**Role of Novel Rat-specific Fc Receptor in Macrophage Activation Associated with Crescentic Glomerulonephritis**

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**Background:** Fc receptor-mediated macrophage activation is a major cause of glomerular damage in crescentic glomerulonephritis.

**Results:** We investigated the role of a novel rat Fc receptor, Fcgr3-rs, in human and rat macrophage activation.

**Conclusion:** We showed that this receptor prevents the cell signaling of its paralogue (Fcgr3).

**Significance:** These results provide a novel way to inhibit Fc receptor-mediated cell activation in macrophages.

Crescentic glomerulonephritis (Crgn) is a complex disease where the initial insult is often the glomerular deposition of antibodies against intrinsic or deposited antigens in the glomerulus. The role of Fc receptors in the induction and progression of Crgn is increasingly recognized, and our previous studies have shown that copy number variation in Fcgr3 partially explains the genetic susceptibility of the Wistar-Kyoto (WKY) rat to nephrotoxic nephritis, a rat model of Crgn. The Fcgr3-related sequence (Fcgr3-rs) is a novel rat-specific Fc receptor with a cytoplasmic domain 6 amino acids longer than its paralogue, Fcgr3. The Fcgr3-rs gene is deleted from the WKY rat genome, and this deletion is associated with enhanced macrophage activation in this strain. Here, we investigated the mechanism by which the deletion of Fcgr3-rs in the WKY strain leads to increased macrophage activation. By lentivirus-mediated gene delivery, we generated stably transduced U937 cells expressing either Fcgr3-rs or Fcgr3. In these cells, which lack endogenous Fcgr3 receptors, we show that Fcgr3-rs interacts with the common Fc-γ chain but that Fc receptor-mediated phagocytosis and signaling are defective. Furthermore, in primary macrophages, expression of Fcgr3-rs inhibits Fc receptor-mediated functions, because WKY bone marrow-derived macrophages transduced with Fcgr3-rs had significantly reduced phagocytic activity. This inhibitory effect on phagocytosis was mediated by the novel cytoplasmic domain of Fcgr3-rs. These results suggest that Fcgr3-rs may act to inhibit Fcgr3-mediated signaling and phagocytosis and could be considered as a novel mechanism in the modulation of Fc receptor-mediated cell activation in autoimmune diseases.

In crescentic glomerulonephritis, leukocytes, especially monocytes/macrophages, infiltrating the glomerulus play a critical role in the development of the disease by recognizing immunoglobulins and complement factors via cell surface receptors (1). One of the checkpoints regulating the immune response in glomerular inflammation is through leukocyte cell surface receptors for the Fc (fragment crystallizable) portion of IgG. The family of Fc receptors can regulate the fine balance between pro-inflammatory versus anti-inflammatory states of cell activation, and there is increasing evidence suggesting that genetic susceptibility to autoimmune diseases is partly dependent on Fc receptor-mediated activating and inhibitory signaling pathways determining the net signal (pro- or anti-inflammatory) in the progression of the disease (2, 3). In systemic lupus erythematosus, aberrant expression or the presence of allelic variants of FcγRs with altered functionality have been reported to contribute to the pathogenesis of the disease (4).

There are two distinct classes of Fc receptors: the activatory and the inhibitory receptors. Most activatory receptors serve as the ligand-binding component of a receptor that also includes a signal transducing molecule containing an immunoreceptor tyrosine-based activation motif (ITAM)² (5). In monocytes and macrophages, this role is fulfilled by the common γ chain. Signaling via its ITAM motif triggers the activation of multiple downstream signaling pathways including the activation of Src and Syk kinases, calcium mobilization, and NF-κB activation, leading ultimately to cellular responses such as phagocytosis and cytokine production. The importance of activatory Fcγ receptors in glomerulonephritis has been shown using mice engineered to lack the ITAM-bearing Fc receptor γ chain. We and others have shown that these mice are protected from glomerulonephritis induced by antibodies to glomerular basement membrane (6–8). In contrast, the inhibitory Fc receptor FcγRIIB has an immunoreceptor tyrosine-based inhibitory

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² The abbreviations used are: ITAM, immunoreceptor tyrosine-based activation motif; FcR, Fc receptor; BMDM, bone marrow-derived macrophage; TNF, nephrotoxic nephritis; Crgn, Crescentic glomerulonephritis; WKY, Wistar-Kyoto; qRT, quantitative RT; PMA, phorbol 12-myristate 13-acetate.
motif and does not use the common γ chain (3). Activatory and inhibitory Fcγ receptors are often expressed on the same cell, and when co-aggregated by IgG antibody, the net signal to the cell depends on the sum total of the activator and inhibitor signals, thus setting thresholds for effector cell responses (3).

