



# Review FcERI: A Master Regulator of Mast Cell Functions

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**Abstract:** Mast cells (MCs) perform multiple functions thought to underlie different manifestations of allergies. Various aspects of antigens (Ags) and their interactions with immunoglobulin E (IgE) cause diverse responses in MCs. Fc $\epsilon$ RI, a high-affinity IgE receptor, deciphers the Ag–IgE interaction and drives allergic responses. Fc $\epsilon$ RI clustering is essential for signal transduction and, therefore, determines the quality of MC responses. Ag properties precisely regulate Fc $\epsilon$ RI dynamics, which consequently initiates differential outcomes by switching the intracellular-signaling pathway, suggesting that Ag properties can control MC responses, both qualitatively and quantitatively. Thus, the therapeutic benefits of Fc $\epsilon$ RI-targeting strategies have long been examined. Disrupting IgE–Fc $\epsilon$ RI interactions is a potential therapeutic strategy because the binding affinity between IgE and Fc $\epsilon$ RI is extremely high. Specifically, Fc $\epsilon$ RI desensitization, due to internalization, is also a potential therapeutic target that is involved in the mechanisms of allergen-specific immunotherapy. Several recent findings have suggested that silent internalization is strongly associated with Fc $\epsilon$ RI dynamics. A comprehensive understanding of the role of Fc $\epsilon$ RI may lead to the development of novel therapies for allergies. Here, we review the qualitatively diverse responses of MCs that impact the attenuation/development of allergies with a focus on the role of Fc $\epsilon$ RI toward Ag exposure.

Keywords: allergy; antigen; IgE; FccRI; mast cell; desensitization



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# 1. Introduction

Recently, allergies have emerged as a public health problem, as over one-quarter of the population in industrialized countries is affected and the prevalence of allergies is increasing [1]. Pathophysiological features of allergy are characterized by a diverse set of clinical symptoms, and the disease burden reflects the long-term consequences of chronic allergic inflammation at sites of persistent or repetitive exposure to antigens (Ags) [2]. Different phenotypes/endotypes have been recognized with the recent increase in the heterogeneity of Ag contents [3].

Mast cells (MCs) initially respond rapidly to exogenous Ags, which is the principal initial effector immune response involved in modulating allergic inflammation [2,4,5]. Ag-induced MC degranulation is the basis of anaphylaxis and other severe allergic reactions [6]. Ag binding to immunoglobulin E (IgE) that is bound to its receptor (FcɛRI) causes the crosslinking of IgE–FcɛRI complexes, resulting in the release of biologically active mediators such as histamine, serotonin, and leukotrienes within minutes. Cytokines, chemokines, and growth factors that are transcriptionally upregulated in MCs are secreted over a period of hours after initial MC activation, leading to prolonged inflammation [7,8].

Fc $\epsilon$ RI, the high-affinity IgE receptor, distinguishes Ag–IgE interactions and drives cellular allergic responses [9]. Fc $\epsilon$ RI is a unique molecular target that initiates different functional outcomes of MC responses and allergic inflammation [10,11]. In this review, we highlight recent advances in the understanding of Fc $\epsilon$ RI as a master regulator of MC functions, with a focus on Fc $\epsilon$ RI behavior in response to various Ags and subsequent cellular responses, including desensitization, which could be a potential strategy for alleviating allergies.

## **2. Fc**εRI

# 2.1. Crosslinking of the IgE–FceRI Complex with Antigen

The high-affinity IgE receptor, Fc $\epsilon$ RI, is primarily expressed on MCs, basophils, and Ag-presenting cells, and mainly exists as the heterotetramer  $\alpha\beta\gamma2$ . However, there are differences among species; an alternate trimeric form  $\alpha\gamma2$  is expressed on human, but not rodent. The  $\alpha\beta\gamma2$  consists of a single-membrane-spanning  $\alpha$ -subunit that contains the IgE-binding domain, a tetra-spanning  $\beta$ -subunit that contains a single immunoreceptor tyrosine-based activation motif (ITAM), and two  $\gamma$ -subunits that exist as a disulfide-linked homodimer and also contain one ITAM each. The importance of the  $\alpha$ -subunit in the Fc $\epsilon$ RI-mediated allergic reaction was demonstrated by the absence of allergic reactions in  $\alpha$ -subunit-deficient mice [12]. The Fc $\epsilon$ RI $\alpha$  binds to the Fc fragment of IgE at a 1:1 ratio to form the IgE–Fc complex. Two hydrophobic regions in C $\epsilon$ 3 of the IgE–Fc complex asymmetrically bind to two Ig-like extracellular domains of the  $\alpha$ -subunit. This asymmetric interaction follows a 1:1 stoichiometry, and both distinct integrations increase the binding avidity, leading to high-affinity binding [13–15].

