, is not a hallmark for all clusters of asthma

limiting the generalizability of the allergen challenge model, *Table 2*. Other limitations include the need of experienced and well trained staff and the difficulty of recruiting eligible subjects.

**CHAPTER 4** Several biomarkers sampling methods are often used within the same clinical trials, in the same individual on the same day and sometimes even repeatedly. Most of them are validated and have proven importance in drug development studies. But what if the complementary use of research tools causes them to interact with each other?

Two frequently used sampling methods in asthma drug development studies are sputum induction (s<sub>1</sub>) with hypertonic saline and fractional exhaled nitric oxide (F<sub>ENO</sub>). Both methods assess the airway inflammation for the lower airways in response to e.g. new anti-inflammatory therapies. In *chapter 4* the effect of s<sub>1</sub> on F<sub>ENO</sub> measurements was investigated in healthy chronic smoking and non-smoking subjects. The conclusion is that hypertonic saline decreased F<sub>ENO</sub> levels in both groups. As a result of the observed interference of s<sub>1</sub> with F<sub>ENO</sub> measurements, the latter should always be measured before the s<sub>1</sub>. There are several explanations for this observation: repeated spirometry measures as part of the s<sub>1</sub> are known to result in decreased F<sub>ENO</sub> level [22]; mucus accumulation resulting from the s<sub>1</sub> procedure could inhibit NO diffusion from bronchial epithelial cells [23]; and osmotic changes in bronchial epithelium evoked by inhalation of hypertonic saline could have an effect on the inducible isoform of NO synthase (NOS) [24].

The effect of s<sub>1</sub> on F<sub>ENO</sub> concentrations was already known; it was studied in healthy volunteers [25], in children [26] and in asthmatics [22]. Data demonstrating this effect in a group of chronic smokers was scarce. Therefore this study confirms and extends previous observations.

Besides the direct effect of s<sub>1</sub> procedures, smoking itself also has an effect on baseline F<sub>ENO</sub> concentrations. Indeed, in our study the F<sub>ENO</sub> levels were generally lower compared to the non-smokers group. In a more recent study, Hillas and colleagues showed that F<sub>ENO</sub> levels were significantly lower in asthmatic smokers compared to non-smokers [27]. It would be interesting to assess the observed interaction between s<sub>1</sub> and F<sub>ENO</sub> in a group of asthmatic smokers in a future study.

**CHAPTER 5** One of the hallmarks for biomarkers is that they must be reliable and reproducible. *Chapter 5* evaluates the reproducibility of soluble markers obtained from sputum induction and serum in a group of chronic smokers and a group of non-smokers. Soluble markers in blood and sputum samples were obtained on 2 separate visits and analyzed with ELISA. The results were somewhat disappointing.

Although it was found that chronic smokers had higher concentrations of IL-8 in induced sputum and Pulmonary Surfactant Associated Protein D (SP-D) in serum, only SP-D out of a set of markers fulfilled the predefined criteria for reproducibility.

**ASSESSMENT OF THE FEASIBILITY OF BLOCKING IL-8** ✱ Although this study was carried out in a group of healthy chronic smokers, smoking plays a prominent role in the pathophysiology of COPD, and is associated with ~80% of the cases. In smokers, the disease is characterized as an inflammation that involves increased numbers of CD8+ (cytotoxic) Tc lymphocytes. These cells, together with neutrophils and macrophages, release inflammatory mediators like IL-8. [28;29]. Increased concentrations of IL-8 have been found in sputum and broncho alveolar lavage (BAL) of patients with COPD. IL-8, also known as neutrophil chemotactic factor or CXCL8, is regarded as a pro-inflammatory factor. It induces chemotaxis in target cells, primarily neutrophils but also other granulocytes, causing them to migrate toward the site of infection. IL-8 also induces phagocytosis once they have arrived. Activation of the IL-8 receptor, CXCR1 and CXCR2, has similar effects. Despite the biological plausibility, several CXCR2 antagonists and IL-8 specific antibodies have been largely ineffective in clinical studies in patients with COPD. As an example, the CXCR2 antagonist SB-656933 reduces sputum neutrophils in patients with COPD. Larger studies however, have shown no clinical benefit [30;31].

