Asthma and Genetic Factors

Asthma is common chronic inflammatory disease of the airways characterised by variable and attacks of cough and breathlessness, usually precipitated by an environmental trigger (air pollution, cold, dry air, smoke, etc). The prevalence and severity for allergic asthma have increased markedly in the last several decades. Dysregulated expression patterns of pro- and anti-inflammatory mechanisms are thought to be responsible for the development of chronic inflammation. The sequence of immunopathogenesis is unclear but there is clearly a genetic predisposition. Asthma is caused by multiple interacting genes, some having a protective effect and others contributing to the disease pathogenesis, with each gene having its own tendency to be influenced by the environment. By the end of 2010, 100 genes had been associated with asthma in six or more separate populations, including GSTM1, IL-1 (α,β), IL-1RN, TSLP-R, IL-1R1, IL-8RA, IL-3,4,5,9,10,12,13, NAT2, CTLA-4, SPINK5, V-CAM-1, TNF-α, ARG1, GSTM1, A3AR, CHIA, LELP1, TGFβ1, SOD-1, EGFR, GPRA, CCR2, FceRIβ, PHF11, ACE, IRAK-3, C6D9, IL-18, MUC-2, eNOS; NOS3, CMA1, and ADAM33, among others [1-3]. The risk of developing asthma has a strong genetic component, with estimated heritability ranging from 35 to 85%. Asthma is described as a complex disease arising from the contribution of multiple genetic and environmental factors [4,5]. Some of these genes may also be involved with other phenotypes such as helminthic infections (FceRIβ and IL-4), COPD, cardiovascular diseases, congenital thrombotic thrombocytopenia, Crohn’s disease (ADAM33) renal cell carcinoma, blood malignancies (PHF11), tuberculosis (TB), hyperparathyroidism, prostate cancer, insulin dependent diabetes mellitus, leprosy and chronic hepatitis B infection (vitamin D receptor:VDR) [2]. Thymic stromal lymphopoietin (TSLP) is an epithelial-derived cytokine similar to IL-7, whose gene is located on chromosome 5q22. 1 and it exerts its biological function through the TSLP-Receptor (TSLP-R). TSLP is expressed primarily by epithelial cells at barrier surfaces such as the skin, gut and lung in response to danger signals. TSLP genetic variants and its dysregulated expression have been linked to atopic diseases such as atopic dermatitis, asthma, allergic rhinitis and eosinophilic esophagitis [6].

The response to drug treatment in asthma is a complex trait and is markedly variable even in patients with apparently similar clinical features. Pharmacogenomics, which is the study of variations of human genome characteristics as related to drug response, can play a role in asthma therapy. Both a traditional candidate-gene approach to conducting genetic association studies and genome-wide association studies have provided an increasing list of genes and variants associated with the three major classes of asthma medications: β2-agonists, inhaled corticosteroids, and leukotriene modifiers. Moreover, a recent integrative, systems-level approach has offered a promising opportunity to identify important pharmacogenomics loci in asthma treatment [7].

The polymorphisms within the ADRB2 gene that are potentially associated with obesity and asthma include Arg16Gly and Glu27Gln, and probably to some extent Thr164Ile, as well. The pleiotropic nature of ADRB2 makes it a good candidate for such an association. It also has an established role in the development of both conditions separately. The underlying mechanism seems to depend on both the alternations in lung function and the metabolic effect associated with ADRB2. However, it could also be linked to immunological functions related to ADRB2 expressions on leukocytes [8].

A genome-wide association study (GWAS) identified a previously unknown asthma-susceptibility locus on chromosome 17q21, harboring the adjacent genes ORM13L3 (ORM13L spingolipid biosynthesis regulator 3) and GSDMB (Gasdemin B). This genetic association has been confirmed in ethnically diverse populations, and gene–environment interactions have been detected between susceptibility alleles and exposure to cigarette smoke and furry pets. Acevedo et al. showed that, significant differences in the DNA methylation levels of the ORM13L3promoter of asthmatic children, independent of age, gender, genotype and differential leukocyte cell counts, which might partially explain the increased ORM13L3 expression observed in cases. Their results


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strongly support the role of both genetic and epigenetic factors contributing to asthma susceptibility in the 17q21 locus [9,10].