Multivalent Ags cause cross-linking of the IgE–Fc $\epsilon$ RI complex, triggering signaling cascades that result in the release of preformed mediators stored in granules. In response to Ag, Lyn kinase phosphorylates the ITAMs of Fc $\epsilon$ RI $\beta$  and Fc $\epsilon$ RI $\gamma$ , which initiates a complex signaling cascade involving a series of membrane-associated and cytoplasmic signaling molecules [8,16–18]. Furthermore, bindings of monomeric IgE to Fc $\epsilon$ RI, in the absence of Ags, is capable of inducing phosphorylation of signaling molecules and MC responses (MC survival and granules maturation) without degranulation [19].

### 2.2. FceRI Clustering: Size, Stability, and Mobility Dynamics

In the cell membrane, the IgE–Fc $\epsilon$ RI complex diffuses freely under resting conditions. Multivalent Ag binding to IgE leads to the reorganization of  $Fc \in RI$  into large aggregates on the cell surface within seconds to minutes, which causes a transition in the receptor from a diffuse state to a highly immobile state [20]. This behavior of the Fc $\epsilon$ RI cluster, including its mobility, kinetics, and size, has been highlighted as a potential feature that governs signal initiation. The relationship between the  $Fc \in RI$  cluster size and mobility has been clearly demonstrated in rat basophilic leukemia (RBL-2H3) cells and bone marrow-derived MCs (BMMCs). The imaging studies using quantum dot (QD)-based probes revealed that the small aggregates remain mobile on cell surface, whereas there is abrupt immobilization in large aggregates, where the actin cytoskeleton plays an important role in cross-link-induced immobilization of Fc $\epsilon$ RI [21,22]. And rews et al. showed that Ag-induced Fc $\epsilon$ RI immobilization was required for receptor internalization, but not for signal transduction, and they presented evidence that FceRI immobilization could cause signal termination. Although the small  $Fc \in RI$  aggregates induced by sub- and supra-optimal concentrations/Ag valency did not lead to strong degranulation,  $Fc \in RI\beta$  phosphorylation was detectable, indicating that signaling was competent. They concluded that the size of receptor clusters regulates mobility, signaling competence, and  $Fc \in RI$  internalization [22].

The same group also investigated the spatial/density regulation of the Fc $\epsilon$ RI clusters of MC, and they demonstrated that the Fc $\epsilon$ RI redistribution in regions of the membrane after Ag stimulation is also remarkable for the complexity of the Fc $\epsilon$ RI dynamics and signaling initiation. Sequential images of IgE–Fc $\epsilon$ RI complexes, assessed by total internal reflection fluorescence (TIRF) microscopy, have been used to observe the dynamics of receptor redistribution upon contact with the ligand. Importantly, Carroll-Portillo et al. demonstrated that small mobile clusters coalesced into large, cholesterol-rich patches that occupied the central portion of the contacting membrane and are likely composed of smaller aggregates in the same region, suggesting that dense populations of mobile receptors could initiate low-level degranulation [23]. These findings highlight the importance of Fc $\epsilon$ RI dynamics not only in the temporal behavior of internalization but also in membrane topography. Furthermore, they investigated the topographical details of IgE–Fc $\epsilon$ RI complexes on the plasma membrane at the nanoscale level using high-resolution microscopy [24]. They observed a coalesced large patch, termed a signaling patch, and confirmed signaling competency downstream of the patch [24]. These findings contribute to the understanding of the sophisticated signaling mechanisms associated with Fc $\epsilon$ RI and seem to be indicative of relationships between receptor diffusion, clustering, signal initiation, and MC activation. Additionally, a quantitative study was performed to measure Fc $\epsilon$ RI motion within clusters and coalesced patches using single-particle tracking, the results of which provided significantly support for a mechanism whereby Fc $\epsilon$ RI clusters diffuse and then coalesce to form large patches in specific regions of the plasma membrane [25].

These examples illustrate the importance of FccRI dynamics and subsequent cellular responses at a low Ag concentration or valency, which in turn drive the formation of small and mobile  $Fc \in RI$  clusters that were previously considered a simple and diminished (i.e., weak) response. Signaling diversity is significantly related to  $Fc\epsilon RI$  dynamics. Srchomology-2-containing inositol phosphatase (SHIP) is a negative regulator of MCs [26], and its relationship with small  $Fc \in RI$  cluster formation has been proposed [27]. Mahajan et al. performed experiments with a new structurally defined trivalent ligand, DF3, and clearly showed the significant participation of SHIP1 in the small  $Fc \in RI$  cluster, which forms under low-concentration and low-valency conditions [27]. Furthermore, the signaling consequences of differential FceRI aggregation were investigated through continuous stimulation of RBL-2H3 cells with DF3. The results revealed that Syk and SHIP1 drove rapid short-lived positive signals and slower long-lived negative signals, respectively [28], and time-dependent signaling following stimulations that likely operate during differential  $Fc\epsilon RI$  cluster initiation [28]. These timescale observations further strengthen the view that FcεRI dynamics may regulate signaling events and initiate cellular outcomes. Recent findings from Bag et al. demonstrated a direct relationship between the mobility of small or large clusters and the formation of lipid raft domains [29]. Importantly, they found that stabilized liquid ordered (Lo)-like nanodomains around the  $Fc \in RI$  cluster (which are accessible by Lyn kinase, but not transmembrane phosphatase), provide essential spatial filtration that augments Lyn binding and FccRI phosphorylation, while suppressing dephosphorylation by phosphatases [29].

Those studies were performed using concentration- and valency-adjusted agents. The Ag properties and their interactions with IgE are important for determining  $Fc \in RI$  dynamics and the initiation of subsequent cellular reactions, as discussed below in Section 2.3.

#### 2.3. Antigen Properties: Concentration, Valency, and Affinity

Naturally encountered allergens are typically complex and structurally heterogeneous proteins with multiple epitopes [30]. Moreover, naturally occurring IgEs display the heterogeneity of antibodies with different affinities against Ags [31]. In a recent study, naturally occurring pollen allergens (not synthetic allergens) were studied to demonstrate the complexity of allergen sources and investigate their interaction with IgE [32]. Earlier studies identified a variety of olive pollen allergens [33] and house dust mite allergens [34,35], whose variations are recognized as isoallergens. These characteristics contribute to the overall complexity of allergens and the variety of associated disease symptoms. The magnitude of the MC responses indeed depends on the allergen properties. The concentrations, valencies, and affinities of Ags affect the interaction with IgE and MC responses, which may cause complex allergic responses.

The influence of the Ag concentration on Fc $\epsilon$ RI dynamics has been elucidated; degranulation in MCs displays a remarkable bell-shaped response with increasing Ag concentrations [36,37]. This phenomenon is a particular feature of Fc $\epsilon$ RI and differs from many other receptors that reach a plateau phase in response to high ligand concentrations. The suppressed responses at supra-optimal concentrations have been suggested to be not due to reduced Fc $\epsilon$ RI crosslinking [38,39]. There is a Fc $\epsilon$ RI-mediated active turn-off mechanism in the descending portion of the specific bell-shaped degranulation responses in MCs (reviewed in [40]). Indeed, MC responses to supra-optimal Ag concentrations are not simply weaker than those at the optimal Ag concentration. Ca<sup>2+</sup> mobilization in MCs, which results from intracellular  $Ca^{2+}$  release from the endoplasmic reticulum (ER) and subsequent extracellular Ca<sup>2+</sup> influx, is important for degranulation. Comparing the intracellular Ca<sup>2+</sup> responses under optimal and supra-optimal concentration stimulation demonstrated markedly different  $Ca^{2+}$ -mobilization patterns [36]. In particular,  $Ca^{2+}$  release from the ER was slightly stronger at a supra-optimal Ag concentration. The supra-optimal Ag concentration resulted in a transient increase in intracellular  $Ca^{2+}$ , whereas the optimal Ag concentration showed a sustained pattern [36], suggesting that active negative regulation occurs. Consistently, the results of that study also revealed that protein tyrosine phosphorylation was stimulated slightly more with a supra-optimal Ag concentration the optimal Ag concentration. This negative signaling was mainly mediated by the activation of SHIP by Lyn [36]. However, the precise mechanisms of SHIP recruitment remain under investigation. One accepted mechanism is that  $Fc \in RI\beta$  can interact with SHIP1 via its unique ITAM sequence upon supra-optimal stimulation [17,41]. The results of another recent study demonstrated that stimulating IgE-sensitized MCs with a supra-optimal Ag concentration induced FcyRIIb tyrosine phosphorylation in an IgG-independent manner [42]. Intriguingly, the co-aggregation of  $Fc\epsilon RI$  and  $Fc\gamma RIIb$  was associated with a submembranous F-actin meshwork that negatively interfered with MC activation at supra-optimal Ag concentrations [42]. Taken together, these data suggest that tightly controlled cellular reactions may reflect a complementary mechanism that activates or compensates for allergic responses to different intensities of stimulation.

