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The human immunoglobulin A Fc receptor $Fc\alpha RI$: a multifaceted regulator of mucosal immunity

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Immunoglobulin A (IgA) is commonly recognized as the most prevalent antibody (Ab) at mucosal sites with an important role in defense by shielding mucosal surfaces from invasion by pathogens. However, its potential to both actively dampen excessive immune responses or to initiate potent proinflammatory cellular processes is less well known. Interestingly, either functional outcome is mediated through interaction with the myeloid IgA Fc receptor $Fc\alpha RI$ (CD89). Monomeric interaction of IgA with $Fc\alpha RI$ triggers inhibitory signals that block activation via other receptors, whereas multimeric $Fc\alpha RI$ crosslinking induces phagocytosis, reactive oxygen species production, antigen presentation, Ab-dependent cellular cytotoxicity, and cytokine release. Thus, $Fc\alpha RI$ acts as a regulator between anti- and proinflammatory responses of IgA. As such, the biology of $Fc\alpha RI$, and its multifaceted role in immunity will be the focus of this review.

INTRODUCTION

Immunoglobulin A (IgA) is the predominant antibody (Ab) class present in mucosal areas, where it has a key role in mucosal defense.¹ Mucosal surfaces represent a vast interface of over 400 m² that protects internal tissues from external influences. They are continuously exposed to inhaled or ingested antigens and pathogens, and are colonized by a commensal microbiota, which must be prevented from penetrating the underlying tissues. As such, mucosal homeostasis requires a delicate balance between avoiding disproportionate responses against innocuous antigens, whereas at the same time effective immunological responses against pathogenic microorganisms must be maintained.² Interestingly, depending on the circumstances, IgA is involved in anti- as well as proinflammatory responses, although the view of IgA as an anti- or non-inflammatory Ab commonly prevails.^{3,4}

At mucosal sites, IgA is produced in the LP by local plasma cells as dimeric molecules (dimeric IgA; dIgA), containing a joining J chain. It serves as an intermediary, and can bind to the pIgR that is expressed on the basolateral membrane of epithelial cells. Subsequently, it is transported through the epithelial cells and released into the lumen as SIgA.^{5–7} Apical cleavage of pIgR ensures continual attachment of a part of this receptor—referred to as SC—which renders increased stability and prevents rapid breakdown of SIgA in the hostile gut lumen milieu. In addition, 1–3 mg ml⁻¹ IgA is present in the circulation as a monomer, and as such it is the second prevalent Ab in serum.

Multiple types of cellular IgA receptors have currently been characterized. In addition to pIgR,⁸ $Fc\alpha/\mu$ receptors,⁹ asialoglycoprotein receptors,¹⁰ transferrin receptors (CD71),¹¹ SC receptors,¹² and M-cell receptors¹³ have been described that can bind the IgA Fc tail, carbohydrate side chains or accessory molecules such as the J chain and SC. The functions of a number of these receptors have not yet been completely elucidated.

In addition, in humans, a myeloid Fc receptor for IgA ($Fc\alpha RI$; CD89) has been described. Thus far, $Fc\alpha RI$ gene homologs have been identified in primates, horses, cattle, hamsters, gerbils, and rats, but not in mice, which is attributable to a gene translocation.^{14–19} The absence of $Fc\alpha RI$ in mice, which are frequently used for experimental work, has likely influenced the earlier dogma of IgA as an anti-inflammatory Ab class. However, a main role for $Fc\alpha RI$ in immune defense is supported, as bacterial evolution has led to the development of proteins that interfere with IgA binding to $Fc\alpha RI$ (e.g., IgA-binding M-like proteins Arp4, Sir22, B-antigen, and members of the staphylococcal superantigen-like proteins family),²⁰ resulting in an important evasion strategy for pathogens to escape IgA-mediated phagocytosis. In this review we will therefore address the biology, function and therapeutic potential of $Fc\alpha RI$.

$Fc\alpha RI$: GENETICS, TRANSCRIPTS, AND PROTEIN STRUCTURE

$Fc\alpha RI$ is a member of the Fc receptor Ig superfamily, although distinct differences can be observed compared with other Fc

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receptors. For instance, the *Fc α RI* gene is located on chromosome 19 (at 19q13.4) and lies within the so-called leukocyte receptor cluster (LRC),^{21,22} whereas other FcR genes, like *Fc γ R*s and *Fc ϵ RI*, map on chromosome 1.^{23,24} The human LRC includes no other Fc receptor genes, but instead it encodes killer cell immunoglobulin-like receptors and leukocyte Ig-like receptors. Fc α RI shows more amino acid (aa) sequence similarities with these receptors than with other human Fc receptors.²⁵ In addition, the murine paired Ig-like receptor-A that was identified in mice on the basis of the homology with Fc α RI, shares sequence similarity with both human Fc α RI and killer cell immunoglobulin-like receptors.²⁶

Fc α RI consists of two extracellular (EC) Ig-like domains of each 206 aa, a 19-aa transmembrane region (TM), which is crucial for association to the signaling FcR γ -chain, and a short (41 aa) cytoplasmic tail. The two EC domains are folded with an angle of approximately 90° to each other. The Fc α RI gene (*FCAR*) coding for this protein consists of five exons. The first two exons encode the leader peptide (S1, 34 bp, and S2, 36 bp), whereas exons 3 (291 bp) and 4 (288 bp) code for EC1 and EC2, respectively. The transmembrane and intracellular domains are both encoded by exon 5 (215 bp).^{27–29} Furthermore, a 78-bp insertion sequence (S3) between S2 and EC1 has been reported.³⁰ The Fc α RI promoter was identified in a 929-bp fragment of Fc α RI 5'-flanking sequence, in which tissue-specific gene expression is regulated by 259 bp, proximal to the translation initiation site.³¹ The sequence between 59 and 197 bp, downstream of the major transcription start site, is essential for promoter activity and contains multiple potential binding sites for transcription factors ((C/EBP) binding sites, nuclear factor- κ B binding site, Spl site, Ets family protein consensus binding site, and a Myb binding site), which have been reported to function in myeloid-specific gene expression.^{32,33} The core protein has a predicted molecular weight of 30 kDa. However, Fc α RI that is expressed on the cell surface has an apparent molecular weight between 50 and 75 kDa, with the exception of eosinophil Fc α RI, which runs between 70 and 100 kDa. This heterogeneous pattern of surface-expressed Fc α RI is a consequence of both N- and O-glycosylation.^{34–36}