It is difficult to pinpoint the reason for these failures. However, as is the case for asthma, numerous mediators are involved in the pathophysiology of COPD. It thus appears that blocking a single mediator is unlikely to provide a major clinical benefit, unless it is found to be solely responsible for orchestrating a cascade of inflammation. This also highlights that the pathophysiology of COPD is poorly understood. Therefore, it is unclear which cells could be promising therapeutical targets. It thus remains crucial to increase understanding of the underlying inflammatory mechanisms of COPD, and to identify biomarkers of disease activity that predict the clinical efficacy of anti-inflammatory treatments. In this respect, analysis of sputum and serum parameters as described in *chapter 5* may be useful.

**ASSESSMENT OF THE FEASIBILITY OF SP-D** ✱ The story is different for Surfactant Pulmonary Associated Protein D (SP-D) than for IL-8. The molecule is produced predominantly in the lungs and has been recognized as an important regulator of innate immune system, capable of binding pathogens and facilitating phagocytosis [32]. It has microbicidal effect on certain bacteria and fungi and promotes the elimination of viruses [33-35]. The lower levels of SP-D found in broncho alveolar lavage fluids in smokers may thus be a cause of weakened lung immunity.

In addition, SPD is able to suppress inflammatory responses caused by lipopolysaccharide (LPS) [36]. SP-D is one of the few lung specific proteins that can be measured in serum, bypassing sometimes difficult and inconvenient sampling methods like BAL and sputum induction. Furthermore, COPD and COPD-exacerbations are associated with higher levels of serum SP-D, which due to some unknown mechanism is inversely correlated with BALF SP-D [37].

In our study we found that levels of serum SP-D in chronic smokers are up to twice as high compared to the levels of non-smokers. In addition, SPD proved to be the only marker of a set of biomarkers to be reproducible in both smokers and non-smokers. Moreover, it has been found that regular inhalation of salmeterol and fluticasone lowers serum SP-D levels in COPD patients [38].

Taken together, SP-D has an important role in the innate immunity, can be easily assessed in serum, is specific for COPD, levels change with severity and exacerbations, and last but not least it is sensitive to effective treatment. This does not necessarily mean that a SP-D compound should be developed. However, it does seem that SP-D may be a potential candidate marker with characteristics of a validated biomarker for COPD.

**LARGE SCALE PROTEIN DISCOVERY STUDIES** ✱ The search for multiple disease specific protein biomarkers is sometimes called “a fishing expedition”, an uncomplimentary term indicating that scientists have no idea what the catch will be. *Chapter 5* describes the results of identification of a broad array of analytes in sputum using ELISA technology. However, newer analytical technologies like multiplex analysis and peptidomics have exponentially increased the surface of the net allowing more fish to be caught. In fact these two technologies were also applied in the study, but not described in the manuscript.

In our study, Multiplexed analysis (Rules Based Medicine) was used to measure a total of 92 soluble markers in serum. The analysis showed increased levels of Macrophage Derived Chemokine (MDC), Carcino-Embryonic Antigen and Apolipoprotein H in smokers compared to non-smokers. All other biomarkers were not significantly different for the two study groups. Reproducibility was not assessed and therefore this analysis was not mentioned in the article.

For proteomics analysis, mass spectrometry MALDI-TOF-TOF was used to assess the feasibility for primary screening of yet to be discovered biomarkers. Once a peptide marker is identified and its sequence validated, the next step is to develop a standard ELISA assay. The ELISA can then be applied to the same clinical samples and deliver quantitative measures of the biomarker. In our study, proteomics analysis identified 147 proteins in the supernatant of induced sputum demonstrating

the feasibility of the mass spectrometry MALDI-TOF-TOF as a tool for biomarker discovery in induced sputum and plasma.

What would the advantage of such technologies be? Although still in their infancy compared with other methodologies, large scale protein discovery studies could result in protein fingerprints which potentially could stratify patients for the treatment of COPD and may provide a tool to assess treatment response [39;40]. However it is unlikely that these findings can be used plausibly or reproducibly as markers for the activity of a new medicine unless they are properly validated and underpinned by a mechanistic understanding of the changes in these profiles. If this is not done the risk of statistically spurious findings is probably high.