Omalizumab

Omalizumab, a humanized mAb that binds to the CH3 domain, near the binding site for the high-affinity type-I IgE Fc receptors of human IgE, can neutralize free IgE and inhibit the IgE allergic pathway without sensitizing mast cell and basophils. Omalizumab, which has been conceptualized for treating IgE mediated allergic diseases and approved for treating patients with severe persistent allergic asthma in many countries, can neutralize IgE, impede the IgE allergic pathway, and render mast cells and basophils insensitive to activation through IgE/FcεRI. In addition to asthma, Omalizumab has been investigated in various other conditions including chronic urticaria (CU), perennial and seasonal allergic rhinitis (AR), pruritic bullous pemphigoid, latex allergy, peanut allergy, idiopathic anaphylaxis, hyper-IgE syndrome, chronic rhinosinusitis, interstitial cystitis, aspirin sensitivity, mastocytosis, eosinophilic gastroenteritis and atopic dermatitis. Most patients with chronic urticaria have an autoimmune cause: some patients produce IgE autoantibodies against autoantigens, such as thymoperoxidase or doublestranded DNA, whereas other patients make IgG autoantibodies against FceRI, IgE, or both, which might chronically activate mast cells and basophils. In the remainder of patients with CU, the nature of the abnormalities has not yet been identified. Accumulating evidence has shown that IgE, by binding to FcεRI on mast cells bearing the receptor, making those cells insensitive to the stimulation by incoming allergens [16-19].

In a patient with an allergic disease caused by type I hypersensitivity toward specific external antigens, omalizumab induces multifactorial therapeutic effects. Omalizumab depletes free IgE in the blood and interstitial space and inhibits IgE binding to FcεRI on basophils, mast cells, and dendritic cells. Omalizumab cannot bind to IgE that is already bound to FcεRI and does not have a direct effect on FcεRI levels. However, the depletion of free IgE results in the downregulation of FcεRI on cells bearing the receptor, making those cells insensitive to the stimulation by incoming allergens [16-19].

Most asthmatic individuals respond satisfactorily to inhaled corticosteroids and β-adrenergic agonists; however, 5-10% of them have severe, persistent symptoms that respond poorly to such treatment. The introduction of Omalizumab as an add on therapy for inadequately controlled moderate-to-severe or severe persistent allergic asthma (SPA) provided a valuable new treatment option for patients. Given the importance of anti-inflammatory therapy for control of SPA, it is important to determine the effects of omalizumab on markers of inflammation. The interaction between Omalizumab and free IgE interrupts a key step in the allergic inflammatory cascade, preventing IgE from binding to mast cells, basophils, and dendritic cells, and down-regulating IgE receptor expression on these inflammatory cells thereby inhibiting degranulation and the release of inflammatory mediators [19-22]. Omalizumab has been approved in over 120 countries for treating patients with SPA. These pharmaceutical developments have validated the IgE pathway as an effective therapeutic target for treating IgE-mediated allergic disease [23-25].

For the very first time, we used omalizumab in symptomatic therapy of recurrent laryngeal oedema attacks in a patient with post operative pulmonary carcinoid tumor for 4 months. During the 3 years of follow-up no recurrences was noted in tumor.

Omalizumab effects on sApo-2 L and allergen specific immunotherapy

Tumor necrosis factor related apoptosis-inducing ligand (TRAIL: Apo-2L) is used as a marker for apoptosis. TRAIL (Apo-2L) is a transmembrane (type II) glycoprotein belonging to the TNF superfamily. The extracellular domain of TRAIL is homologous to that of other family members and shows a homotrimeric subunit structure. Like TNF and FasL, sApo-2L also exists physiologically in a biologically active soluble homotrimeric form. An increase in eosinophil levels has been reported in allergic asthma, and is thought to reflect an increase in peripheral blood eosinophil survival promoted by Apo-2 L.

In our previous study we showed that soluble Apo-2 L levels in patients with severe persistent allergic asthma decreased after anti-IgE treatment using omalizumab. These results suggested that sApo-2L may act as a soluble effector molecule, and that the decrease in levels after omalizumab treatment may allow us to use this marker to monitor clinical improvement. Combination therapy with omalizumab and specific subcutaneous immunotherapy (SCIT) in patients with severe persistent asthma also suggest that omalizumab is an effective therapy in such individuals. Omalizumab reduces serum IgE levels and FcεRI receptor expression on key cells in the inflammatory cascade. The consequences of these processes are the inhibition of the release of inflammatory mediators from mast cells, and diminished recruitment of inflammatory cells, especially eosinophils, into the airways [21,26-28].