Several isoforms have been described in primary cells in addition to the full-length Fc α RI protein. Fc α RIa.2 bears deletion of 66 aa in EC2, and it is the only isoform expressed on alveolar macrophages, which might have a physiological relevance in IgA-mediated host defense in the lung.³⁷ In addition, the isoform Fc α Rb lacks the transmembrane and intracellular domain, which is replaced by 23 new aa. This is the consequence of alternative splicing that skips the 3-splice site at the end of the EC2 exon, resulting in the insertion of 23 new aa before the stop codon. Fc α Rb has been described in both neutrophils and eosinophils.³⁸ Although Fc α Rb can be expressed on the cell membrane—which is probably due to the 23 aa insertion—it is presented mainly as a soluble form. Furthermore, cell-surface-expressed Fc α Rb is unable to associate with FcR γ -chain and therefore unable to exert cellular signaling after binding to IgA complexes. This suggests that full-length Fc α RI and soluble Fc α Rb on granulocytes may compete for IgA binding.

As such, Fc α Rb may downregulate signaling via full-length Fc α RI. A Fc α Rb alternative spliced transcript (Fc α Rb Δ S2) has been described as well, in which the leader peptide 2 is deleted. However, it is not clear whether this transcript is translated into a functional protein. A second soluble Fc α RI has been described that is specifically expressed by monocytes, and which is the result of crosslinking of full-length Fc α RI, which triggers FcR γ -chain-dependent shedding.³⁹ Interestingly, release of soluble Fc α RI—which is likely due to proteolytic cleavage—is induced by IgA aggregates, suggesting a regulatory effect on Fc α RI effector functions. A 2–3 kDa smaller protein variant of full-length Fc α RI (with a core protein of 29–30 kDa) was described in neutrophils, as well.⁴⁰ The nature of this variant is not clear at present.

In addition, mRNA encoding Fc α RIa.3 that lacks EC2 has been identified in granulocytes and monocytes.⁴¹ Cells, which were transfected with this variant, were able to bind SIgA, but not serum IgA. Furthermore, tumor necrosis factor- α specifically increased or decreased Fc α RIa.3 transcripts compared with full-length Fc α RIa.1 transcripts in neutrophils or monocytes, respectively.⁴² Other alternative spliced transcripts have been described on mRNA, but not at protein level, in primary cells or cell lines.^{34,37,41–44} These include the Δ S2 transcript, in which the leader peptide 2 is deleted, Δ S2EC1, in which both leader peptide 2 and EC1 are eliminated, and transcripts with (part of) EC2 deletions, including Δ S2EC2 (deletion of leader peptide 2 and EC2), Δ S266EC2 (deletion of part leader peptide 2 and 66 bp of EC2), and Δ S274EC2 (missing leader peptide 2 and 74 bp of EC2). Biological significance is not yet completely understood, although different levels of full-length and alternative spliced Fc α RI transcripts have been documented in several diseases.^{30,42} For example, neutrophils from patients suffering from pneumonia have lower Fc α RIa.3 transcripts, whereas monocytes from patients with IgA nephropathy solely express full-length Fc α RI. As such, protein isoforms may diversify Fc α RI structure and function in immunoregulation of IgA-mediated host defense.

Several single-nucleotide polymorphisms (SNPs) have been described in the *FCAR* gene as well.^{29,31,45–51} These constitute three different SNPs within the *FCAR* gene promoter region, namely –340G/A, –311T/C, and –142T/C, of which the latter two show lower promoter activity within the T allele. Furthermore, two non-coding SNPs (324A/G and 363A/G) are documented within the EC1. The SNP 376G/A results in an aa change from aspartic acid into asparagine. The effect of this variation on IgA binding is not known, although it is located closely to the ligand binding site.⁵² Finally, a functional SNP has been identified in the intracellular domain (844A/G), which results in an aa transition from serine to glycine (S248–G248).⁵⁰ IgA-mediated crosslinking of neutrophil Fc α RI–G248 triggered significantly more interleukin (IL)-6 release than equivalent crosslinking of the Fc α RI–S248 variant. Remarkably, only Fc α RI–G248, is capable of inducing cytokine release in the absence of FcR γ -chain, which is presumed due (at least in part) to its ability to interact directly with the Src family member Lyn, an important component of the Fc α RI signaling cascade.⁵⁰

Association of *FCAR* gene polymorphisms with diseases has been investigated to some extent. Until now, none of the SNPs were associated with allergy,⁵² but altered susceptibility to aggressive periodontitis (324A/G SNP),⁴⁸ systemic lupus erythematosus (844A/G SNP),⁵⁰ and chronic HCV infection (–311T/C and –142T/C SNPs) have been described.⁴⁹ Furthermore, the (844A/G SNP) influences susceptibility to systemic lupus erythematosus, but not systemic sclerosis or rheumatoid arthritis,^{50,53} whereas controversy exists for *FcαRI* polymorphisms in relation to susceptibility to IgA nephropathy.^{31,46,54}

EXPRESSION AND MODULATION

FcαRI expression is already observed at the promyelocyte stage in differentiation, and is restricted to cells of the myeloid lineage, including neutrophils, eosinophils, monocytes, and several macrophage subsets (e.g., alveolar, tonsillar, and splenic, but not small intestine macrophages). *FcαRI* is furthermore expressed on Kupffer cells and on interstitial, CD34⁺-derived dendritic cells (DCs) and monocyte-derived DCs (although the latter may be reflected by *FcαRI* expression on monocytes, which decreases during DC differentiation).^{34–37,55–59} Expression has recently been described on human platelets as well, whereas *FcαRI* is not expressed on mast cells or basophils.⁶⁰ *FcαRI* expression is constitutive and independent of its ligand, which is demonstrated in IgA-deficient patients who still express *FcαRI*.⁶¹ However, depending on the cell type, expression and function of *FcαRI* can be modulated by lipopolysaccharide, chemoattractants, inflammatory cytokines, or adapter protein, binding to the intracellular domain of *FcαRI*.^{25,62} For instance, upregulation is induced by IL-1β, tumor necrosis factor-α, granulocyte-macrophage-colony stimulating factor, and IL-8,^{63–66} whereas expression is downregulated by transforming growth factor-β, interferon-γ, or the ligand polymeric IgA.^{67,68} Upregulation of neutrophil *FcαRI* expression levels can be the result from either *de novo* synthesis or transport from intracellular stores to the cell surface.⁴⁰ Full-length *FcαRI* is present in both secretory and tertiary granules, whereas the 2–3 kDa smaller variant (core protein of 29–30 kDa) is also present in tertiary granules. As these granules are differently mobilized during neutrophil activation or inflammatory responses, distinct biological functions for *FcαRI* are suggested, but this has not yet been investigated thoroughly. Tissue distribution of *FcαRI* is mostly defined by the presence of neutrophils and few emigrated macrophages.⁶⁹ These cells are evident as clusters in tonsils and appendix, and are scattered in varying numbers in lymph nodes, kidneys, livers, intestinal mucosa, bronchoalveolar lavages, or peritoneal fluid. Inflamed intestines display major influxes of *FcαRI*-positive neutrophils. The level of *FcαRI* on neutrophils was similar in tissue compared with blood neutrophils, whereas *FcαRI* expression on monocytes was much lower in tissues than in blood. Altered *FcαRI* expression has been reported in diseases such as ankylosing spondylitis, allergic diseases, human immunodeficiency virus (HIV) infection, or bacterial infections.^{36,70–72}