In our previous studies of nephrotic nephritis (NTN), a model of crescentic glomerulonephritis in the Wistar-Kyoto (WKY) rat, we performed genome-wide linkage analysis for glomerular crescents, macrophage infiltration, and proteinuria in an F2 population derived from NTN-susceptible WKY and NTN-resistant Lewis rats (9). The most significant linkage (logarithm of odds > 8) was obtained with a quantitative trait locus mapping to chromosome 13 (Crgn1) accounting, respectively, for 21.8, 16.7, and 12.9% of the genetic variance in crescent formation, proteinuria, and macrophage infiltration. Positional cloning of Crgn1 led to the identification and functional characterization of the Fcgr3-related sequence (Fcgr3-rs), a novel rat-specific activatory Fc receptor arising from copy number variation in the Fcgr3 gene (9). The genomic rearrangement in the Fcgr3 gene is such that there is a duplication of exon 5 in the NTN-resistant Lewis genomic DNA with the presence of a shorter exon lacking a sequence of 226 bp in its 3′-untranslated region. This shorter exon 5 was deleted in the WKY genome, suggesting a copy number variation in the Fcgr3 gene. Sequence analysis of this shorter copy of exon 5 revealed deletion of a single guanine nucleotide at position 129 (ΔG129) in the coding sequence of the cytoplasmic domain, resulting in a frameshift and generating a novel cytoplasmic domain 6 amino acids longer than that encoded by Fcgr3. We designated this copy number variant Fcgr3-related sequence (Fcgr3-rs) and showed that this novel variant is transcribed and translated in the Lewis strain, whereas it is deleted from the WKY genome (9).

Given the increased macrophage activity and associated NTN susceptibility in the WKY rat that lacks the Fcgr3-rs molecule, it is important to understand how the presence of this molecule might be influencing macrophage activity. Thus, we have studied the role of Fcgr3-rs in macrophage activation. We first investigated whether the genomic duplication/deletion event giving rise to Fcgr3-rs occurred during the derivation of inbred Wistar-related colonies. We have then generated U937 cells stably expressing either Fcgr3 or Fcgr3-rs and have studied the role of Fcgr3-rs in macrophage activation both in U937 cells and WKY bone marrow-derived macrophages (BMDMs) transduced with Fcgr3-rs. Our results show that the absence of Fcgr3-rs is not specific to the WKY strain but is widely distributed throughout the rat phylogenetic tree. Furthermore, in both human monocyte cell lines and primary rat macrophages, expression of Fcgr3-rs inhibits Fcgr3-mediated signaling and phagocytosis, suggesting that the rat possesses a unique mechanism by which Fc receptor-mediated cell activation can be modulated.

**EXPERIMENTAL PROCEDURES**

**Animals**—WKY (WKY/NCrl) rats were purchased from Charles River. All of the procedures were performed in accordance with the United Kingdom Animals (Scientific Procedures) Act.
after 16 h. Media containing virus were collected after a subsequent 24 and 48 h of culture. Insoluble debris was removed by centrifugation and sterilized by filtration through 0.2-μm filters. This crude viral supernatant was stored at -80 °C until use. Titers of viral preparations produced in this way were established by infection of U937 cells as follows. Briefly, U937 cells were mixed with varying dilutions of viral supernatant (typically 1/5 to 1/500) in the presence of 8 mg/ml polybrene (hexadimethrine bromide; Sigma). The cells were “spinculated” by spinning at 2200 rpm for 2 h at 30 °C. After 72 h, the percentage of GFP-positive cells (FITC channel) was determined by FACS analysis (using a LSRII FACS). The lowest dilution factor with GFP+ score of <20% was used to calculate viral titer.

U937 Lentiviral Line Production—Human monoblastic leukemia U937 cells were infected with lentivirus carrying the gene of interest at a ratio of 1:1 (cells:virus) as described above. The cells were then cultured overnight before removal of virus containing media. Cell surface Myc-positive cells (Myc-Fcgr3 or Myc-Fcgr3RS) were isolated by panning on 10-cm Petri dishes coated with anti-Myc antibody (Millipore). Adherent cells were expanded in culture and used for subsequent experiments. GFP-positive cells expressing pCSGW (no surface Myc) were isolated by cell sorting based on FITC fluorescence. Subsequent analysis of cell lines by FACS analysis confirmed 90–100% of cells to be FITC-positive.

FcR Stimulation—The cells were resuspended in ice-cold serum-free RPMI and anti-Myc antibody (Millipore) added to a final concentration of 130 μg/ml. After 60 min of incubation on ice, the cells were washed three times and resuspended in serum free RPMI with 8 mg/ml goat anti-mouse IgGF(ab')2. The cells were then incubated at 37 °C for the indicated times. The reaction was stopped by the addition of gel sample buffer (2% SDS, 10% glycerol, 0.5% β-mercaptoethanol, 50 mM Tris, pH 6.8).

Immunoprecipitations and Western Blot Analysis—The cells were lysed in ice-cold lysis buffer (25 mM Tris, pH 7.6, 150 mM NaCl, 1 mM EDTA, 0.1% Triton X-100, pH 7.8, 1% digitonin), containing protease inhibitor mixture (Sigma) with 1 mM DTT, 10 mM NaF, 100 mM sodium orthovanadate). The nuclei and cell debris were removed by centrifugation, and supernatants were precleared with protein G-Sepharose. Myc-labeled proteins were then precipitated with anti-Myc antibody (Millipore) or isotype matched control antibody and protein G-Sepharose for 1.5 h. Immunoprecipitated complexes were washed in lysis buffer and were resolved on 4–20% SDS-PAGE gels (Criterion).

After electrotransfer on to nitrocellulose membrane, the membranes were blocked in 2% BSA/TBS/Tween (0.05%) and probed with the appropriate antibodies at 4 °C overnight followed by HRP-conjugated secondary antibody. Blots were developed using ECL (Amersham Biosciences).

ELISAs—At 24 h after stimulation, the supernatants were harvested. The concentrations of TNFα and IL-10 were determined by ELISA (PharMingen) according to the manufacturer’s instructions. Absorbance was read and analyzed at 450 nm on a spectrophotometric ELISA plate reader (Labsystems Multiskan Biochromic) using the Ascent software program.

RESULTS

The Phylogeny of Genomic Deletion of Fcgr3-rs in Laboratory Rat—Exon 5 of Fcgr3-rs contains a deletion of a single guanine nucleotide at position 129 (ΔG129) in the coding sequence of the cytoplasmic domain (9). When compared with Fcgr3, this ΔG129 variant results in a frameshift in the Fcgr3-rs coding sequence and generates a novel cytoplasmic domain 6 amino acids longer than that encoded by Fcgr3 that was not found in any other species (Fig. 1). We have previously shown that Fcgr3-rs was deleted from NTN-susceptible Kyoto-derived strains (9), but to investigate the distribution of this deletion throughout the rat genealogical tree, we carried out a phylogenetic tree analysis of 94 inbred laboratory strains and 40 of their substrains (Fig. 2). The results show that the deletion of Fcgr3-rs is widespread throughout the rat phylogeny (32 of 94 strains), suggesting that the origin of the deletion event is earlier than the derivation of these strains. We also investigated the relative levels of Fcgr3 and Fcgr3-rs mRNA levels in Lewis rat BMDMs by qRT-PCR and showed that although there is relatively
increased Fcgr3 expression when compared with Fcgr3-rs, LPS stimulation did not change the relative levels of both receptors (Table 1).

**U937 Cells Expressing Either Fcgr3 or Fcgr3-rs: Surface Expression and Interaction with γ Chain**—The U937 cell line does not express any endogenous FcγRIIIA (11) and is therefore an ideal tool for the analysis of FcγRIII receptors, allowing direct functional comparison between the rat orthologues Fcgr3 and Fcgr3-rs. Thus, to study the role of Fcgr3-rs in macrophage activation, we generated U937 cells stably transfected with lentivirus constructs expressing the protein of interest. U937 cells were infected with either Myc-tagged Fcgr3 (Myc-Fcgr3), Myc tagged-Fcgr3-rs (Myc-Fcgr3RS), or empty (pCSGW) lentiviral vectors. All of the lentiviral vectors contained GFP. Expression of Myc-Fcgr3 and Myc-Fcgr3-rs was confirmed by fluorescence microscopy, detecting GFP + cells and also by a Myc Western blot on lysates from stably transduced U937 cells (Fig. 3A). We then investigated whether Myc-Fcgr3 and Myc-Fcgr3-rs are expressed on the surface of U937 cells. Flow cytometry analysis using an anti-Myc-APC mouse antibody (mAb) showed that both Myc-Fcgr3 and Myc-Fcgr3-rs are expressed on the cell surface at similar levels in these cells (Fig. 3B). Because Fcgr3 transport to and expression at the cell surface is known to be dependent on its association with the γ chain (12), these findings implied that both the Fcgr3 and Fcgr3-rs molecules were associated with the γ chain at the cell surface. Co-immunoprecipitation analysis of U937 cells expressing either Fcgr3 or Fcgr3-rs demonstrated that this was indeed the case (Fig. 3C). Thus, although Fcgr3-rs has an additional cytoplasmic domain 6 amino acids longer than its parologue Fcgr3, it binds the γ-chain on the surface of stably transfected and PMA-differentiated U937 macrophages.

**Fcgr3-rs-mediated Signaling and Phagocytosis in U937 Cells**—In most macrophage Fc receptor complexes (with the exception of FcγRII), the ITAM-containing common γ-chain is necessary for downstream signaling (13, 14). Thus, the dimeric γ-chain forms multichain complexes with the ligand binding α-chain of FcγRII to allow signal transduction (3). This signaling starts with tyrosine phosphorylation of the ITAMs by kinases of the SRC family (15, 16) and leads to the recruitment and subsequent phosphorylation of the Syk kinase, followed by the activation of various downstream targets such as PI3K and NF-κB. Subsequent changes in gene expression led to the induction of inflammatory cytokines such as TNFα. We were therefore keen to understand whether the Fcgr3-rs is able to transduce signals similar to those described for Fcgr3. However, despite its interaction with the γ-chain, cross-linking of Fcgr3-rs with anti-Myc antibody does not lead to phosphorylation of Syk or the degradation of 1kDa (Fig. 4A). This was not the case for the U937 macrophages stably transfused with Fcgr3 (Fig. 4B).

Furthermore, Fc receptor-mediated TNFα production and IgG-opsonized bead phagocytosis were also significantly less in U937 macrophages expressing Fcgr3-rs when compared with U937 macrophages stably transfused with Fcgr3 (Fig. 4B). Taken together, these results suggest that the rat-specific Fcgr3-rs is unable to mediate the same intracellular signals as the wild type receptor.

**Fcgr3-rs and TLR4-mediated TNFα and IL-1β Expression Levels**—To test whether the relative reduction in TNFα production in U937 cells transfected with Myc-Fcgr3RS is specific to receptor cross-linking, we have stimulated the stably transfused FcγRIIIA, FcγRIIA, FcγRIIIA, and FCGR3 in other species. Fcgr3-rs has lost the terminal PDKQ sequence (highlighted) that is highly conserved across species. *a, amino acid prediction based on human FCGR3A using Ensembl Genome Browser.

**FIGURE 1. Rat-specific Fcgr3-rs is a novel Fc receptor with a unique cytoplasmic domain 6 amino acids longer than Fcgr3**. Shown is a comparison of amino acid sequences of transmembrane and cytoplasmic domains of rat Fcgr3 and Fcgr3-rs, human Fcgr3, and FCGR3 in other species. Fcgr3-rs has lost the terminal PQDK sequence (highlighted) that is highly conserved across species. *a, amino acid prediction based on human FCGR3A using Ensembl Genome Browser.
result of altered FcR binding. Thus, we also generated an additional lentiviral construct encoding a chimeric protein to specifically test the effect of the novel cytoplasmic domain of Fcgr3-rs on Fc receptor-mediated phagocytosis. Fcgr3-ΔG contains the wild type Fcgr3 extracellular, transmembrane, and cytoplasmic domains with the presence of the Fcgr3-rs-specific novel cytoplasmic domain (9) (Fig. 6A).

We first measured the expression of Fcgr3 by using primers specific for the transmembrane domain, thereby allowing differential amplification between Fcgr3 and Fcgr3-rs transcripts (Fig. 6B). We then measured the number of opsonized beads phagocytosed by 200 WKY BMDMs transduced with each different lentiviral construct.

Phagocytosis was significantly reduced in Fcgr3-rs transduced WKY BMDMs when compared with those transduced with pCSGW (Fig. 6C). This inhibitory effect is clearly mediated by the novel cytoplasmic domain of Fcgr3-rs, because similar reductions in phagocytic activity were seen in WKY BMDMs transduced either with Fcgr3-ΔG or with Fcgr3-rs (Fig. 6C). Taken together, these data suggest that the presence of FcRs containing the related sequence cytoplasmic domain is able to reduce phagocytosis mediated by endogenous FcRs in the WKY BMDMs.

To further understand the mechanism of action of the Fcgr3-rs, we then investigated how the Fcgr3 and Fcgr3-rs
interact with TLR4-mediated TNFα production following LPS stimulation. U937 cells transduced with FcRs (U937 + Myc-Fcgr3, U937 + Myc-Fcgr3RS, and U937 + Myc-Fcgr3ΔG) were cross-linked with anti-Myc antibodies and IgG F(ab’)_2. Following washing, the cells were stimulated with LPS for 5 h. TNFα and IL-10 protein levels were measured by sandwich ELISA.

In the absence of any additional FcR expression in these cells, treatment with Ab-Fab complexes resulted in a 38% decrease in TNF production. This was further decreased in
the presence of the Fcgr3 molecule to 50% of that seen in
cells that are not cross-linked, revealing that signals from
Fcgr3 act to reduce TLR4-induced TNF production in U937
cells (Fig. 6D). In contrast, the presence of either the Fcgr3-rs
or FcgR3ΔG did not result in any changes in TNFα produc-
tion compared with cells that are not cross-linked. Levels of
IL-10 produced in these cells did not differ significantly
between Fcgr3 and FcgR3-rs or FcgR3-ΔG cells (Fig. 6E),
revealing that the Fcgr3-rs is not functioning in these cells by
increasing levels of this anti-inflammatory cytokine.

DISCUSSION

There is considerable evidence from experimental models
of crescentic glomerulonephritis that Fcγ receptors play a central
role in disease progression. In mice, the Fcγ chain-deficient
mice (Fcγγ−/−) lack surface expression of both murine activa-
tor Fc receptors I, III, and IV, and the high affinity IgE receptor
(FcRI) (18). Several studies have shown that Fcγγ−/− mice are
protected from disease in accelerated nephrotic nephritis (6,
7, 19), and we have shown that the disease is mediated by Fcγ
receptors on circulating leukocytes and not intrinsic renal cells
(8). Our previous studies in the NTN model in the Wistar Kyoto
rat showed that the disease architecture is complex and highly
heritable, with seven quantitative trait loci (Crgn1–7) linked to
the percentage of glomerular crescents and controlling mainly
macrophage activation through distinct transcriptome profiles
(9, 20–22). In this model, we showed that copy number varia-
tion in rat Fcgr3 locus led to the deletion of Fcgr3-rs in the WKY
rat, predisposing this strain to enhanced macrophage activity
(9).

The purpose of the current study was to understand the
mechanism by which the genomic deletion of Fcgr3-rs leads to
macrophage overactivation and whether the novel cytoplasmic
domain binds to the common γ chain to trigger downstream
signaling pathways. We first investigated the distribution of
Fcgr3-rs deletion throughout the rat phylogeny and found that
the deletion was widespread. It was previously reported that
five outbred Wistar and one Long-Evans stocks have Fcgr3-rs
(23), suggesting that the deletion has originated from the ances-
tor outbred Wistar colony in the Wistar institute in the United
States. We now confirm and extend these findings because all of
the laboratory rat strains that we genotyped for Fcgr3 copy

FIGURE 4. Fc receptor-mediated signaling and phagocytosis in U937 cells
stably transduced with Myc-Fcgr3RS. A, Fc receptor stimulation after cross-
linking with anti-Myc Ab for a maximum of 20 min and analysis of Syk phos-
phorylation and IkBα degradation in stably transduced U937-derived macro-
phages. B, TNFα levels were assessed by sandwich ELISA after cross-linking
stably transduced U937 macrophages by anti-Myc mAb. Supernatants were
collected after 48 h of incubation. Fc receptor-mediated bead phagocytosis
was assayed by incubating macrophages with IgG opsonized latex beads.
Nonopsonized beads did not show any phagocytosis. *, p < 0.001 when com-
pared with U937 + Myc-Fcgr3.

FIGURE 5. Fcgr3-rs does not affect TLR4-mediated Tnfa and Il1b1 expressions in stably transduced U937 cells. Stably transduced U937 cells were
differentiated in macrophages after adding PMA (100 nM) for 48 h and stimulated with LPS (1 μg/ml) for 3 h. Tnfa and Il1b1 gene expression were measured by
qRT-PCR. ns, not significant.
number variation are derived from outbred Wistar and Long-Evans stocks.

Previously, to investigate how Fcgr3-rs exerts its inhibitory effect on Fc receptor signaling, we have transfected COS7 cells and have shown that although Fcgr3-rs is expressed on the cell surface, it is less efficient than Fcgr3 in mediating phagocytosis (9). It also had an inhibitory effect on Fcgr3-mediated phagocytosis when both receptors were co-transfected in COS7 cells.

**FIGURE 6.** Fcgr3-rs inhibits Fc receptor-mediated phagocytosis in WKY BMDMs. A, a schematic representation of the functional differences between Fcgr3 and Fcgr3-rs. Although there are amino acid changes in the Fcgr3-rs extracellular, transmembrane, and cytoplasmic domains (indicated by bars) when compared with Fcgr3, the major structural difference is the presence of a novel cytoplasmic domain. The chimeric Fcgr3-ΔG has the extracellular transmembrane and cytoplasmic domains identical to Fcgr3, the only difference being the presence of the novel cytoplasmic domain. B, Fcgr3 expression was assessed by qRT-PCR in stably transduced U937 cells using specific primers designed based on polymorphisms in the transmembrane domain differentiating between Fcgr3 and Fcgr3-rs. The expression of Myc-Fcgr3-ΔG was also assessed and showed similar expression levels as Fcgr3. This is because Fcgr3 and Fcgr3-ΔG share the same sequence in the transmembrane domain. C, WKY bone marrow-derived cells were transfected with lentiviral constructs co-expressing GFP with either Myc-tagged Fcgr3-rs (Myc-Fcgr3RS) or Myc-tagged wild type Fcgr3 (Myc-Fcgr3) or Myc-tagged Fcgr3-ΔG. The lentiviral construct expressing GFP alone (without any transgene) was used as control (pCSGW). The cells were then differentiated into BM-derived macrophages by culturing in DMEM containing L929, and Fc receptor-mediated phagocytosis was performed. **, p < 0.01; *, p < 0.05; ns, not significant when compared with pCSGW. D and E, LPS-induced TNFα (D) and IL-10 (E) levels were assessed by sandwich ELISA after cross-linking stably transduced U937 macrophages by anti-Myc mAb and F(ab’). Following cross-linking, the cells were washed, and supernatants were collected after 5 h of incubation with LPS (1 μg/ml). TNFα and IL-10 production were expressed as a percentage of that produced in LPS-stimulated cells without FcR cross-linking (100% of cytokine production). **, p < 0.001 when compared with LPS-stimulated cells without FcR cross-linking (100% of cytokine production; histogram in gray).
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(9). However, the COS7 cells are not primarily phagocytic cells and do not endogenously express the γ chain, making it difficult to draw mechanistic conclusions concerning the role of Fcgr3-rs in Fc receptor-mediated signaling and activation. Here we have examined the signaling function of Fcgr3-rs in a human monocyte cell line U937, a well characterized cell line previously used to study Fc receptor function (11). U937 cells constitutively express only Fcgr1 and Fcgr2a but not Fcgr3 (11). We have used lentivirus-mediated gene transfer to generate stably transduced U937 cell lines. Our results showed that Fcgr3-rs associates with the common γ chain in these cells, but Fc receptor-mediated signaling and phagocytosis are defective. Importantly, WKY BMDMs transduced with the Fcgr3-rs also showed reduced bead phagocytosis, and this was mediated through the rat-specific novel cytoplasmic domain.

