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Plasma levels of the soluble form of the FcyRIIa receptor vary with receptor

polymorphisms and are elevated in rheumatoid arthritis

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Running title: Soluble FcyRIIa levels and polymorphisms

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Abstract

Soluble forms of the low-affinity immunoglobulin receptor FcyRIIa (sFcyRIIa) lacking the cytoplasmic tail have been reported in plasma however the mechanism and functional consequences are unknown. This study aimed to evaluate mechanisms of FcyRIIa release compared to GPVI release from platelets, and examine whether genetic polymorphisms at positions 27 and 131 within FcyRIIa correlate with platelet FcyRIIa stability and function. Enzyme-linked immunosorbent assays (ELISAs) were used to measure plasma sFcyRIIa and sGPVI levels. FcyRIIa genotype at positions 27 and 131 was evaluated. sFcyRIIa levels were not significantly different between non-131HH and 131HH, but were significantly lower in 27W/131H than non-27W/131H. Treatment of platelets with aggregated immunoglobulin (Ig) G induced release of FcyRIIa and GPVI, but only sGPVI release was statistically significant, required functional FcyRIIa, and was blocked by inhibitors of signalling pathways and metalloproteinases. This indicated that sFcyRIIa was not released from platelets by metalloproteolysis. sFcyRIIa levels were not correlated with sGPVI levels in healthy individuals however levels of sFcyRIIa and sGPVI in plasma from patients with rheumatoid arthritis (RA) were significantly elevated above levels found in healthy individuals. Elevated level of sFcyRIIa in RA patients may reflect active immune-based arthritis and be predictive of active inflammation.

Key words: FcyRIIa; polymorphism; platelet; GPVI; rheumatoid arthritis

Introduction

FcyRIIa is the only IgG Fc-receptor expressed on the surface of platelets but is also found on neutrophils, monocytes, and macrophages [1, 2]. It is a low affinity receptor for Ig unique to humans and non-human primates, and composed of two extracellular Ig-like domains, containing the binding site for the Fc portion of IgG, a transmembrane domain and a cytoplasmic tail that contains a binding site for calmodulin and an immunoreceptor tyrosinebased activation motif (ITAM) domain that is essential for signal transduction and platelet activation. Both alleles of the gene encoding FcyRIIa are transcribed and translated, generating equal amounts of protein. A polymorphism, resulting from the amino acid substitution from arginine (R) to histidine (H) at position 131 in the second Ig-like domain of FcyRIIa has been previously shown to alter the binding affinity of the receptor for different IgG subclasses. FcyRIIa-H131 binds to human IgG2 with higher affinity than FcyRIIa-R131; however, FcyRIIa-R131 has a higher binding affinity for murine IgG1 than FcyRIIa-H131 [3-5]. Increasing evidence [6-9] demonstrated that the FcyRIIa polymorphism (H/R131) was associated with immune diseases due to different IgG binding affinity. A further polymorphism in FcyRIIa at position 27 (glutamine or tryptophan) was shown not to be linked with the polymorphism at position 131 and in isolation did not affect receptor function or IgG binding [10] but may have some impact on FcyRIIa signalling events [11].

Previous studies *in vitro* demonstrated that engagement of either FcγRIIa or GPVI on platelets by their respective ligands induced both intracellular calpain-mediated cleavage of FcγRIIa and extracellular metalloproteinase-mediated ectodomain shedding of GPVI [12]. Increased levels of sGPVI are also observed in patients with immune thrombocytopenia [13, 14] and in patients with seropositive rheumatoid arthritis [15]. It is postulated that increased release of sGPVI could be induced by engagement of platelet receptor FcγRIIa by the Fc Page 5 of 27

Platelets

portion of anti-platelet autoantibodies in the plasma of ITP patients. The cytoplasmic domain of Fc γ RIIa binds calmodulin [12], a hallmark of many metalloproteolytically shed membrane receptors including GPVI [16], GPV [17] and L-selectin [18]. Two Fc γ RIIa transcripts (with or without the transmembrane domain encoding exon, termed Fc γ RIIa1 and Fc γ RIIa2 respectively), resulting from alternative RNA splicing, have been found in human platelets and a megakaryocytic cell line [19]. A third transcript, Fc γ RIIa3 with a 19-amino acid cytoplasmic insertion was identified in peripheral blood mononuclear cells [20]. Further, there are other transcript variants (X1 to X10) which are predicted by automated computational analysis. Soluble forms of Fc γ RIIa (Fc γ RIIa2) reported in serum samples from healthy individuals and patients with chronic lymphocytic leukaemia are released from thrombin-activated human platelets [21] and megakaryocyte cell lines [22]. Activated neutrophils from systemic lupus erythematosus patients demonstrated a serine proteinase-dependent cleavage of Fc γ RIIa that was correlated with markers of disease severity and complement activation [23]. However, the mechanism by which Fc γ RIIa is released from blood cells and whether ligand-induced release of Fc γ RIIa similar to that observed with GPVI shedding occurs *in vivo* has not been addressed.

This study aimed to evaluate the proteolytic mechanisms of FcγRIIa release compared to GPVI release and examine whether genetic polymorphisms at positions 27 and 131 within FcγRIIa correlate with platelet FcγRIIa stability and function.