LIGAND BINDING

Although all forms of IgA are ligands for *FcαRI*, the binding capacities differ. Although monomeric IgA and dimeric IgA are capable of binding *FcαRI* with moderate affinity ($K_a = \sim 10^6 \text{ M}^{-1}$) in the boundaries of Cα2 and Cα3, IgA-immune complexes bind avidly.^{73–77} Furthermore, because of the partial overlap of the IgA binding site for either *FcαRI* or pIgR, interaction of SIgA to *FcαRI* is (partly) hampered because of steric hindrance of SC.^{75,78–80} SIgA binding is however increased when complement receptor 3 functions as coreceptor.

Crystallographic studies demonstrated that one IgA molecule can simultaneously bind two *FcαRI* molecules (**Figure 1**).^{75,76} This is in contrast to *FcγRIII* and *FcεRI*, for which a 1:1 stoichiometry with their respective ligands was described, again emphasizing dissimilarities between *FcαRI* and other members of the Fc receptor family.^{81–85} Moreover, the IgA binding site on *FcαRI* is located in EC1 (**Figure 1**, depicted as yellow aa),^{75,76,79} which is different compared with *FcεRI* and *FcγRs*, as these FcRs bind their ligands in EC2. Residues Y35 (in the BC loop), R52, R53, L54, K55 (in the D strand), F56, W57, N58 (in the DE loop), Y81, R82, I83, G84, H85, and Y86 (in the FG loop) within EC1 are involved in IgA binding.^{73–75} Surprisingly, alternative spliced *FcαRI*, which lacks EC2, does not bind serum IgA, but binds SIgA comparable to full-length *FcαRI*.⁴¹

Residues within IgA that are involved in *FcαRI* binding (**Figure 1**, depicted as blue aa) are L256, L257, and L258 in the α-helix of the AB loop of Cα2. Within Cα3 E348 (A-strand), R382, L384 (C-strand), S387, E389 (CC' loop), M433, H436 (F-strand), E437, A438, L439, P440, L441, A442 (FG loop), F443, T444, and Q445 (G strand) are involved.^{75,86} Immune complexes with optimal binding contain five to six molecules of IgA per complex.⁸⁷ A number of conformational changes have been observed within the *FcαRI*–EC1 domain (in the D-strand, DE and FG loop) after binding to IgA, but the approximately 90° orientation of the two EC domains do not change significantly.⁷⁵ Furthermore, both *FcαRI* and IgA are heavily glycosylated proteins. *FcαRI* harbors six N-glycosylation sites and several putative O-linked glycosylation sites, and it has been demonstrated that deglycosylation of *FcαRI* N58 increases IgA binding.⁸⁸ By contrast, IgA Fc glycosylation is not critical for binding to *FcαRI*.^{88–90} Several monoclonal Abs (mAbs) have been described that bind *FcαRI* (**Figure 1**, right part) in EC1 (My43, 2E6, 2D11, 7G4, 2H8, and MIP8a), EC2 (A59, A77, A62, and 7D7), or have binding sites in both ECs (A3).^{34,91,92} All EC1 recognizing mAbs block IgA binding, whereas mAb A62 recognizes low glycosylated *FcαRI*. Recently, it was described that the pentraxin C-reactive protein binds to *FcαRI*, which induces cellular activation. C-reactive protein binds to distinct regions of *FcαRI* compared with IgA, suggesting simultaneous binding of both molecules to *FcαRI*.⁹³

Ligand binding to *FcαRI* is regulated via a mechanism referred to as inside-out signaling, which entails that cytokine stimulation of cells rapidly, modulates binding capacity in response to intracellular signals, without affecting receptor expression levels.^{94–98} Experiments with eosinophils, monocytes, and