It is well established that the transmembrane domain is required for association with the γ chain and consequently cell surface expression of Fcgr3 (24). Because of the amino acid changes between Fcgr3 and Fcgr3-rs in the transmembrane domains, it could therefore be argued that the amino acid changes simply prevented Fcgr3-rs expression. This, however, was not the case, and in both U937 cells and primary rat macrophages, Fcgr3-rs was expressed at the cell surface at similar levels to Fcgr3. The use of the ΔG chimeric molecule, which contains the extracellular and transmembrane domains of Fcgr3 combined with the cytoplasmic domain of Fcgr3-rs, also demonstrates that the differences seen between Fcgr3 and Fcgr3-rs functions are mediated only by the novel cytoplasmic domain of Fcgr3-rs and cannot be explained by point mutations in the extracellular domain of Fcgr3-rs, which might simply have prevented Fc binding of IgG.

The majority of signaling cascades emanating from the Fcgr3 receptor complex are thought to be derived from the γ chain via its ITAM motifs. Thus, removal of the γ chain cytoplasmic domain or mutations of tyrosine residues within the ITAM motifs affects intracellular Ca²⁺ levels, PI3K activity and receptor-induced phosphorylation events (25). In particular there is compelling evidence that activation of Syk is essential for subsequent events such as receptor-induced phagocytosis (26, 27). However, the cytoplasmic tail of the Fcgr3 molecule is unlikely to be completely redundant in this process, and Hou et al. (28) have suggested that sequences within the cytoplasmic domain of Fcgr3, in particular those close to the transmembrane domain, are required for signaling. Our findings demonstrate that signaling from the Fcgr3-rs is clearly different from that mediated by Fcgr3, because Fcgr3-rs γ chain complexes do not generate phosphorylated Syk molecules or activate the NF-κB transcription factor that is required for transcription of Fc-responsive genes such as TNFα. Accordingly, IgG-induced TNFα production and phagocytosis are both significantly lower in U937 cells expressing Fcgr3-rs compared with Fcgr3. Moreover, expressing Fcgr3-rs in primary rat WKY BMDMs resulted in lower levels of phagocytosis than cells expressing Fcgr3 alone, suggesting that Fcgr3-rs has an inhibitory effect on Fcgr3-mediated phagocytosis in these cells.

Although the Fcgr3-rs cytoplasmic domain includes a novel sequence with a single tyrosine molecule, this is not an immunoreceptor tyrosine-based inhibitory motif in the same way as that found in the FcγRIIb receptor. Elegant studies by Kim et al. (29) have shown that specific regions of the transmembrane region of Fcγ receptors are important for signaling leading to phagocytosis. The same group has provided further evidence on differences in the signaling properties of FcγR1/γ and FcγRIIA such as their interaction with Syk and Src-related tyrosine kinases (30). The region of the cytoplasmic domain shown by Hou et al. (28) to be important in signaling is present in the Fcgr3-rs; however, it is possible that the altered sequence of the remainder of the cytoplasmic domain may render this region inaccessible to essential downstream signaling molecules, thereby rendering it inactive. The question of whether the Fcgr3-rs exerts its effects by acting as a decoy receptor, binding immunoglobulins, but not signaling, or whether its novel cytoplasmic domain induces distinct signaling pathways that act to inhibit those from the wt Fcgr3 is therefore an interesting one.

Our demonstration herein that the -rs does not induce Syk or NF-κB activation suggests that this molecule may not signal, and this is underscored by the demonstration that LPS-induced TNF and IL-10 production is unaffected in U937 cells by the presence of the -rs. Taken together, these data add further weight to the argument that Fcgr3-rs is a decoy receptor that is able to bind Fc but unable to signal such that its expression inhibits co-expressed Fc receptor-mediated signaling and cell activation. However, we cannot completely rule out a potential additional role of this receptor, which will lead to inhibition by a novel negative signaling pathway. Investigations into this question are continuing in this laboratory.

In conclusion, our results suggest that the rat-specific Fcgr3-rs acts to inhibit Fcγ-mediated signaling and phagocytosis in both rodent and human cells and could be considered as a novel therapeutic mechanism for the modulation of Fc receptor-mediated cell activation in macrophage-mediated diseases such as crescentic glomerulonephritis.

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