Allergen Specific Immunotherapy (SIT) has the advantage of being the only causal treatment of allergic controlled asthma and rhinitis but is fraught with the dangers of severe systemic or local side effects and anaphylaxis [29-32]. Omalizumab can possibly overcome these limitations by binding exclusively to circulating IgE molecules and reducing the levels of circulating IgE regardless of allergen specificity by binding to the constant region of circulating IgE molecule. This prevents free IgE from interacting with the high- and low-affinity IgE receptors (FcεRI and FcεRII) on mast cells, basophils, macrophages, dendritic cells, B lymphocytes, and subsequently leads to a decrease in the release of the mediators of the IgE mediated allergic response, namely, cytokines, histamines and leukotrienes [33,34].

The first clinical trial looking for the clinical effects of a combined therapy of SIT and Omalizumab was performed in grass- and birch-pollen allergic children and adolescents in Germany. Kuehr and
colleagues recruited 221 children and adolescents to evaluate the efficacy and safety of Omalizumab with SIT on birch pollen induced allergic rhinitis (AR) [35]. SIT plus Omalizumab-treated subjects were reported to have a 48% reduction in allergen-induced symptom load over two pollen seasons independent of the allergen. Furthermore, rescue medication use, number of days with symptoms and symptom severity were significantly lower in the SIT plus Omalizumab groups compared with SIT alone. A post hoc sub-analysis of this study to assess the effects of each treatment (SIT or Omalizumab) demonstrated that SIT alone did not significantly reduce either symptoms severity score [36]. Hence, combination therapy may be complimentary, providing the superior effect compared to individual treatments. Recently, there have been trials of Omalizumab and treatments in patients with AR and co-morbid asthma. In the trial by Kopp and colleagues, a significant reduction of 40% in symptom load was observed in favor of SIT plus Omalizumab compared with SIT alone (p = 0.04) [37]. Another study showed that the tolerability of SIT after pretreatment with Omalizumab or placebo in patients with symptomatic asthma was not adequately controlled with inhaled corticosteroids. A total of 13.5% of patients treated with Omalizumab showed systemic allergic reactions to SIT compared to 27% in those receiving placebo (p = 0.017). More patients were able to reach the target maintenance SIT dose (p = 0.004) in the Omalizumab group compared to placebo [38,39], suggesting that pre-treatment with Omalizumab was associated with fewer systemic allergic reactions to SIT and enabled more patients to achieve the target immunotherapy maintenance dose.

Studies in allergic rhinitis and asthma have shown that pre-treatment with Omalizumab may be an effective option to reduce systemic anaphylactic reactions and achieve a higher dose of allergen immunotherapy in a safe way. This can be of specific relevance to hymenoptera venom immunotherapy. Although there are no controlled trials, there are case reports of anti-IgE therapy with Omalizumab reducing the risk of systemic reaction during induction of venom immunotherapy in patients who have either failed treatment or in those with mastocytosis [40-42].

Omalizumab effects on oxidative stress markers, vitamin-D and homocysteine

An imbalance between oxidative stress and anti-oxidative capacity may play an important role in the development and progression of bronchial asthma (BA) and chronic obstructive pulmonary disease (COPD). The systemic oxidant-antioxidant status changes during exacerbation versus stable periods in patients with BA and COPD. During an exacerbation period of BA, despite the decreases in glutathione peroxidase (GSH-Px), glutathione reductase (GRd) and melatonin levels, malondialdehyde (MDA) and catalase (CAT) levels, and the white blood cell count, the percentage of eosinophils is significantly higher than in the stable period. MDA and superoxide dismutase (SOD) values are higher in the exacerbation period than in the stable period although GSH-Px, GRd, melatonin, pH, and pO₂ values are lower in the exacerbation period than in the stable period. Blood counts and respiratory function tests were reported not to change between exacerbation and stable periods in patients with COPD. Thus episodes of BA or COPD might be associated with elevated levels of oxidative stress.