Materials and Methods

Reagents

N-ethylmaleimide (NEM), GM6001 (a broad-range hydroxamic acid-based metalloproteinase inhibitor), calmodulin inhibitor W7 (N-(6-aminohexyl)-5-chloro-1-aphthalenesulfonamide)

Platelets

and the membrane permeable calpain inhibitor E64d (L-3-carboxy-trans-2, 3-epoxypropionyl-L-leucylamido-(3-methyl) butane) were from Calbiochem (La Jolla, CA, USA). The hydroxamate-based inhibitor GI254023 has been previously described and characterized for preferential inhibition of ADAM10 over ADAM17 [24]. The Syk inhibitor, 2-(7-(3,4dimethoxyphenyl)-imidazo[1,2-c]pyrimidin-5-ylamino)-nicotinamide dihydrochloride (BAY61-3606) was from Enzo Life Sciences (Farmingdale, NY, USA).

Antibodies

Anti-FcyRIIa monoclonal antibody 8.26 recognising a non-ligand binding region and HRPconjugated anti-FcyRIIa monoclonal antibody 8.7 recognising a ligand binding region within domain 2 have been described [25]. 14A2, which binds to tetraspanin CD151 and engages FcyRIIa via its Fc domain, has been previously described [26, 27]. The anti-FcyRIIa monoclonal antibody, IV.3 [28] was purified from hybridoma medium on protein A-Sepharose Z.C (Amersham, United Kingdom).

Measurement of sFcyRIIa by ELISA

The ELISA method for measuring sFcyRIIa was based on an established sandwich ELISA used to measure sGPVI in plasma [29]. Briefly, 96-well microtitre plates were coated with a "capturing" mouse monoclonal anti-FcyRIIa IgG 8.26 (2 µg/ml) in coating buffer (0.05 M bicarbonate, pH 9.6) overnight at 4°C. Wells were washed six times with PBS (0.01 M sodium phosphate, 0.15 M sodium chloride, pH 7.4) containing 0.2% (v/v) Tween-20 (PBS-T), blocked with 1% (w/v) BSA in PBS for 1 h at room temperature, then washed six times with PBS-T, before addition of samples diluted 10% (v/v) in PBS in triplicate wells. After 1 h incubation at room temperature, wells were washed six times with PBS-T and the HRP-conjugated anti-FcyRIIa monoclonal antibody 8.7 was added at 1 µg/ml (100 µl/well). After further 1 h incubation, wells were washed six times, 100 µl of a one-half dilution of Super Signal ELISA

Platelets

Pico chemiluminescent substrate (Pierce Rockford IL, USA) was added and light emission (stable after 1 min) was measured using a Wallac-Victor2 luminescence plate reader.

Blood collection and platelet aggregation

Experiments were carried out with the approval of the Monash University Standing committee on Ethics in Research Involving Humans, and informed consent was obtained from all participants in accordance with the Declaration of Helsinki. Platelet-rich plasma (PRP) was obtained from blood collected into 3.2% (w/v) trisodium citrate and centrifuged for 20 min at 160 x g. Platelet-poor plasma (PPP) was obtained by centrifuging PRP at 1,350 x g for 15 min. Washed platelets were prepared as previously described [30]. Platelet aggregation in citrated PRP was carried out at 37°C in a ChronoLog lumiaggregometer (Havertown, PA, USA) stirred at 900 rpm using PPP as control. Donors were of broad ethnicity and included donors with Caucasian, Asian, and Middle Eastern heritage. The donors were not preselected and all experimental work was completed before the genotype of each sample was revealed.

DNA sequencing

Genomic DNA was sequenced using the BigDye Terminator v1.1 cycle kit (Applied Biosystems, CA, USA). The sequenced region of the FcyRIIa protein included the two extracellular Ig domains. The sequences of primer 1 for DNA sequencing (the first Ig domain) were as follows: Forward: 5'-CCCCCAAAGGCTGTGCTGAAACTTGAGC-3'; Reverse: 5'-TCATTGTTGTTGGCCTTGAACCTGTAG-3'. The sequences of primer 2 for DNA sequencing (the second Ig domain) were: Forward: 5'-CATCATGCTGAGGTGCCACAGC-3'; Reverse: 5'-GATGAGAACAGCGTGTAGCCTATGTTTC-3'. For FcyRIIa genotype displayed in this study, at amino acid position 27 (located within the first Ig domain), the polymorphic residues are Q: Glutamine, W: Tryptophan; at position 131 (within the second Ig

domain) the polymorphic residues are H: Histidine, R: Arginine. For example QW/HR represents heterozygosity at both positions 27 (Q/W) and 131 (H/R) polymorphic sites.

Analysis of sFcyRIIa in immune-related disease

To compare levels of sFcyRIIa and sGPVI in plasma from individuals with circulating immune complexes, citrated plasma samples were obtained from consecutively recruited patients with an established diagnosis of RA, or healthy donors. The samples were not matched for age or gender. These samples were obtained with institutional ethics approval at the Mater Misericordiae University Hospital, Dublin, