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**Cutting Edge: Human FcRL4 and FcRL5 Are Receptors for IgA and IgG**

Timothy J. Wilson, Anja Fuchs, and Marco Colonna

Fc receptor-like (FcRL) proteins are a family of cellular receptors homologous to FcγRI and are predominantly expressed by B cells. They function to costimulate or inhibit BCR signaling through consensus ITAMs and ITIMs; however, the extracellular ligands of these receptors remain unknown or controversial. In this study, we tested the ability of human FcRL proteins to bind Igs and found FcRL4 and FcRL5 to be bona fide Fc receptors. In cellular binding assays, FcRL4 bound efficiently to IgA and FcRL5 binds all IgG isotypes with varied efficiency. Additionally, we generated mAbs capable of specifically blocking these interactions. Given their expression on activated B cells and potential for inhibitory signaling, FcRL4 and FcRL5 are likely to be important for immune complex-dependent human B cell regulation, and they represent novel therapeutic targets for receptor blockade therapies. *The Journal of Immunology, 2012, 188: 4741–4745.*

Receptors for Ig are essential mediators of immune responses. They bridge innate and adaptive immune responses by mediating phagocytosis of opsonized pathogens, Ab-dependent cell-mediated cytotoxicity, trans-epithelial Ig transport, and immune complex (IC) retention by follicular dendritic cells, as well as enhancing Ag presentation and regulating leukocyte activation (1, 2).

Human Fc receptor-like (FcRL) proteins are a family of receptors homologous to FcγRI in one or more of their Ig superfamily (IgSF) domains. Most are predominantly expressed by B cells (3–13), whereas FcRL3 can be additionally found on NK cells (14) and regulatory T cells (15), and FcRL6 is found mainly on NK cells and effector cytotoxic T cells (7, 8). Subsets analyses have refined our knowledge of the expression of each of the receptors. For example, FcRL5 appears to be broadly expressed by B cell subsets (3–5, 14), whereas FcRL4 expression is limited to a unique subset of tissue memory B cells, concentrated in the subepithelial and marginal zones of mucosal lymphoid tissues (16–18).

FcRL1–6 contain consensus ITAMs and/or ITIMs in their cytoplasmic tails. In addition to the six cell surface FcRL proteins are two intracellular receptors, FcRLA and FcRLB. These receptors lack transmembrane domains but contain a flexible C-terminal proline-rich stalk followed by a coiled-coil region (9–13, 19). Among the cell surface FcRs, only FcRL1 has been shown to exhibit costimulatory activity with the BCR (20). Of the others, FcRL2–5 have all been shown to recruit SHP-1 and inhibit BCR signaling (16, 21–23), whereas the functional outcome of tyrosine phosphorylation in FcRL6 remains elusive (7, 8).

Beyond their potential signaling capabilities, the biological roles of these receptors have remained unknown, largely due to the absence of knowledge about their extracellular ligands. Recently, FcRL6 has been reported to bind HLA-DR (24), and we and others have shown that intracellular FcRLA associates with multiple Ig isotypes within the lumen of the endoplasmic reticulum (25, 26). Given the homology of FcRL3–5 with FcγRI and the association between FcRLA and Igs, we decided to examine whether any of the FcRL proteins could bind Ig. In fact, we found that FcRL4 is an IgA receptor and FcRL5 is an IgG receptor. The ability of these receptors to bind Ig, along with their demonstrated ability to regulate B cell Ag receptor signaling, suggests a role for FcRL4 and FcRL5 in the regulation of B cell responses by Abs.