A decrease in NO during omalizumab therapy was also previously described by Silko et al. [44]. Down regulation of ET-1 in EBC significantly correlates with a decrease in the markers of allergic (and eosinophilic) inflammation, such as NO, ECP or blood eosinophil counts, as well as increase in spirometric indices. These changes were observed after 16 weeks of therapy. A follow-up observation performed after 52 weeks of treatment revealed a further significant fall in ET-1 concentrations in EBC; however, the improvement of other markers of allergic inflammation was less pronounced. This could indicate that anti-IgE therapy has its greatest influence on eosinophilic inflammation during the first 16 weeks of therapy. Nevertheless the effects of many other immunological mechanisms related to remodeling, as well the known action and interactions of ET-1 observed in the first period of treatment, are thought to continue over time. This suggests that longer-term anti-IgE therapy with omalizumab in asthmatic patients could significantly limit the development of inflammation and bronchial structural changes.

In our previous study we investigated changes in total antioxidant capacity in asthmatic patients treated with Omalizumab. Our data suggested that ongoing therapy with Omalizumab, already proven to be clinically effective in treatment of severe allergic asthma. Anti-IgE therapy is an innovative and promising treatment modality that mediates its effects in part at least through decreased inflammation following improved anti-oxidant capability. In turn, our study was suggesting that measuring of the latter may prove to be useful surrogate markers to monitor efficacy of treatment in patients suffering from this disease [46].

Alternatively, the development of atopy may also be a direct effect of elevated homocysteine or some of its metabolites, which appears to exert a number of diverse effects on immune function. In addition, total homocysteine (Hcy) has been shown to increase in response to immune activation and cell proliferation during a non-allergic Th1-type immune response. Although much less is known about the health effects of sustained post load homocysteine concentrations, there is evidence that it has negative effects on platelet aggregation and endothelial function. A number of studies have indicated that homocysteine may contribute to the development and progression of atherosclerosis, a risk factor for cardiovascular diseases. However, the mechanisms by which Hcy can induce vascular dysfunction are not fully understood [48-53].

Vitamin-D (25(OH)D) has effects on the innate and adaptive immune system. 25(OH)D levels are associated with poor asthma control, reduced pulmonary function, increased medication intake and exacerbations. Little is known about 25(OH)D in adult asthma patients or its association with asthma severity [54,55]. More than that, 25(OH)D triggers a Hcy metabolizing enzyme and data from the Longitudinal Aging Study Amsterdam suggested a correlation between 25(OH)D status and Hcy levels [56]. The decrease in Hcy concentrations and increase in 25(OH)D also supports the possible vascular endothelial protection mechanism.

Omalizumab effects on pruritic bullous pemphigoid

Bullous pemphigoid (BP) is an acquired, autoimmune, bullous
disease that is characterized by autoantibodies against the 230-kDa bullous pemphigoid antigen within basal keratinocytes and the 180-kDa type XVII collagen within the basement membrane zone lying between the epidermis and dermis. In addition to skin blisters, patients with BP often experience pruritus and erythematous urticaria-like skin lesions.

CD200 (OX-2) is a novel immune-effective molecule, both cell membrane-bound and also existing in a soluble form in serum (sCD200, sOX-2), which acts both as a pro-inflammatory through its receptor [11,14,47]. In our previous study, we reported a patient who had a pruritic bullous pemphigoid and very high levels of total IgE (5000 kU/L) who was refractory to the aggressive immunosuppressive regimens for bullous pemphigoid but responded rapidly to systemic anti-IgE. The circulating level of sOX-2 was 48.45 pg/mL in serum and 243 pg/mL in blister fluid. Soluble OX-2 levels were higher in blister fluid than in serum. During the second month of follow-up, the patient’s sOX-2 level decreased to 26.7 pg/mL. Clinical improvement was demonstrated as histologically re-epithelialization. Optimal treatment modalities need to be clarified in such situations. After the second round of omalizumab (300 mg), frequency of exacerbations decreased and after 13th round it was completely disappeared [58].

Reduction in serum levels sOX-2 with anti-IgE treatment suggests that OX-2 could be pro-inflammatory [21,24,47,57,58]. Soluble OX-2 might also play a role in immune response in the pathogenesis of autoimmune and inflammatory skin disorders [60,61].