## SECTION 2

### CLINICAL STUDIES IN HEALTHY VOLUNTEERS, ALLERGIC RHINITIS AND ALLERGIC ASTHMA; WITH USE OF BIOMARKER

In the second part of this thesis, the focus was on allergic asthma and the biomarker assisted development of the new anti-asthmatic drug RPL554.

Asthma and chronic obstructive pulmonary disease (COPD) are commonly treated with a combination of a bronchodilator and an anti-inflammatory drug, usually an anti-inflammatory glucocorticosteroid. A drug that combines clinically relevant symptom control with substantial anti-inflammatory activity in one molecule would therefore be very welcome.

Phosphodiesterases (PDE's) are a large family of II enzyme subtypes, modulating intracellular concentrations like cAMP. Increased levels of cAMP are related to airway smooth muscle relaxation and anti-inflammatory actions. PDE3 inhibitors cause airway smooth muscle relaxation in vitro [41;42], bronchodilation in vivo [43], and increases in forced expiratory volume in 1 second (FEV1) in patients with asthma [44]. PDE4 inhibitors inhibit the function of a wide range of inflammatory cells in vitro and have pronounced anti-inflammatory effects in patients with inflammatory airway disease [45;46]. Experimental data suggest that combined inhibition of PDE3 and PDE4 iso-enzymes may have synergistic effects [47]. RPL554 is a dual PDE3 and PDE4 inhibitor that has shown bronchodilator and anti-inflammatory effects in animals in vitro and in vivo [48].

**CHAPTER 6** describes an experiment in which the effects of nebulized RPL554 were assessed in a series of adaptive proof of concept studies. Both its bronchodilator

and anti-inflammatory properties were investigated for the first time in humans. This single dose study was split into three stages. Safety of RPL554 was assessed in healthy individuals (stage 1), and safety and efficacy was assessed in otherwise healthy subjects with allergic asthma or allergic rhinitis (stage 2 & 3).

This experiment showed that RPL554 was well tolerated and had mild adverse events that were equal to placebo. In allergic asthmatic patients RPL554 at 0,018 MG/KG resulted in an increase in PC20Mch of around 1,5 doubling doses and substantial bronchodilation at doses between 0.009 and 0.072 MG/KG with a mean increase at 1 hour of 520 mL in FEV1. The latter effect is comparable to that of salbutamol, a widely used  $\beta$ -agonist. Increases in eosinophil count after the nasal allergen challenge were reduced by RPL554- compared to placebo (7.1%) although this did not reach statistical significance.

**CHAPTER 7** In another study with RPL554, described in *chapter 7*, the reproducibility of the bronchodilator response to a daily dose of nebulized RPL554 at 0,018 MG/KG for 6 consecutive days in a placebo controlled study in 12 healthy men with allergic asthma was examined. This study showed that RPL554 had a similar maximum mean increase on day 1 (555mL) day 3 (505 mL) and day 6 (485 mL), suggesting that acute tolerance to the effect of RPL554 did not develop.

**RPL554'S FURTHER DEVELOPMENT FOR ASTHMA** ✱ As explained in the introduction of this thesis, the underlying pathophysiology of asthma is often – but not always – TH2 driven and a consequence of the contribution of numerous cells including the epithelium, dendritic cells, T lymphocytes, eosinophils, mast cells and airway smooth muscle. Many of these cells express PDE4 and inhibition of this enzyme could suppress the function of these cells [49].

When a new pharmaceutical agent for asthma is studied, its ability to block the early and late phase asthmatic response and subsequent airway hyperresponsiveness is important evidence for effectiveness [1]. So far, a few PDE4 inhibitors have been assessed for their effect on the acute allergen induced bronchoconstriction. An inhibition of the late phase response of around 30% was observed for CDP840 and roflumilast after several days of treatment [50;51]. In addition, the effect of GSK256066, an inhaled PDE4 inhibitor, was regarded as modest [52]. However, roflumilast showed a 30–50% reduction in sputum eosinophil number indicating anti-inflammatory activity in asthma. However, the effect of roflumilast on allergen induced airway hyperresponsiveness was limited; around 1 doubling dose of methacholine [53]. These effects of roflumilast are modest when compared to the effect of fluticasone (*chapter 2*), and consistent with the lack of demonstrable action of PDE4 inhibition on mast cell [54].