### Materials and Methods

**Ig binding assays**

To test for Ig binding by FcRL family members, cDNA encoding mouse FcRH3/FcRL5, human CD200R, FcRL1, FcRL3, FcRL4, FcRL5 (Open Biosystems), and FcRL6 (7) were ligated into pFLAG-CMV-3 (Sigma-Aldrich, St. Louis, MO). cDNA encoding human CD32 was ligated into pEF6 (Invitrogen). Proteins were expressed in 293 cells by transient transfection using Lipofectamine 2000. Transiently transfected 293 cells were used for Ab binding assays 36–42 h after transfection. Purified human Igs were obtained from Sigma-Aldrich in the following formats: IgG and IgM from serum; IgA from colostrum; and IgG1, IgG2, IgG3, and IgG4 (all k L chain) from myeloma plasma. Purified human IgA from serum was from Bethyl Laboratories (Montgomery, TX). For the heat aggregation assay, Igs were aggregated by heating to 60°C for 30 min. Igs were then diluted to 100 μg/ml in PBS/1% BSA. The 293 cells were incubated for 30 min on ice with the Igs and washed four times, followed by incubation with biotin-conjugated goat
FcR(λ); anti-human κ L chain Ab for IgG1, IgG2, IgG3, and IgG4, or a combination of biotinylated goat F(ab')2; anti-human κ and goat F(ab')2; anti-human λ Ab for total IgG and IgA (SouthernBiotech) for 20 min on ice. Cells were washed three times and incubated with a combination of allophycocyanin-conjugated streptavidin (Invitrogen) and FITC-conjugated anti-Flag Ab (M2; Sigma-Aldrich) for 20 min on ice. To detect CD32-expressing cells, a PE-conjugated anti-human CD32 Ab (Beckman Coulter) was added to CD32-transfected samples. Cells were washed twice and analyzed by flow cytometry on a FACS Calibur (BD Biosciences) for Ab binding. Dead cells were excluded by propidium iodide staining.

As an alternative to heat aggregation, IgG1-A was preincubated with equimolar quantities of biotinylated F(ab')2 fragments specific for human λ L chains to simulate ICs. The binding assay was performed as described. Secondary detection of bound Ig was achieved using streptavidin-PE.

Generation of mAbs
Mouse P815 cells stably expressing surface FcRL4 or FcRL5 were used for immunization. BALB/c mice were immunized three times at 2-wk intervals with paraformaldehyde-fixed FcRL4 or FcRL5 stable transfectants, first in CFA, then twice inIFA plus CpG 1826. Three days prior to fusion with SP2/0 cells, the final immunization was performed by i.p. injection of irradiated cells in PBS. Hybridoma clones were screened by FACS for the ability to stain FcRL4- or FcRL5-expressing P815 cells, but not the P815 parent cell line. Blocking Abs for FcRL4 were identified by FACS using a 1:1 mixture of FcRL4-expressing and nonexpressing P815 cells preincubated with anti-FcRL4 Ab. CFSE-labeled colostral IgA was heat aggregated and incubated with cells as described above. Blocking Abs for FcRL5 were identified in the primary screen for FcRL5-specific Abs. A 1:1 mixture of FcRL5-expressing and nonexpressing P815 cells was first incubated with supernatants from hybridoma clones. Goat F(ab')2-specific mouse IgG-FITC and anti-mouse IgM-FITC were used for secondary detection of FcRL5-specific mAbs. Blocking Abs were simultaneously identified among FcRL5-specific clones by the ability of FcRL5-specific mAbs to block the nonspecific association of goat anti-rabbit IgG-Alfa Fluor 647 with FcRL5-expressing P815 cells. Animal procedures have been reviewed and approved by the Washington University Animal Care and Use Committee.

For epitope mapping of anti-FcRL5 Abs, FcRL5 cDNA encoding IgSF domains 1–3, 4–6, or 7–9 were amplified by PCR and ligated into the expression vector pDisplay (Clontech). The 293 cells transiently transfected with FcRL4, CD32b, or CD200R were incubated with 293 cells transiently transfected with FcRL proteins or control CD320R or CD32b were incubated with heat-aggregated IgG, IgA, IgE, or IgM. No binding to IgE or IgM was detected for any of the FcRL family members (not shown), whereas IgG bound FcRL5 and CD32, and human serum IgA was found to bind FcRL4 (Fig. 1).