Omalizumab effects on coagulation pathway

More interestingly, as in some of our cases we have observed, one with severe persistent asthma (SPA) patient who had protein C/S deficiency history, multiple massive pulmonary embolus and systemic subacute thrombosis determined in vena saphena parva and in left vena perforantes cruris, underwent Omalizumab treatment. After a long term (20 month) treatment with Omalizumab, he had a decreased fractional exhaled nitric oxide concentrations (FENO), severe lung infection are seen. During the second month of follow-up, the platelet count fell down to 55,000/mm³, the omalizumab treatment was suspended for 4 weeks until the platelet count rose up to 100,000/mm³ [23], (Figures 1 and 2).

Omalizumab effects on hyperimmunoglobulin-E syndrome, eosinophilic gastroenteritis, mastocytosis

Hyperimmunoglobulin E syndrome (HIES), is a heterogeneous group of immune disorders. It is characterized by very high concentrations of the serum antibody IgE. Clinically eczema-like rash, cold staphylococcal infection, severe lung infection are seen. An IgE level greater than 2,000 IU/mL is often considered diagnostic, except patients younger than 6 months of age. Extrinsic pathway of coagulation is activated in response to high level of circulating IgE [69,70]. Abnormal neutrophil chemotaxis due to decreased production of interferon gamma by T lymphocytes is thought to cause the disease. Both autosomal dominant and recessive inheritance have been described [69]. Mutations in molecules DOCK8 have been associated with syndromes that share many features with classical autosomal dominant HIES, which is inherited by an autosomal recessive trait and tend to have a milder clinical picture [70,71]. STAT3 is a key regulator of many immunologic pathways. It is involved in the signal transduction of many cytokines, including but not limited to IL-6, IL-10, IL-21, IL-22, and IL-23 [72]. Animals with a myeloid-specific deletion of STAT3 leads to up-regulation of many Th1 cytokines, such as IFNγ and TNFα, and down-regulation...
of pro-inflammatory and anti-inflammatory responses regulated by IL-6 and IL-10, respectively [70,71]. These cytokines are critical to differentiation of TH17 cells, which are important in inflammatory response to bacterial and fungal pathogens. It was reported that both STAT3 mutation-positive and STAT3 mutation-negative HIES exhibited a profound deficit in TH17 differentiation [73]. Several studies reported clinical improvement in patients with severe atopic eczema with high serum IgE level [73-75].

Eosinophilic gastroenteritis (EGE) is characterized by patchy or diffuse eosinophilic infiltration of any part of gastrointestinal (GI) tract [76,77]. Eosinophils are normally present in gastrointestinal mucosa, but deeper infiltration and more than 30 eosinophils per high-power field in at least five areas are pathologic [78]. Since GI tract is frequently face with external allergens via ingested foods, allergens from food pass the mucosa and trigger an inflammatory response that lead mast cell degranulation and recruit eosinophils. Tissue damage is caused by cytotoxic proteins contained in the cytoplasmic granules of eosinophil. In addition to tissue eosinophil, eosinophils can also mediate proinflammatory effects, ie up-regulation of adhesion systems, modulation of cell trafficking, releasing chemokines (eotaxin), lipid mediators and leukotriene. Eosinophil recruitment into the tissue is regulated by a number of inflammatory cytokines, ie IL-3, IL-4, IL-5, IL-13, granulocyte macrophage colony stimulating factor (GM-CSF) and T helper 2 (Th2) cytokines. A Th2-type immune response seems to be involved in both IgE and non-IgE mediated EGE [79]. Anti-IgE treatment with Omalizumab is associated with a 35–45% drop in peripheral blood eosinophil count as well as decrease in duodenal and antral eosinophil count [80,81]. It also effectively blocks CD23 mediated allergen binding to B cells. But some reports failed to demonstrate in vivo immunomodulatory activity on T cell responses [82].

Mastocytosis is a heterogeneous disorder that results from clonal...
Omalizumab effects on chronic urticaria

Improvement in clinical outcomes. Omalizumab-treated group. Patients on anti-IgE therapy had an randomized, placebo-controlled clinical trial. In addition, they had strikingly decreased levels of TSLP, OX40L, TARC (involved found symptomatic improvement with omalizumab [13]. Recently, case reports investigating anti-IgE therapy in patients with AD (AD) need further research in lowering serum IgE subjects. Several mechanisms of Omalizumab in the treatment of atopic dermatitis compared with the nasal polyp patients without Samter’s syndrome a significantly higher inducible nitric oxide synthase activity when treatment [24]. Nasal polyps from patients with Samter’s triad had our study, no change was seen on nasal polyposis after Omalizumab and functionality of local IgE formation in the airways [92], but in of nasal polyps with comorbid asthma, supporting the importance surgery. Omalizumab demonstrated clinical efficacy in the treatment of nasal polyps with comorbid asthma, supporting the importance and functionality of local IgE formation in the airways [92], but in our study, no change was seen on nasal polyposis after Omalizumab treatment [24]. Nasal polyps from patients with Samter’s triad had a significantly higher inducible nitric oxide synthase activity when compared with the nasal polyp patients without Samter’s syndrome [93].

Omalizumab Effects on atopic dermatitis

Severe refractory atopic dermatitis is a chronic, debilitating condition that is associated with elevated serum IgE levels. The mechanisms of Omalizumab in the treatment of atopic dermatitis (AD) need further research in lowering serum IgE subjects. Several case reports investigating anti-IgE therapy in patients with AD found symptomatic improvement with omalizumab [13]. Recently, Iyengar et al. [84] showed that all patients receiving Omalizumab had strikingly decreased levels of TSLP, OX40L, TARC (involved in Th2 polarization) and interleukin-9 compared to placebo in their randomized, placebo-controlled clinical trial. In addition, they found a marked increase in IL-10, a tolerogenic cytokine, in the Omalizumab-treated group. Patients on anti-IgE therapy had an improvement in clinical outcomes.

Omalizumab effects on chronic urticaria

Metz et al. [90], assessed responder rates, optimal dosage, response to up-/downdosing, time to relief of symptoms, rates of return and time of relapse after omalizumab administration, and safety in 51 CU patients, 20 with chronic spontaneous urticaria (CSU) alone, 21 with different forms of chronic inducible urticaria (CindU) and 10 with both in their clinical analysis. They showed that Omalizumab was a rapidly acting, highly effective and safe drug in CSU and CindU patients in their clinical experience from more than 1250 injections in those patients over four years indicates. Their observations in a real life clinical setting support the recommendation of current EAACI/ GA2LEN/EDF/WAO guideline for the management of urticaria to use Omalizumab on the treatmenof urticaria patients [91].

The activation of mast cells and their release of inflammatory mediators are regarded as the “final common pathway” for a myriad of proinflammatory factors, including those involved in the various types of urticaria [99,100].

The clinical response of urticaria to H1-antihistamines and the finding of increased concentrations of histamine in skin tissue fluid underscore the role of histamine derived from dermal mast cells as a major mediator of urticaria. Although highly unlikely, it cannot be excluded that in some cases of urticaria, the primary abnormality lies in the mast cells themselves. If this were the case, it would be likely that the condition would be systemic rather than confined to the skin. Therefore it is more likely that skin mast cells in patients with urticaria are not intrinsically abnormal but become increasingly sensitive or “unstable” or activated as the result of certain abnormal factors present in their surroundings. Although there are many nonimmunologic factors that might influence mast cell function in the skin, such as components of the complement system and neuropeptides, particularly those related to stress, because this review is primarily concerned with the mechanisms by which omalizumab might be effective, nonimmunologic factors will not be considered in detail [101-106].

In conventional thinking the involvement of IgE in mast cell activation requires the cross-linking of FcεRI-bound IgE by antigen or anti-IgE antibodies. This initiates the aggregation of FcεRI, leading to tyrosine kinase activation and subsequent mast cell activation for secretion. However, in 2001, it was suggested independently by 2 groups that monomeric IgE in the absence of antigen can have multiple effects in murine mast cells, including differentiation, proliferation, survival, and mediator and cytokine generation. These effects, which involve the binding of IgE to FcεRI and the aggregation of FcεRI, occur without the mast cells undergoing degranulation. The finding that monomeric IgE can augment mast cell activity has been confirmed by studies using various techniques. In a transcriptome analysis of 8793 genes, sensitization of mast cells with monoclonal IgE alone, without FcεRI cross-linking, was found to upregulate 58 genes more than 2-fold compared with their levels in unsensitized mast cells. These genes included those for cytokines, such as IL-1β, IL-6, and colony-stimulating factor 1; chemokines, such as CXCL8, CCL4, and CCL7; and cytokine and chemokine receptors. The genes for various immune regulators, adhesion molecules, antiapoptosis proteins, and cytoskeletal elements, such as RAS protein activator like 1 and fibroconnectin leucine-rich transmembrane protein 2, were also upregulated [106-108].