The Ag valency is also an important factor in controlling the signal intensity as well as Ag concentration. Naturally occurring allergens are the most common complex type of multivalent proteins. Despite the small size of the timothy grass pollen allergen, Phl p 5, it is known to be highly allergenic and has multiple independent IgE epitopes [32,43]. In experimental settings, multivalent agents such as DNP-BSA are commonly used to crosslink IgE–FccRI to enhance MC activation as a potent stimulation. The valency and concentration-dependent characteristics (size and mobility) of  $Fc \in RI$  clusters have been demonstrated, as discussed above [22,24]. Typically, high-valence Ags rapidly form large and immobile FccRI clusters. Monovalent Ags generally cannot form aggregates with FceRI, but can stimulate FceRI mobilization/redistribution and form a distinct kind of Fc $\epsilon$ RI patch, termed a signaling patch [23]. Monovalent Ags are capable of competitive inhibition that rapidly influences the binding of multivalent ligands to receptors; thus, monovalent Ags are considered candidates for the rapeutic interventions targeting Fc $\epsilon$ RI. A previous study involving transmission electron microscopy and real-time TIRF imaging of FccRI in monovalent Ag-stimulated RBL-2H3 cells primed with fluorescent-labeled IgE showed significantly different  $Fc \in RI$  membrane topographies between multivalent and monovalent ligands [23]. In that study, the authors determined the Ag mobility by engaging anti-DNP IgE on the surfaces of cells presenting either a monovalent DNP-embedded lipid bilayer (mobile ligand) or a chemically crosslinked multivalent DNP (immobilized ligand). Immobile multivalent DNP<sub>24</sub>-BSA stimulation resulted in the rapid formation of stable, moderately sized clusters with typical levels of RBL-2H3 degranulation. The mobile monovalent ligand showed a high density of mobile receptors, in which thousands of mobile receptors were densely packed in the centralized region of the ventral membrane, despite the lack of direct crosslinking [23]. With regard to the un-crosslinked IgE–Fc $\epsilon$ RI bearing, several studies using IgE clones indicated/discussed mechanisms underlying the effect of IgE in the absence of Ags [44,45], which have been proposed that the IgE–Fc $\epsilon$ RI could be clustering and internalize without Ag, suggested that the importance of Fab regions of IgE for triggering the IgE-mediated robust activation of MCs (reviewed in [46]). Although the mechanism of un-crosslinked internalization of  $Fc \in RI$  is still being investigated, such studies will likely be found to have important implications for further characterization of the size, mobility, and density of  $Fc \in RI$  clusters.

FcεRI functionally distinguishes differences in the affinity of IgE antibodies for different Ags. Naturally occurring polyclonal IgEs show heterogeneity with different affinities against Ags [47]. Differences in binding affinities are also known as differences in binding kinetics. Low-affinity Ags dissociate IgE from FccRI more rapidly than high-affinity Ags. A comparison of MC responses to a high-affinity Ag (dinitrophenyl (DNP)-Ag) and a low-affinity Ag (nitrophenyl (2NP)-Ag) demonstrated that the quantitative difference in the affinities between the IgE and Ag determines the quality of subsequent cellular reactions [48]. The effects of DNP-Ag and 2NP-Ag were compared at concentration that elicited similar Fc $\epsilon$ RI phosphorylation levels. Although both Ags elicited similar degrees of Fc $\epsilon$ RI phosphorylation, the behavior of  $Fc \in RI$  clusters (size, mobility, and distribution) and MC secretory responses (cytokines and chemokines) differed significantly. Live-imaging analysis of FcERI dynamics revealed that 2NP-Ag induced slowed mobility of FcERI clusters, whereas DNP-Ag treatment resulted in rapidly formed  $Fc \in RI$  clusters. These differences in FceRI dynamics were found to be related to distal signaling and functional responses. DNP-Ag elicited a more robust linker of activated T cells 1 (Lyn-Syk-LAT1)-dependent signaling, which led to the release of inflammatory cytokines (e.g., TNF- $\alpha$ , IL-6, and IL-13). In contrast, 2NP-Ag dominantly released chemokines (e.g., CCL2, CCL3, and CCL4) through a pathway that depended on the Src-family kinase Fgr and linker for activation of T cell family member 2 (Fgr-LAT2). A mouse model of passive cutaneous anaphylaxis showed a greater magnitude of skin inflammation with DNP-Ag than with 2NP-Ag. DNP-Ag-stimulated MCs recruited neutrophils to the site of inflammation, whereas 2NP-Ag-stimulated MCs recruited monocytes/macrophages (reviewed in [49]). The role of differential recruitment of neutrophils or monocytes/macrophages remains under investigation, the clarification of which may represent a considerable advance in understanding the regulation of systemic allergic inflammation. The mechanisms responsible for different responses to DNP-Ag or 2NP-Ag, together with other evidence pertaining to  $Fc \in RI$  dynamics, suggest that lowaffinity Ags-stimulated FccRI clusters might remain mobile/diffuse and can induce MC responses (Figure 1).