Because CD32 is known to have differential affinity for the individual IgG subtypes (27), we examined the ability of FcRL5 to bind human IgG isotypes. Human IgG1-κ, IgG2-κ, IgG3-κ, and IgG4-κ were heat aggregated and incubated with 293 cells transiently transfected with FcRL4, FcRL5, CD32b, or CD200R. Whereas no association was consistently found between FcRL4 and IgG1 or IgG2, binding of IgG3 and IgG4 by FcRL4 was weak but detectable. FcRL5 was able to bind all IgG isotypes, showing consistently stronger binding to IgG1 and IgG2 over IgG3, while associating only weakly with IgG4. The binding of CD32b in this assay to IgGs showed the qualitative preferences of IgG3 > IgG1 > IgG4 > IgG2 (Fig. 2A). This pattern precisely matches previously identified binding specificities for CD32b (27), providing validation for the observed binding specificity of FcRL4 and FcRL5.

![Figure 1](http://www.jimmunol.org/). FcRL4 and FcRL5 bind human IgGs. HEK293T cells were transiently transfected with human cDNA for FcRL1, 3, 4, 5, and 6 and CD200R expressed in pFLAG-CMV-3 or CD32 in pEF6. Binding of heat-aggregated human serum IgG and IgA (y-axis) to cells expressing human FcRL4 and FcRL5 was easily observable by FACS. Cell surface expression of each protein was confirmed using FITC-conjugated anti-FLAG M2 Ab or by staining with anti-CD32 Ab. Binding to CD200R and CD32 are included as negative and positive controls, respectively. No binding of human IgM to any of the FcRL proteins was observed (not shown).

**Results**

FcRL4 and FcRL5 are Ig receptors

FcRL proteins are diverse in their expression patterns and functional roles among human B cell subsets. Despite the growing body of literature surrounding the expression patterns of these receptors and their associations with disease, little is known about the extracellular interactions on the cell surface. Because some of the FcRL family members are highly homologous to FcγRI, and anecdotal reports in the literature suggest Ig binding potential (4, 14), the ability of FcRL family members to bind Igs was assayed. In a preliminary test, FcRL5 was able to bind heat-aggregated IgG1. Weak binding of IgG1 to FcRL3 and FcRL4 was also observed, but these interactions were not consistently reproducible. No detectable binding to IgG1 was observed by control transfected or untransfected cells, or by cells transfected with FcRL1 or FcRL6 (not shown). To analyze binding specificity of FcRL proteins, 293 cells transiently expressing FcRL proteins or control CD200R or CD32b were incubated with heat-aggregated IgG, IgA, IgE, or IgM. No binding to IgE or IgM was detected for any of the FcRL family members (not shown), whereas IgG bound FcRL5 and CD32, and human serum IgA was found to bind FcRL4 (Fig. 1).

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The N-terminal domains of FcRL5 are necessary and sufficient for IC binding

FcRL5 contains nine IgSF domains, with the N-terminal three domains sharing homology with those of FcγRI. We predicted that these three domains would be sufficient for Ig binding. To test this hypothesis, cDNA encoding domains 1–3, 4–6, or 7–9 of FcRL5 were expressed transiently in 293 cells using the vector pDisplay, and analyzed for the ability to bind Ig. Whereas surface-expressed forms of FcRL5 domains 4–6 or 7–9 were unable to bind IgG1, cells expressing FcRL5 domains 1–3 readily bound IgG1 mock ICs (Fig. 2B). These data indicate that the first three domains of FcRL5 are necessary and sufficient for Ig binding.