First, omalizumab sequesters monomeric IgE to reduce its priming effect on mast cells. This would be particularly relevant if HC IgE is involved in the pathogenesis of urticaria. Second, in patients with IgG autoantibodies against IgE or FcεRI, the depletion of mast cell–bound IgE by omalizumab and the subsequent downregulation of FcεRI on mast cells and basophils would lead to their decreased state of hyperexcitability. Third, in those patients with IgE autoantibodies against autoallergens, the inhibition of IgE binding to FcεRI by omalizumab and the downregulation of FcεRI would represent a central mechanism of omalizumab [101]. Other studies have followed the suggestion that mouse monoclonal IgE molecules are heterogeneous with respect to their ability to induce survival and activation events in mast cells [109].

Omalizumab and cardiovascular safety

There have been concerns about the cardiovascular safety in patients initiating Omalizumab therapy, because of the most recent study that analyzed the association between Omalizumab and arterial thrombotic events [96-98,110]. We showed that in one of our patient, Doppler ultrasonography did not reveal any thrombus after anti-IgE therapy and the patient did not require lung transplantation and that serum protein S/C levels increased to normal ranges. Exercise stress testing was normal and after initiation of anti-IgE treatment, neither cranial emboli event nor any neurologic complications did occur. Patient did not report any cardiac arrhythmias after initiation of anti-IgE therapy. Besides, exercise stress testing was normal, while the patient was treated with anti-IgE. Aneurysm enlargement or complications were not detected during the treatment with anti-IgE [21].

Omalizumab effects on Diabetes mellitus

The clinical experience during the patient follow-up of omalizumab treated severe persistent allergic asthma patients with type-2 diabetes mellitus is introduced. Omalizumab is generally considered safe. The most common adverse reaction from omalizumab is injection site pain and bruising but the package insert contains warnings regarding malignancies, geohelminth infections, and a “black box” warning that serum protein S/C levels increased to normal ranges. Exercise stress testing was normal and after initiation of anti-IgE therapy. Besides, exercise stress testing was normal, while the patient was treated with anti-IgE. Aneurysm enlargement or complications were not detected during the treatment with anti-IgE [21].

based on post marketing surveillance data and reported such cases indicating that different side effects may occur beyond 2 hours after the injection. Patients with diabetes mellitus should be informed that such a need of insulin dose should be increased due to the possible effect of omalizumab on blood glucose level. In these two patients half of the recommended dosage was given and blood glucose levels were controlled.

Omalizumab is a humazon recombinant anti-IgE monoclonal antibody approved for therapeutic use both in adults and in children aged 6–12 years with severe allergic asthma. The coexistence of severe asthma refractory to the conventional pharmacological approach and sensitization to at least one perennial allergen represent the current indications for Omalizumab prescription. Its efficacy and safety as an add-on therapy is sustained by several data coming from both clinical trials and real-life experiences [111-121] and showing a significant reduction of yearly exacerbation-rate.

To sum up what I would like to express as a conclusion is that, Omalizumab in patients with severe persistent asthma is an effective therapy for asthma and co-morbid conditions (CU, bee venom allegy, latex allergy, atopic dermatitis, food allergy, Samters syndrome) just like I have mentioned above. The mechanism of action of Omalizumab in the treatment of asthma is believed to be multifactorial.

References


13. Yalcin AD, Bisgin A, Gorcynski RM (2012) IL-6, IL-10, TGF-β and GCSF levels were increased in severe persistent allergic asthma patients with the anti-IgE treatment. Mediators Inflamm 2012: 2012.


47. Yalcin AD, Genc GE, Bisgin A. Evaluation of Homocysteine, 25(OH) Vitamin D, Pro-inflammatory IL-1β and Immune Modulator OX-2 Levels in Moderate Allergic Asthma Patients: Association with Biological Treatment and Disease Activity. Jour Clin Pharm and therapy ID: 2366.