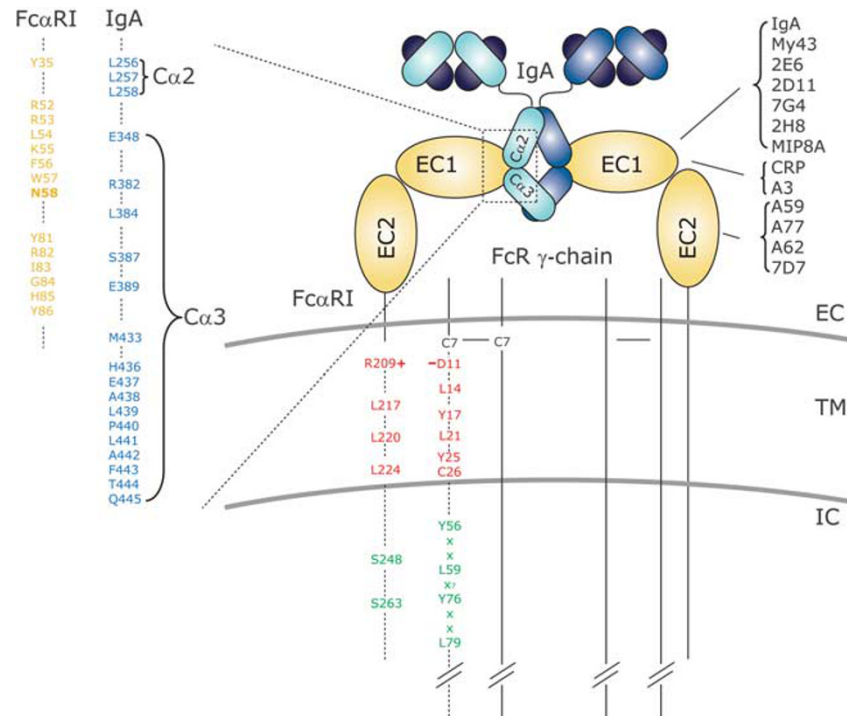


Figure 1 Schematic representation of the FcαRI–FcR γ-chain complex, binding immunoglobulin A (IgA) in a 2:1 stoichiometry. Two FcαRI bind each IgA–Fc part at the Cα2 and Cα3 junction via extracellular (EC) 1. Amino acids (aa) involved in ligand–receptor binding are depicted in yellow for FcαRI–EC1 and in blue for aa in Cα2 and Cα3 of IgA (left part of figure). Deglycosylation of N58 in FcαRI–EC1 (bold) increases IgA binding. Amino acids in the transmembrane regions (TMs) of FcαRI and a FcR γ-chain homodimer, involved in complex formation, are depicted in red, whereas aa involved in signaling are shown in green. FcαRI intracellular (IC) serine 248 and 263, respectively, modulates FcR γ-chain independent interleukin (IL)-6 production and inside–out signaling. FcR γ-chain intracellular immunoreceptor tyrosine-based activation motif (ITAM) consensus (YxxLx₇YxxL) is shown, as well as the disulfide bond between two cysteines at position 7 (solid line). Monoclonal antibodies and C-reactive protein recognizing different ECs of FcαRI are depicted (right part).

transfectants demonstrated that FcαRI shows low capacity to interact with IgA–immune complexes in a resting state, but ligand binding capacity increases profoundly after stimulation with cytokines such as granulocyte-macrophage-colony stimulating factor, and IL-4 or IL-5. As such, FcαRI becomes primed, but surface receptor expression is not augmented.^{96–98} Inside–out signaling critically depends on the intracellular domain of FcαRI and on the presence of an intact cytoskeleton, but it does not require FcR γ-chain.^{96,97} In more detail, phosphorylation of serine 263 (**Figure 1**, depicted in green), and activation of phosphatidylinositol 3-kinase (PI3K) and its downstream target protein kinase C are essential in switching inactive FcαRI into an active, ligand binding receptor (**Figure 2a**). Simultaneously, cytokine stimulation induces binding of the serine–threonine phosphatase protein phosphatase 2A to the intracellular domain of FcαRI, which results in dephosphorylation of the FcαRI intracellular domain, allowing binding of IgA–immune complexes.⁹⁸ Whether FcαRI priming is a result of enhanced lateral movement (avidity) and/or conformational changes within the receptor (affinity), as described for integrins,⁹⁹ is currently unknown. It is not clear whether FcαRI on neutrophils needs priming as well, as activation of neutrophils during isolation has, as of yet, precluded these experiments. However, neutrophils of patients suffering from active, ongoing derma-

titis herpeticiformis (DH) demonstrate increased ability to bind IgA, without increased receptor expression, which is consistent with a pattern of receptor priming.¹⁰⁰ This suggests that priming is the result of ongoing mucosal inflammatory responses and concomitantly systemic cytokine release in patients with dermatitis herpeticiformis.

IgA-MEDIATED FcαRI SIGNALING AND CELLULAR FUNCTIONING

Binding of IgA–immune complexes (containing either monomeric IgA or dimeric IgA) induces proinflammatory responses, which requires association of FcαRI with the FcR γ-chain subunit.^{101–103} Two basic aspects are essential for tethering FcαRI and FcR γ-chain into a stable FcαRI–FcR γ-chain complex. First, dimerization of two γ-chains—through disulfide bond between two cysteines—is required, and second, association between the positively charged arginine 209 (R209) in the TM of FcαRI and an opposite negatively charged aspartic acid 11 (D11) in the TM of FcR γ-chain is essential (**Figure 1**, depicted in red). Moreover, it was demonstrated that orientation of the positive charge within the TM of FcαRI into the vicinity of the FcR γ-chain dimer is important.¹⁰⁴ In addition, two leucines (L14 and L21), two tyrosine (Y17 and Y25), and a cysteine at position 26 (C₂₆), within FcR γ-chain, as well as three leucines on position 217,