**Figure 1.** Characteristics of antigen (Ag)-dependent control of clustering of the high-affinity IgE receptor (Fc $\epsilon$ RI) and subsequent mast cell (MC) responses. The characteristics of Ags (i.e., concentration and affinity) regulate Fc $\epsilon$ RI functions. High-affinity Ags elicit robust bridging between immunoglobulin E (IgE) and large immobile clusters with increasing Ag concentration, resulting in typical MC activation. In contrast, low-affinity Ags likely allow Fc $\epsilon$ RI to remain mobile and diffuse, leading to slower internalization (even at high Ag concentration) within Fc $\epsilon$ RI signalosomes. These differential dynamics of Fc $\epsilon$ RI may represent a characteristic feature that translates Ag properties into differential outcomes.

This information may help in developing novel strategies for controlling Fc $\epsilon$ RI clustering in MCs and the subsequent allergic responses. A recent report by Mahajan et al. demonstrated that not only the valency and concentration of an allergen, but also the Fc $\epsilon$ RI occupancy with IgE, help set the thresholds for MC effector responses [50]. High-affinity IgEs for multimeric Ags effectively induce degranulation. In contrast, low-affinity IgEs can mediate effector responses via the avidity effect of polyvalent Ags [32]. A complicated input is likely governed by Ags and IgEs, whereas Fc $\epsilon$ RI may dynamically respond to the stimulation intensity and translate it into an appropriate response. Many researchers have investigated signaling events and MC secretory responses in response to high affinity antigen with different concentrations. Although there is still being uncovered roles of low affinity Ags appears to alter MC responses and disease outcome.

#### 2.4. Perspective

FcεRI displays diverse dynamics (i.e., receptor-cluster size, mobility, distribution, and naturally occurring allergens) and IgEs show heterogeneous properties. The functional responses of FcεRI may be studied to decipher the characteristics of endogenous Ags and/or their interactions with IgEs, and to regulate MC-dependent allergic responses, thereby enabling further elucidation of the clinical relevance of human allergic responses.

## 3. FccRI: Therapeutic Potential and Benefits

## 3.1. Inhibition of IgE–FceRI Binding

FccRI has been targeted for clinical benefits to inhibit or attenuate MC degranulation, and effective clinical approaches for suppressing IgE-dependent MC activation have emerged [51,52]. Omalizumab, a humanized anti-IgE monoclonal antibody that binds to the C $\epsilon$ 3 domain of IgE, can prevent the binding of circulating IgE to F $\epsilon\epsilon$ RI, is a common and preferred treatment (especially for chronic urticaria), and is effective against asthma [53–55]. Omalizumab binds free IgE, disrupts IgE–Fc $\epsilon$ RI complexes, and decreases Fc $\epsilon$ RI expression [56,57]. IgE binding to  $Fc \in RI$  further increases expression of the receptor, resulting in stable Fc $\epsilon$ RI expression on the MC surface [58,59]. Omalizumab does not crosslink FceRI-bound IgE on the MC surface because of conformational changes in FceRI-bound IgE that mask omalizumab-binding sites [54,60]. Omalizumab could dissociate/remove IgE from  $Fc\epsilon RI$ ; however, only poor disruptive activity has been observed [61,62]. Although the precise mechanisms involved in stripping IgE from FccRI remain to be elucidated, allosteric destabilization and facilitated dissociation of the IgE-FccRI complex have been suspected as potential mechanisms [63,64]. Additionally, omalizumab has been studied as an adjuvant therapy for oral immunotherapy (OIT) against food allergies [65]. Open-label pilot studies in cow's milk and peanuts allergy suggested that 9-12 weeks of omalizumab therapy could facilitate rapid oral desensitization to each allergen in a high-risk patient [66,67]. The mechanisms of the combined therapy of OIT plus omalizumab have been studied. Recent study suggested that the combined therapy promotes allergen desensitization through an initial omalizumab-dependent acute depletion of allergen-reactive effector T cells, and which is followed by an increase in allergen-specific regulatory T (Treg) cell activity due to the reversal of their T helper 2 (Th2) cell-like program [68]. Regarding other antibody antagonism/antibody neutralization, ligelizumab is another potent humanized anti-IgE monoclonal antibody that binds to free IgE with a higher affinity than omalizumab. Ligelizumab binds to the C3 domain of IgE, but unlike omalizumab, it can also bind to IgE-bound to CD23, a low-affinity IgE receptor expressed on B cells [51,69]. In 2020, Gasser et al. demonstrated the structural and mechanistic differences between ligelizumab and omalizumab, where they report epitope differences between the anti-IgE antibodies and indicate the differences contribute to their qualitatively distinct IgE-receptor inhibition profiles [69].