Development of blocking Abs for FcRL4 and FcRL5

To further study the expression and function of FcRL4 and FcRL5, a panel of mAbs specific for these receptors was generated. Hybridoma clones were screened for reactivity to either FcRL4- or FcRL5-transfected P815 cells, as well as the ability to block the binding of Ab to these receptors. Because FcRL5 contains nine Ig domains, anti-FcRL5 mAbs were further characterized for epitope location. Using 293 cells transiently transfected with FcRL5 domains 1–3, 4–6, or 7–9, we determined that mAb 509F6 and mAb 501H9 (both IgG2a) bind FcRL5 domains 1–3 and 4–6, respectively (Fig. 3A). By preincubating these Abs with FcRL5-transfected 293 cells, we determined that mb 509F6 can specifically block the association of FcRL5 with mock ICs, whereas 501H9 is a non-blocking Ab. (Fig. 3B) Similarly, anti-FcRL4 mAb 413D12 is able to block the binding of FcRL4-transfected P815 cells with heat-aggregated IgA, whereas isotype-matched anti-FcRL4 mAb 418C8 (IgG2b) has no blocking potential (Fig. 3C). These data confirm the importance of the FcR-homologous domains of FcRL5 in Ab binding, and further support a role for FcRL4 and FcRL5 as Ig receptors.

FcRL4 expression has been described as limited to a memory B cell population concentrated within mucosal sites, and with variable expression of the memory B cell marker CD27 (4, 14, 17, 18). Staining human tonsil sections and cell suspensions with anti-FcRL4 (413D12) was largely consistent with the published expression patterns. FcRL4+ cells were found to be abundant in the tonsillar submucosa and marginal zones (Supplemental Fig. 1). Additionally, the proportion of FcRL4+ cells that were also CD27+ showed substantial variation among individuals, ranging from 50 to 70% in five tonsillar preparations, whereas FcRL5 expression was found among CD27−, CD27+, and CD27high cells (Fig. 3E and data not shown). Interestingly, all FcRL4+ cells also expressed FcRL5 at a high level (Fig. 3F). The combined expression of these two inhibitory Fc receptors may help explain the reported resistance of FcRL4+ memory cells to stimulation through the BCR (18).

Discussion

Human FcRL proteins are an important family of immunomodulatory receptors. Earlier studies have indicated an inhibitory function for both FcRL4 and FcRL5 through the recruitment of SHP-1 (16, 21), and our own unpublished observations have confirmed these findings. Anecdotal reports have previously suggested that FcRL4 and FcRL5 might be Fc

FIGURE 2. Specificity of FcRL4 and FcRL5 for Igs. (A) Human IgG isotypes were assayed by FACS for binding to FcRL4, FcRL5, and CD32. Dark lines indicate aggregated IgG bound to the indicated receptor-transfected 293 cells. Gray shading indicates binding of Igs to negative control CD200R-transfected cells. Histograms show populations gated on receptor-expressing cells by simultaneous staining with anti-Flag (FcRL4, FcRL5, and CD200R) or anti-CD32 Abs. (B) Portions of the ectodomain of FcRL5 comprising Ig domains 1–3, 4–6, or 7–9 were expressed on the surface of 293 cells. Binding of IgG1 mock ICs to FcRL5 domains 1–3 was measured by FACS.

FIGURE 3. Specific blockade of Ig binding to FcRL4 and FcRL5 by mAbs. (A) FACS staining of FcRL5 domains 1–3 or 4–6 with mAbs 501H9 and 509F6. (B) Preincubation of Ab 509F6 with FcRL5-expressing 293 cells prevents binding of FcRL5-transfected 293 cells to mock ICs. (C) Blockade of IgA binding to FcRL4 by anti-FcRL4 clone 413D12 (IgG2b) but not isotype-matched anti-FcRL4 clone 418C8. (D) Staining of tonsillar B cells with anti-FcRL4 (413D12) and anti-FcRL5 (509F6) mAbs or isotype controls. (Plots show cells following gating on live CD19+ cells.) (E) Anti-FcRL4 and anti-FcRL5 identify mixed populations of CD27+ and CD27− tonsillar B cells. (F) All FcRL4+ tonsillar B cells coexpress FcRL5. For (E) and (F), plots are representative of results obtained with tonsil samples from five different donors.
receptors (4, 14). However, these observations were not initially reproducible by all groups studying these receptors (28), and so there has been no consensus ligand for these receptors. Our data clearly show that FcRL4 can bind IgA and FcRL5 can bind IgG.

Nonspecific binding of soluble monomeric mouse and goat IgG by FcRL5-transfected P815 cells was observed while screening anti-FcRL5 hybridoma clones, leading initially to the conclusion that FcRL5 might be a high-affinity receptor. In contrast, binding of soluble monomeric IgG to FcRL4 or FcRL5-transfected 293 cells was not readily observable in FACs-based assays, indicating that these receptors are likely to be low- to medium-affinity Fc receptors. Because P815 cells express classical IgG Fc receptors, these observations may instead indicate that FcRL5 and classical Fc receptors can bind IgG cooperatively on the cell surface. Structural studies of the interactions between Abs and the classical FcRs have shown a small conformational change in the Ig Fc region upon receptor binding, ensuring a 1:1 IgG/FcR stoichiometry (29).

Although the FcRL proteins are homologous to FcγRI in their three N-terminal IgSF domains, they lack sequence homology with classical receptors at the putative Ig-binding interface (28), leading to the assumption that FcRL proteins would not bind Abs. A model that includes cooperative IgG binding on B cells by FcγRIIb and FcRL5 explains observed binding phenomena and is consistent with the presence of structural similarity between the two receptors, but differences in the specific contact residues.

The finding that human B cells contain additional inhibitory Fc receptors is critically important for understanding the regulation of human B cell responses. Importantly, FcRL4 is the only known inhibitory receptor for IgA. Given the abundance of FcRL4 on mucosal memory B cells, IgA responses to mucosal Ags may be substantially regulated by FcRL4. Although it is possible that FcRL5 is somewhat redundant with FcγRIIb in humans, the potential for simultaneous recruitment of SHIP-1 by FcγRIIb and SHP-1 to FcRL5 provides a substantial barrier to recurrent activation of B cells. Together, FcRL4 and FcRL5 may limit B cell activation against chronic pathogens, self Ag, or commensal flora.

Mouse studies have clearly shown the importance of FcγRIIb in B cell regulation and autoimmune disease (1). Of the mouse cell surface FcR proteins, only FcγRIIIa/FcRL5 has a structure similar to FcγR4 or FcγR5. In our assays, mouse FcγRIII/FcRL5 failed to bind heat-aggregated mouse IgG (data not shown), and so FcγRIIb remains the sole inhibitory Fc receptor on mouse B cells. As a result of this functional difference, the regulatory function of human FcRL5 is unlikely to be determined from the study of FcγR3/FcRL5 knockout mice. Instead, FcRL4 or FcRL5 transgenes, expressed under appropriate promoters, would be more relevant for in vivo studies.

With extracellular ligands known and blocking Abs available, FcRL4 and FcRL5 are attractive as therapeutic targets. First, FcRL4 and FcRL5 are both expressed by previously activated B cells. By blocking their ability to bind existing Abs, the Ab titer and repertoire generated by pathogen or tumor vaccines may be augmented. Second, downregulation of FcRL4 in “exhausted” memory B cells from HIV patients was shown to enhance their responsiveness to HIV (30). Blocking FcRL4 might lead to renewed generation of neutralizing Abs for HIV and other chronic viral infections. Lastly, the expression of FcRL5 by Ag-experienced B cells makes it an attractive target for Ab-mediated cellular depletion. Autoimmune disorders that are dependent on autoantibodies for their pathology might be successfully managed by depleting FcRL5+ cells. Additionally, the prevalence of FcRL5 on lymphomas with translocations at 1q21 (4) provides a specific target for depletion of tumors of this type. In conclusion, further study of the fundamental biology of FcRL proteins in general, and of FcRL4 and FcRL5 in particular, offer exciting possibilities for understanding human B cell biology and mucosal immunity.

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Disclosures

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