Anti-inflammatory effects of RPL554 in the studies presented in *chapter 6 en 7* were difficult to establish. PC20MCH increased with 1,5 doubling dose compared to placebo. On the other hand, methacholine challenge only has moderate correlation with eosinophilic airway inflammation [55]. Also, RPL554 did not show a significant effect on eosinophil cell count in patients with allergic rhinitis after a nasal allergen challenge, (*chapter 6*). Furthermore, RPL554 at a dose of 0,018 MG/KG continued to induce sustained bronchodilation in a group of mild allergic asthmatic volunteers, but it did not improve FEV1 values at baseline at each study day, (*chapter 7*).

In conclusion, the bronchodilator effects of RPL554 are promising. However, its anti-inflammatory effects in asthma based on the observations in our studies and based on the limited effects of PDE4 inhibitors on mast cells in general, will need to be determined in further clinical studies.

**RPL 554 FOR COPD** ✱ COPD is another inflammatory disease but the nature of the inflammatory response is different from asthma. COPD is characterized amongst other reactions, by activation of macrophages and cells recruitment of neutrophils in the lung. Lipopolysaccharide (LPS) challenges mimic the airway neutrophilia and have been used to evaluate the efficacy of new therapeutic agents for the treatment of COPD. The inhibiting effect of RPL554 (0,018 MG/KG) on LPS-induced sputum neutrophils was demonstrated in a placebo-controlled crossover trial with 21 healthy men [7], showing reduction in absolute numbers of neutrophils after 6 hours. The effect was similar to that observed with roflumilast in the sputum of patients with COPD [46].

In addition, an open-label, placebo controlled crossover trial, in 12 men with mild-to-moderate COPD showed a bronchodilator effect of RPL554 (0,018 MG/KG) with a mean maximum FEV1 increase of 17,2% [7]. These effects are comparable with peak effects reported in such patients with inhaled  $\beta_2$  agonists [56], and suggest potential therapeutic use of RPL554 in COPD.

## OVERALL CONCLUSION

This thesis describes attempts to identify novel pathophysiology based biomarkers for chronic airway disease by applying newer detection techniques. It appears that such an approach may be worthwhile to pursue. Further, it shows that application of biomarkers allows effective development of new therapies in chronic airway disease.

**Table 1** Summary of potential new therapies for asthma.

|                                       |  |
|---------------------------------------|--|
| Ultra long acting $\beta_2$ -agonists | Effective for COPD, possible available in near future for asthma - especially when combined with ICS [3;4] |
| Long acting muscarine-antagonist      | Effective for COPD, possible available in near future for asthma - especially when combined with ICS [5;6] |
| PDE-Inhibitors                        | Favorable early studies [7]  |
| Bronchial thermoplasty                | Effective but has possible immediate complications associated with the                                     |
| TNF- $\alpha$                         | Risk of side effects [9]   |
| IL-9                                  | Not effective [10]   |
| IL-17                                 | Not effective [11]   |
| IL-5                                  | Effective in asthmatic patients with high eosinophil levels [12]   |
| IL-13                                 | Effective in asthmatic patients with high periostin levels [13]  |
| IL-4                                  | Promising early studies [14]   |

**Table 2** Clusters of asthma

|           |  |
|-----------|--|
| Cluster 1 | Early onset allergic asthma with Th2 signature: high levels of eosinophils, mast cells, IGE and FENO                                   |
| Cluster 2 | Adult onset non allergic asthma. Th2 signature with eosinophilia and high levels of IL-4, IL-5, IL-13, but absence of allergic disease |
| Cluster 3 | Exercise induced asthma. Mast cells play an important role   |
| Cluster 4 | Prominent obesity and minimal Th2 response   |
| Cluster 5 | Sputum neutrophilia with Th17 response   |

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