DARPins are small molecules and highly stable non-antibody protein scaffolds. Anti-IgE DARPins, which are genetically engineered proteins comprising a varying number of

stacked ankyrin-repeat domains, also block IgE binding to FccRI. Ankyrin-repeat domain binding can affect the stability and effector functions of target proteins. The motivation for engineering DARPins was to generate binding proteins that can be used to target proteins with high affinity and specificity, essentially replacing the use of monoclonal antibodies [51,70]. Therefore, the advantage of anti-IgE DARPins is that they not only neutralize free IgE but also actively disrupt pre-formed IgE–FccRI complexes through a facilitated dissociation mechanism [63,71]. The disruption of IgE–FccRI complexes by anti-IgE DARPins has been shown to be 10,000-fold more efficient than omalizumab in both in vitro and ex vivo human tissues [61]. Recently, newly engineered highly potent disruptive IgE inhibitors have been developed based on DARPin scaffolds, and their efficacies has been assessed [72].

Natural products, including herbal medicines and nutritional supplements, have been reported to inhibit MC degranulation. The underlying mechanisms appear to involve the blockade of IgE–Fc $\epsilon$ RI binding or IgE-mediated Fc $\epsilon$ RI signaling. For example, carotenoids suppress Ag-induced FccRI aggregation and inhibit FccRI-mediated intracellular signaling [73]. Procyanidin-enriched extracts from apples inhibit IgE binding to  $Fc \in RI$  and its subsequent signaling [74,75]. Medicinal herbs, such as Rubia cordifolia and Dianthus superbus in China; KOTMIN13 (composed of Inula japonica flowers), Trichosanthes kirilowii semen, *Peucedanum praeruptorum* radix, and *Allium macrostemon* bulbs in Korea [76]; and *Paeonia* radix and Zanthoxyli fructus in Japan, suppress MC degranulation and murine allergic inflammation [77,78]. These observations highlight the potential of plants and herbs as sources of bioactive compounds against allergic responses. However, their precise mechanisms are not fully understood, especially in the context of regulating  $Fc \in RI$  dynamics. We previously reported the potential of a natural product for regulating FccRI functions; an ephedra herb extract, Mao, in Japanese, significantly suppressed Ag-induced MC degranulation [79]. Mao significantly induced  $Fc \in RI$  internalization in MCs, thereby inhibiting Ag/IgE-dependent MC degranulation. We also showed that Mao induced  $Fc \in RI$  reorganization into distinct small clusters, without degranulation. These effects in Mao-treated MCs mimicked Ag-induced rapid desensitization states, suggesting that Mao has both therapeutic potential and unknown mechanisms, specifically in terms of  $Fc\epsilon RI$  cluster formation, which might be involved in MC desensitization [79].

#### 3.2. FceRI–Fc $\gamma$ RIIb Coaggregation and Inhibitory Signal against FceRI

Inhibitory signaling against FccRI is another candidate for FccRI-targeting therapy, which is mediated by co-crosslinking ITIM-containing inhibitory IgG receptor (FcyRIIb) with Fc $\epsilon$ RI. Fc $\gamma$ RIIb is highly expressed in murine MCs, murine basophils, and human basophils. Previous findings demonstrated that MCs isolated from human skin only express the activating FcyR and FcyRIIa proteins, whereas human cord blood-derived cultured MCs only express FcyRIIb [80,81], implying potential heterogeneity among human MCs with respect to the expression of  $Fc\gamma Rs$  [82].  $Fc\epsilon RI$  and  $Fc\gamma RIIb$  were co-crosslinked separately in Ag-independent and Ag-dependent manners. Ag-independent crosslinking of these receptors was achieved with bispecific antibodies against  $Fc \in RI$  and  $Fc \gamma RIIb$  or  $Fc \in -Fc \gamma$ fusion proteins to prevent allergic reactivity [83,84]. In contrast, specific IgG antibodies can induce allergen-specific IgE–Fc $\epsilon$ RI–Fc $\gamma$ RIIb crosslinking in the presence of specific allergens and competitively inhibit allergen binding to the specific IgE [85]. A recent clinical trial of cat allergen-specific IgG cocktails for respiratory allergies showed the efficacy of therapeutic applications of IgG, and demonstrated that passive immunization with allergenspecific IgG monoclonal antibodies potently suppresses allergic symptoms. Although receptor-mediated inhibition has not been clarified in the literature, these inhibitory effects might be due to IgG-FcyRIIb-dependent and/or IgG works as a neutralizing antibody for allergen [86].

# 3.3. FceRI Internalization in Allergen-Induced FceRI Desensitization

Allergen-induced  $Fc \in RI$  desensitization, which is achieved by sequentially increasing allergen stimulation from a suboptimal concentration and is accompanied by  $Fc \in RI$  internalization without MC (basophil) activation, is an important mechanism underlying allergenspecific immunotherapy [87]. This is the only disease-modifying therapy/intervention available for the treatment of allergies. Successful allergen-specific FccRI desensitization renders these effector cells less responsive or non-responsive, and the altered magnitude of mediators released from MCs represents the first step in a series of events following the induction of therapy. This could eventually lead to immunological tolerance to a specific allergen by modulating the subsequent development of Th2-biased responses. Such therapy provides a measurable benefit and is currently the only potential treatment for allergies. However, depending on disease phenotypes, the safety and efficacy are not sufficient in terms of the risk of undesirable adverse effects and the long duration of therapy [88]. Data from a recent clinical study showed a low compliance rate by patients over long periods of allergen-specific immunotherapy [89]. The authors of that study demonstrated that the compliance rates in the third year of treatment were 58.7% among patients diagnosed with allergic rhinitis (with or without allergic asthma) who were on subcutaneous immunotherapy, and 11.6% among those on sublingual immunotherapy [89]. These obvious limitations of allergen-specific immunotherapy have driven research toward alternative strategies. Several efforts aimed at improving therapeutic efficacies have been developed, including novel routes of immunotherapy, combining Ags with immunostimulatory adjuvants, and the use of modified Ags. For example, several combined therapies, such as omalizumab plus OIT, have been studied for treating food allergies. Traditional herbal medicines, which potently induce intestinal Treg cells, have been proposed as potential agents for improving the therapeutic efficacy when combined with specific Ags [90–93]. In experimental models of food allergy, combined therapies using herbal medicine suppressed MC activation at the inflammatory site and increased the therapeutic efficacy compared to OIT alone [92,93].

Modifying the properties of Ags has also been recognized as a potential strategy for improving immunotherapeutic efficacies [94]. A wide variety of structurally modified Ags have been generated, including chemically modified [95], adjuvant-bound [96], and recombinant hypoallergenic Ags (based on modifying the sequence of wild-type Ags) [97]. A recent study of the shrimp allergen tropomyosin showed that modifying tropomyosin by glycation decreases its allergenicity, where hypoallergenic glycated tropomyosin suppressed MC responses and allergic symptoms [98]. These modified Ags were designed to reduce adverse effects while maintaining immunogenicity at the systemic level. With regard to the local cellular events in MCs, these ideal Ags do not provoke robust MC activation, but induce  $Fc \in RI$  internalization. This silent  $Fc \in RI$  internalization could be attributed to a certain degree of FccRI dynamics in the context of the above-mentioned findings. Desensitized MCs, whose FceRI dynamics likely represent small and mobile  $Fc \in RI$  clusters and slow internalization (Figure 2), are discussed further below. Although limited data are available regarding the relationship between the desensitization efficacy and Fc $\epsilon$ RI dynamics, some novel mechanisms responsible for controlling Fc $\epsilon$ RI dynamics based on the above studies may provide important information for improving the efficacy of FccRI desensitization.