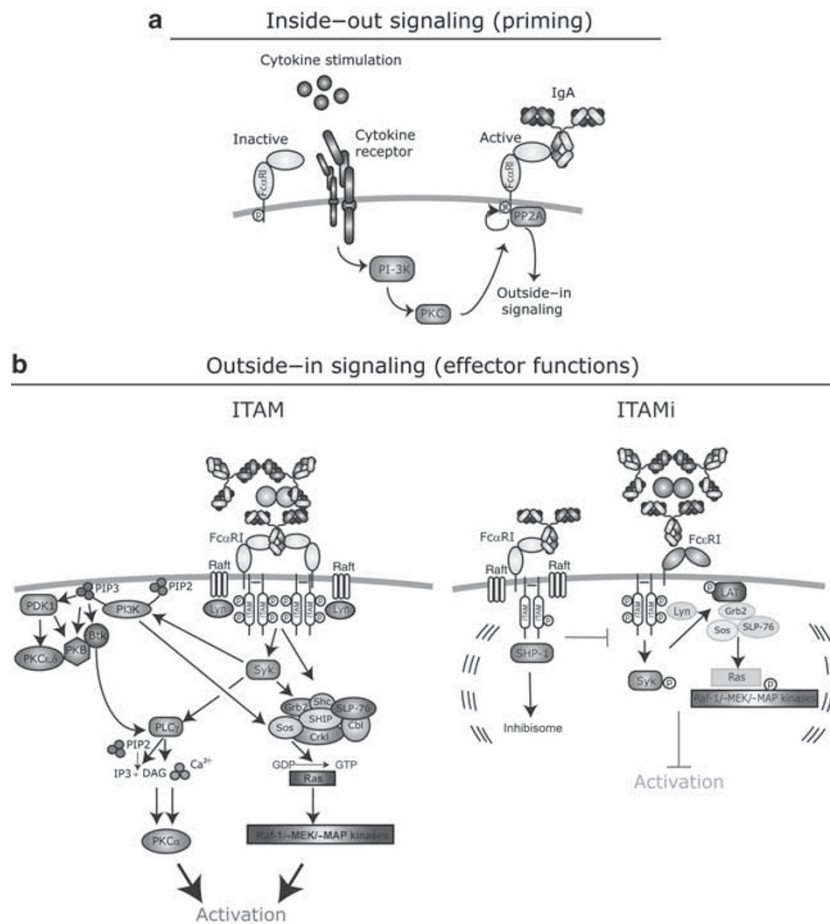


Figure 2 Simplified scheme of signaling pathways involved in FcαRI functioning. **(a)** Inside-out signaling or priming of FcαRI. The intracellular domain of FcαRI, an intact cytoskeleton, phosphorylation of serine 263, phosphatidylinositol 3 kinase (PI3K), and its downstream target protein kinase C (PKC) and serine/threonine phosphatase protein protein phosphatase 2A (PP2A) are involved in switching inactive FcαRI into an active, ligand binding receptor after cytokine stimulation. **(b, Left)** Crosslinking of FcαRI by immunoglobulin A (IgA)-immune complexes induces redistribution of FcαRI to plasma membrane rafts. Src kinase Lyn phosphorylates the tyrosines within the associated FcR γ-chain-immunoreceptor tyrosine-based activation motif (ITAM). These then serve as “docking” sites for recruitment of B lymphocyte kinase (Blk), Syk, phospholipase (PLC)-γ, Shc, and growth factor receptor-bound protein 2 (Grb2), which facilitates activation of multiple (and subsequent) targets such as PI3K, PLC-γ, and components of a Grb2 containing multimolecular adapter protein complex. This results in cellular functions such as phagocytosis, Ab-dependent-cellular cytotoxicity, respiratory burst, degranulation, antigen-presentation, and release of cytokines and inflammatory mediators. **(Right)** Triggering FcαRI with monomeric serum IgA (not crosslinking FcαRI) transduces inhibitory signals through FcαRI-FcR γ-chain complex via inhibitory capacity through FcR γ-chain ITAMi (ITAMi), which downregulates other activating Fc receptors. The inhibitory signal involves recruitment of Src homology region 2 domain-containing phosphatase-1 (SHP-1) to FcαRI, and formation of inhibisome clusters (dotted lines), which impair phosphorylation of Syk, LAT, and ERK. Ca²⁺, calcium; Cbl, Casitas B-lineage lymphoma; DAG, diacylglycerol; GDP, guanosine diphosphate; GTP, guanosine triphosphate; LAT, linker of activated T cells; p, phosphate; PIP2, phosphatidylinositol 4,5-bisphosphate; PIP3, phosphatidylinositol (3,4,5)-triphosphate; SLP-76, SH2 domain containing leukocyte protein of 76 kDa; SHIP, Src homology-2-containing inositol 5'-phosphatase.

220, and 224, within FcαRI TM, contribute to the stabilization of the FcαRI-FcR γ-chain complex^{105,106} (Figure 1, depicted in red). *In vivo* studies in FcαRI transgenic-FcR γ-chain knock-out mice demonstrate that FcR γ-chain association is essential for FcαRI surface expression.¹⁰³ However, FcαRI expression in the absence of FcR γ-chains (FcαRI “γ-less”) was observed in transfected cell lines, and selective human monocyte and neutrophil populations.^{107–109} Functionality of FcαRI “γ-less” receptors is limited to ligand binding (inside-out signaling) and receptor recycling via early endosomes, although FcαRI-Gly248 was shown to trigger IL-6 production in the absence of FcR γ-chain.⁵⁰

The mechanism of FcαRI binding to FcR γ-chain underscores the difference with other FcRs, but shows similarities with members of the LRC family.^{22,74} Within the LRC family, an N-terminal positively charged arginine is highly conserved among all activatory receptors, with the exception of killer cell activatory receptors, which bear a positively charged lysine in the center of their TM. Interestingly, killer cell activatory receptors interact with the signaling molecule DAP12, whereas other LRC stimulatory receptors associate (like FcαRI) with the FcR γ-chain.^{110,111} In addition, paired Ig-like receptor-A associates with the FcR γ-chain via an transmembrane N-terminal arginine. By contrast, FcR γ-chain association of other activating Fc receptors is based

on the residues in the C-terminal part of their TMs.^{112–114} For example, Fc γ RI requires a domain of 10 aa, including an asparagine and Fc ϵ RI, and Fc γ RIIIa require a C-terminal aspartic acid for FcR γ -chain association.

Crosslinking of Fc α RI by IgA-immune complexes induce FcR γ -chain independent redistribution of Fc α RI to plasma membrane rafts.¹¹⁵ Furthermore, both the tyrosine kinase Bruton's tyrosine kinase and *src* family kinase Lyn are recruited to these signaling platforms. Fc α RI crosslinking furthermore initiates immunoreceptor tyrosine-based activation motif (ITAM)-dependent signaling of the Fc α RI-associated FcR γ -chain (**Figure 2b**, left part).^{23,116} The FcR γ -chain ITAMs consists of a conserved stretch of aa of paired tyrosines and leucines in a consensus sequence (YxxLx₇YxxL, **Figure 1**, depicted in green). *Src* kinase Lyn phosphorylates the tyrosines within the associated FcR γ -chain-ITAM. These then serve as the “docking” sites for recruitment of other tyrosine kinases, including B lymphocyte kinase and the Src homology 2 (SH2)-domain-containing proteins Syk, phospholipase C- γ , Shc, and growth factor receptor-bound protein 2, which facilitates the activation of multiple targets such as PI3K (with downstream PDK1, protein kinase C ϵ and δ , PKB α , and Bruton's tyrosine kinase activation) and phospholipase C- γ (with downstream release of IP3 and diacylglycerol to trigger calcium release and activation of calcium- and diacylglycerol-dependent protein kinase C α) (**Figure 2b**, left part).^{115,117–120} Of note, these pathways are interconnected, for example, Bruton's tyrosine kinase can potentiate calcium signaling—via phospholipase C γ —but is activated by binding to the PI3K product phosphatidylinositol (3,4,5)-triphosphate via its PH domain. Moreover, *Src* family kinases, that are activated after Fc α RI stimulation, also induces the formation of multimolecular adapter protein complexes consisting of the adapter growth factor receptor-bound protein 2—which is constitutively bound to Sos—and recruited upon phosphorylation of Shc. Furthermore, this complex contains Src homology-2-containing inositol 5'-phosphatase, Casitas B-lineage lymphoma, SH2 domain containing leukocyte protein of 76 kDa, and Crkl.¹²¹ Through this adapter complex, GDP-Ras is exchanged to active GTP-Ras by Sos (a guanine nucleotide exchange factor), that is activated by PI3K, which, in turn, activates Raf-1-MEK-MAP serine-threonine kinases by sequential phosphorylation. The interconnected signaling pathways couple upstream FcR γ -chain ITAM phosphorylation to different cellular processes, such as gene expression by activation of several transcription factors (including nuclear factor- κ B, AP-1, and Sp1), phagocytosis, Ab-dependent-cellular cytotoxicity, respiratory burst, degranulation, antigen presentation, and release of cytokines and inflammatory lipid mediators.^{103,109,119–130} Of note, depending on cell type or cell stimulation, Fc α RI activation may trigger cell type-specific signaling and functional responses.

Intriguingly, it was demonstrated that non-targeted monomeric serum IgA (not crosslinking Fc α RI) transduces inhibitory signals through the Fc α RI-FcR γ -chain complex, which downregulates IgE- or IgG-, Fc-receptor-mediated phagocytosis, chemotaxis, bacterial activity, oxidative burst activity, and

cytokine release.^{131–139} The underlying molecular mechanisms involves ERK-dependent recruitment of tyrosine phosphatase Src homology region 2 domain-containing phosphatase-1 and Fc α RI to lipid rafts (**Figure 2b**, right part).^{131,140} In a second step, ligation of the activating Fc receptor results in colocalization with Fc α RI and Src homology region 2 domain-containing phosphatase-1 in rafts. These assembled proteins form intracellular structures called “inhibisomes,” which induce impairment of Syk, LAT, and ERK phosphorylation and functionality of activating Fc receptors. Formation of inhibisomes requires Src homology region 2 domain-containing phosphatase-1-dependent depolarization of actin. Sustained aggregation of Fc α RI by multimeric ligands by contrast stimulates cell activation by recruiting high amounts of Syk (described above), whereas Src homology region 2 domain-containing phosphatase-1 binding is aborted.¹³¹ The inhibitory capacity through FcR γ -chain ITAM is referred to as ITAMi, which differs from the classical immunoreceptor tyrosine-based inhibitory motif signaling of inhibitory Fc receptors, which requires co-aggregation with the activatory Fc receptor.^{23,141} Thus, both IgA-induced activating and inhibitory signals depend on Fc α RI-FcR γ -chain ITAM, but differ in the recruitment of tyrosine kinases versus tyrosine phosphatases, respectively, (**Figure 2b**). As such, it has been proposed that crosslinking of Fc α RI during infection with IgA-opsonized pathogens results in proinflammatory responses, whereas naturally occurring serum IgA (not complexed with an antigen) induces inhibitory signals through Fc α RI to dampen excessive immune responses (initiated by other Ig-immune complexes).

Fc α RI AND IgA IN MUCOSAL IMMUNOLOGY

Only few Fc α RI-positive cells are observed in mucosal areas in homeostatic conditions. For instance, intestinal macrophages lack Fc α RI expression,⁵⁸ which is consistent with an anti-inflammatory role of IgA to protect mucosal integrity. Furthermore, mucosal Langerhans cells do not express Fc α RI. However, low Fc α RI levels were observed on *in vitro* cultured monocyte-derived DCs, which resemble interstitial DC human epithelial interstitial-type DC. A study by Heystek *et al.*⁵⁹ demonstrated low level expression of Fc α RI on immature monocyte-derived DCs as well. Crosslinking of Fc α RI led to internalization of (serum) IgA complexes and antigen presentation through the major histocompatibility complex class II pathway, but not via major histocompatibility compatibility class I cross-presentation, which resulted in monocyte-derived DCs maturation and production of IL-10.^{57,120,142} Because IL-10 mediates IgA isotype switching,¹⁴³ a role for Fc α RI-positive DC in mounting specific immune responses is supported. However, it should be noted that, although Fc α RI is able to mediate efficient antigen presentation in Fc α RI-expressing transfectants,¹⁰⁹ uptake of IgA immune complexes by DCs, expressing low levels of Fc α RI, led to poor antigen presentation.¹⁴⁴ Efficient uptake of SIgA was demonstrated, which was partially blocked by anti-mannose receptor mAb, but not by anti-Fc α RI blocking mAbs. This indicated that monocyte-derived DCs did not internalize SIgA via Fc α RI, but through

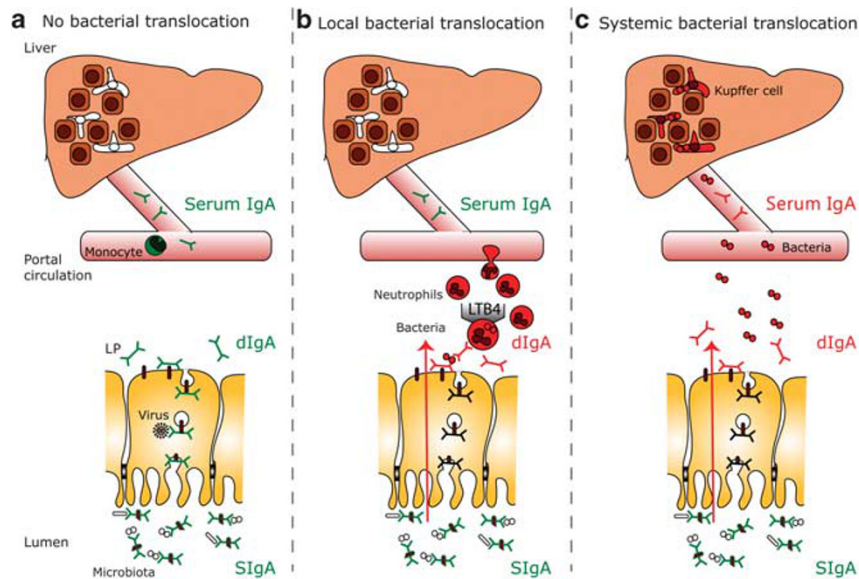


Figure 3 Model for the role of $Fc\alpha RI$ in mucosal immunity. In humans, immunoglobulin A (IgA) is expressed in three different forms. **(a)** In homeostatic conditions dimeric IgA (dIgA) functions as intermediary molecule that is secreted as SIgA, which inhibits bacterial invasion. Monomeric serum IgA functions as anti-inflammatory molecule through targeting $Fc\alpha RI$ —inhibitory capacity through FcR γ -chain ITAM (ITAMi) signaling. **(b)** When microorganisms have been able to breach the epithelial barrier, dIgA can opsonize these pathogens. Recruited neutrophils that express $Fc\alpha RI$ will clear the infection through phagocytosis and release of leukotrien B4 (LTB4, via active immunoreceptor tyrosine-based activation motif (ITAM) signaling), which may lead to a self-controlled positive feedback loop, until pathogens have been eliminated. Hence, in case of local bacterial translocation, dIgA represents a proinflammatory molecule and functions as the second line of innate mucosal immune defense. **(c)** Finally, pathogens that have entered the portal circulation are opsonized by serum IgA, and subsequently phagocytosed by $Fc\alpha RI$ -positive Kupffer cells. As such, serum IgA can function as proinflammatory antibody when systemic bacterial translocation occurs, and interaction with $Fc\alpha RI$ on Kupffer cells represent a third line of defence at the interface of mucosal and systemic immunity. Green, non- or anti-inflammatory and red, proinflammatory.

interaction with carbohydrate-recognizing receptors. Because uptake of SIgA was not accompanied by DC maturation, it was suggested that internalization of SIgA by DC might have a role in maintaining self-tolerance against commensal bacteria. SIgA furthermore has an important role as the first line of defense by preventing penetration of mucosal surfaces by microorganisms or foreign antigens (**Figure 3a**).¹ SIgA inhibits adherence of microorganisms, can agglutinate microbes, and interfere with bacterial motility by interacting with their flagella. Moreover, SIgA neutralizes bacterial products such as enzymes and toxins. However, opsonic activity is poor compared with dIgA or serum IgA as a result of (partial) blockage of the $Fc\alpha RI$ binding site by SC, which is consistent with a more anti-inflammatory role of SIgA. By contrast, both dIgA and serum IgA have a dual role in immunity, as they can show both anti- and proinflammatory roles.

In homeostatic conditions, dIgA serves as an intermediary that is transported through epithelial cells and released as SIgA. Although antibodies in general have a limited benefit against intracellular pathogens, dIgA can passively neutralize intracellular viruses because of trans-epithelial cell transport by intersecting virus particles and interfering with virus replication or assembly. IgA–virus complexes are subsequently removed via the lumen (**Figure 3a**). It was demonstrated that addition of specific anti-viral IgA to the basolateral surface of polarized epithelial cells *in vitro* decreased virus titers of Sendai virus, rotavirus, influenza, or HIV.^{145–149} Furthermore, culture with

polymeric IgA against toxin A of *Clostridium difficile* prevented destruction of epithelial monolayers.¹⁵⁰ *In vivo* evidence supporting that IgA transcytosis is required for viral inactivation was obtained when mice were only protected against rotavirus infection when IgA mAbs were given systemically, but not when they were provided via the lumen of the intestinal tract.¹⁵¹ A proinflammatory role for dIgA was recently proposed to eliminate pathogens that have successfully evaded the epithelial barrier. Bacteria, which are opsonized with dIgA, are efficiently phagocytosed by neutrophils.¹²³ Moreover, interaction of dIgA with neutrophil $Fc\alpha RI$ leads to release of leukotrien B4, which is a potent neutrophil chemoattractant (**Figure 3b**).¹³⁰ As such, it is hypothesized that a self-contained positive feedback loop is initiated, which results in enhanced recruitment of neutrophils, until clearance of invading pathogens has been achieved. Thus, dIgA functions as active second line of defense at mucosal surfaces by recruiting neutrophils.

The function of serum IgA in immunity is even more complicated and incompletely understood. It was demonstrated that IgA has the ability to downregulate IgG-mediated effector functions by transducing inhibitory signals through $Fc\alpha RI$.^{133–139} However, Kupffer cells in the liver, which are essential for elimination of bacteria in the portal circulation that have invaded via the gut, were demonstrated to express $Fc\alpha RI$.⁵⁶ In addition, serum IgA-opsonized *Escherichia coli* bacteria were efficiently phagocytosed by Kupffer cells, supporting a role for serum IgA in clearance of pathogens at the interface between

mucosal and systemic immunity. Thus, crosslinking of Fc α RI on Kupffer cells by serum IgA functions as the third line of defense (**Figure 3c**).

The importance of Fc α RI in mucosal infections was supported by two *in vivo* studies in human Fc α RI transgenic mice, which express Fc α RI on myeloid cells, similarly as in humans.^{103,152} When mice were infected with *Bordetella pertussis* that had been opsonized with human IgA, enhanced bacterial clearance in lungs of human Fc α RI transgenic mice was observed compared with non-transgenic littermates.¹²⁴ Furthermore, it was recently shown that passive transfer of human IgA mAbs against *Mycobacterium tuberculosis* protected human Fc α RI transgenic mice, but not Fc α RI-negative control mice, against *M. tuberculosis* infection.¹⁵³ To date, enhanced *in vitro* uptake of *E. coli*, *Streptococcus pneumoniae*, *Staphylococcus aureus*, *Porphyromonas gingivalis*, *Candida albicans*, *B. pertussis*, and *Neisseria meningitidis* by neutrophils after targeting of Fc α RI have been demonstrated.^{56,123,124,130,154–156}

Fc α RI AS THERAPEUTIC TARGET

Because Fc α RI has a dual role in immunity, as naturally occurring serum IgA induces inhibitory signals to dampen excessive immune responses, whereas crosslinking of Fc α RI during infection with IgA-opsonized pathogens results in proinflammatory responses; manipulation of Fc α RI function may offer novel promising therapeutic strategies. Until now, no (bispecific) Ab (BsAb)-based therapies that target Fc α RI exist in the clinic. Most of the research to investigate the potential of IgA mAbs for immunotherapeutic approaches has been based on *in vitro* experiments, but the increasing availability of suitable mouse models will greatly facilitate future *in vivo* studies. Two human Fc α RI transgenic mouse models are currently available, in which Fc α RI is either expressed on monocytes/macrophages or preferentially on neutrophilic granulocytes.^{103,157} The complementary Fc α RI expression in these models will allow studying the involvement of either cell type in disease or therapeutic efficacy.

Although it has been a challenge in the past to generate sufficient amounts of purified human IgA, technologies to reclone IgG mAbs, phage display, or transgenic plant technology has resulted in the generation of specific human IgA mAbs (as generation of sufficient BsAb is labor intensive with low yield, and as such may be unrealistic for clinical use). Moreover, the development of human IgA knock-in mouse model useable for standard hybridoma technology,¹⁵⁸ and new approaches in IgA purification techniques^{159–161} will furthermore allow the production of a continuous source of antigen-specific human IgA.

Fc α RI AS ANTI-INFLAMMATORY THERAPEUTIC TOOL

Two experimental inflammatory disease models have been described that demonstrate the potential of dampening excessive immune responses (initiated by other Ig-immune complexes) by inducing ITAMi signaling (**Figure 2b**) through Fc α RI. First, Fc α RI transgenic mice (expressing Fc α RI on monocytes/macrophages) that were immunized with IgE-

immune complexes developed bronchial hyper-reactivity after challenge with the antigen. Treatment with anti-Fc α RI Fab, which targets Fc α RI monovalently, hereby inducing ITAMi signaling—significantly reduced peribronchial inflammatory cell infiltration as well as symptoms.¹³¹ Second, both, decreased inflammatory cell infiltrates and fibrosis were observed in Fc α RI mice that were treated with anti-Fc α RI Fab in kidney inflammation models.¹⁶² Moreover, anti-Fc α RI Fabs were shown to induce apoptosis in Fc α RI-expressing mast cell transfectants, which prevented tumor development and halted the growth of established tumors.¹⁶³ As such, targeting ITAMi via Fc α RI can initiate either inhibitory signals or apoptosis, which may help to control disproportionate inflammation or tumor development. The therapeutic potential of targeting ITAMi has recently been reviewed by Monteiro *et al.*^{141,164}

TARGETING Fc α RI FOR TREATMENT OF INFECTIOUS DISEASES AND CANCER

Because IgA is predominantly present in mucosal areas where it shows several important functions (see above), therapies that aim to increase specific IgA titers against mucosal pathogens may help to fight (mucosal) infection. For instance, treatment with specific IgA protected mice against rotavirus, which is a diarrhea-causing pathogen.¹⁵¹ Interestingly, protective effect was only observed when IgA mAbs were given systemically, but not when IgA was presented via the lumen of the intestinal tract. These results therefore not only support the hypothesis that IgA transcytosis is required for intracellular viral inactivation but also suggest that transport of systemically delivered IgA via the pIgR route is not hampered by locally produced mucosal IgA.¹⁵¹ Similarly, increased presence of mucosal IgA by either passive transfer with specific IgA or through oral immunization prevented *Helicobacter felis*,¹⁶⁵ *Helicobacter pylori*,¹⁶⁶ influenza,^{167,168} or *Shigella flexneri*¹⁶⁹ infection. Importantly, as these experiments were performed in mice, which lack Fc α RI, the protective effect of IgA is presumably even more pronounced in humans. Mucosal administration of an HIV-1 vaccine resulted in resistance to the virus and production of virus-specific IgA with HIV-1 transcytosis-blocking properties in *Macaca mulatta* monkeys.¹⁷⁰ However, it was recently demonstrated that targeting Fc α RI directed neutrophils to destroy HIV-infected target cells.¹⁷¹ Because *M. mulatta* monkeys express Fc α RI, an active role for Fc α RI in eliciting protection is suggested, in addition to HIV neutralizing IgA Ab. Treatment of human Fc α RI transgenic mice with specific IgA induced enhanced protection against *B. pertussis* or *M. tuberculosis* infection, compared with non-transgenic littermates.^{124,153}

Fc α RI was furthermore proposed as a novel trigger molecule for mAb-based anti-cancer therapy.^{172,173} Although low expression of Fc α RI on DCs, poor ability of efficient Ag presentation, and no cross-presentation limits Fc α RI targeting on DCs for development of cancer vaccines, it was demonstrated that targeting Fc α RI efficiently recruits neutrophils as effector cells. *In vitro* experiments using therapeutic IgA1, IgA2, dIgA, chimeric IgA, or Fc α RI BsAb have provided promising results.^{127,128,172–186} For instance, neutrophils killed tumor cells much more effectively in

the presence of anti-(HER2/neu × FcαRI) BsAb, or anti-EpCAM IgA mAbs compared with an IgG counterpart. Similar superior ability of FcαRI to induce neutrophil-mediated tumor cell killing has now also been demonstrated for epidermal growth factor receptor, human leukocyte antigen class II, CD20, CD30, and carcinoembryonic antigen.^{127,172–174,176–186} One explanation for increased Ab-dependent-cellular cytotoxicity after targeting with IgA mAb or FcαRI BsAb may be the induction of neutrophil migration in the presence of IgA. Neutrophil accumulation and destruction of either mamma carcinoma or colon carcinoma colonies in a three-dimensional culture system was only observed when FcαRI was targeted, which is likely the result of leukotrien B4 release after crosslinking of FcαRI.^{127,130,187} However, it was previously shown that immature bone marrow neutrophils were not capable of killing tumor cells via FcγRI, whereas FcαRI efficiently induced Ab-dependent-cellular cytotoxicity.¹²⁸ It is therefore likely that the amplitude of signals mediated through FcαRI or FcγR also differs, as interaction of FcαRI with FcR γ-chain is stronger because of an electrostatic interaction (**Figure 2**) that is absent for FcγR.¹⁰² An attractive feature of recruiting neutrophils as effector cells is the fact that targeting FcαRI on neutrophils was recently demonstrated to induce autophagic tumor cell death (and necrosis to a lesser extend).¹⁸⁸ As such, neutrophils may be able to kill tumor cells with mutations in apoptotic pathways. Moreover, neutrophils attract Th17 cells, which have been shown to have a role in anti-tumor immunity,^{189,190} and moreover secrete cytokines and chemokines that attract other immune cells (monocytes or DCs), which may result in more generalized anti-tumor immune responses.¹⁹¹

CONCLUSION

FcαRI has a significant role *in vivo* for maintaining appropriate immune responses in both systemic and mucosal compartments. On one hand, FcαRI is involved in the prevention of superfluous immune responses that are initiated through other activating receptors. On the other hand, IgA can potently trigger protective immunity by crosslinking FcαRI on myeloid immune cells. The lack of FcαRI expression in mice has previously seriously hampered *in vivo* experiments to further elucidate the involvement of FcαRI in anti- and proinflammatory functions. Currently increasing availability of suitable genetically engineered mouse models will significantly facilitate future *in vivo* studies to establish the complex role of FcαRI in (mucosal) immunity and its potential as therapeutic target for human diseases.

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DISCLOSURE

The authors declared no conflict of interest.

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