To explore the mechanisms underlying Fc $\epsilon$ RI desensitization in MCs and cell-specific local responses during allergen-specific immunotherapy, in vitro desensitization studies have been performed by several different groups [99–101]. IgE-sensitized MCs were subjected to sequential stimulation with incrementally increasing concentrations of Ag, which started from a subthreshold concentration and reached an optimal level. Desensitized MCs show decreases in IgE-bound Fc $\epsilon$ RI, which could be explained by internalization of the IgE–Fc $\epsilon$ RI complex [99,100]. Ang et al. investigated aberrant rearrangements of cytoskeletal actin fibers that inhibited Fc $\epsilon$ RI-mediated Ca<sup>2+</sup> flux and intracellular vesicle tracking [101]. Furthermore, recent data demonstrated activated tolerogenic capacities in the desensitized state [102] and that desensitized MCs showed tolerogenic features by releasing anti-inflammatory cytokines during the desensitization process, which were suggested to induce Treg cells in the local inflammatory site [102]. However, the precise mechanisms of successful desensitization, especially in the context of desensitized Fc $\epsilon$ RI dynamics, have not been thoroughly investigated. We recently performed in vitro MC desensitization using both DNP-HSA and ovalbumin (OVA) Ags and investigated the formation and localization of Fc $\epsilon$ RI clusters in MCs. We demonstrated that Fc $\epsilon$ RI clusters in vitro desensitized MCs were significantly smaller, despite sufficient Fc $\epsilon$ RI internalization as a consequence of desensitization [79,103]. The small Fc $\epsilon$ RI clusters in desensitized MCs may be mobile and slowly internalized. Additionally, when we determined the Ag properties (i.e., antigenicity) using heated OVA and naïve OVA, heated OVA showed higher allergenicity, possibly because the thermal treatment caused OVA aggregation. Heated OVA failed to induce desensitization at the same concentration as naïve OVA. Heated OVA showed undesirable degranulation during desensitization, where significantly larger clusters were formed [103]. Collectively, these results suggest that manipulation of the allergenicity is critical for successful desensitization.



**Figure 2.** FccRI desensitization and hypothesized mechanism of silent FccRI internalization. Ag–IgE–FccRI complexes are capable of internalization without MC activation. (**Left**) High-intensity stimulation (e.g., modest to high concentrations of high-affinity Ag) causes a rapid transition of FccRI from a diffuse state to a highly immobile state, resulting in the formation of large aggregates. The rapidly internalized FccRI clusters are observed obviously large, in which rapid short-lived positive signals are derived. (**Right**) During FccRI desensitization with incremental Ag stimulation, suboptimal stimulations might allow FccRI to remain mobile and diffuse, and form smaller clusters, followed by slower internalization. There is no considerable degranulation, but a sufficient degree of internalization and signals may result.

Taken together, these data reviewed above indicate that specific features of  $Fc\epsilon RI$  dynamics appear to be involved in  $Fc\epsilon RI$  desensitization in MCs. With therapeutic interventions based on  $Fc\epsilon RI$  desensitization, ideal Ags may be those that can flexibly and generally induce small  $Fc\epsilon RI$  clusters and regulate moderate  $Fc\epsilon RI$  dynamics, which may contribute to the appropriate regulation of  $Fc\epsilon RI$  clusters that are small and may be mobile, diffuse, and compete for internalization (Figure 2). However, one important limitation/question is whether small cluster induction and internalization, which certainly prevent rapid and robust degranulation, might induce other types of signaling or sustained/prolonged immunological responses. Indeed, the MC response shows diverse reactivity upon weak stimulation [104]. Investigating such mechanisms may lead to discoveries that influence effective desensitization and produce more reliable information. Further development

of this topic might help in understanding the regulatory mechanisms involved in related therapies, which may offer insights into optimizing Ag immunotherapy and rapid desensitization strategies.

#### 3.4. Perspective

Adequate regulation of Fc $\epsilon$ RI dynamics is a potential intervention strategy for controlling MC qualitative functions and allergic inflammation. Fc $\epsilon$ RI may actively adjust cellular responses even when no considerable degranulation occurs immediately after Ag challenge, which is likely to contribute to improving Fc $\epsilon$ RI targeting therapy.

# 4. Conclusions

Fc $\epsilon$ RI displays dynamic changes in response to the properties of Ags and IgEs, which induce differential signaling pathways that regulate MC functions. A complementary mechanism may exist, even after weak or simple input patterns. Fc $\epsilon$ RI on the MC responds to the stimulation intensity/quality and translates it into an appropriate MC response. Taking advantage of this Fc $\epsilon$ RI characteristic may improve Fc $\epsilon$ RI-targeting therapeutics, especially Fc $\epsilon$ RI desensitization and Ag-specific immunotherapy for allergies.

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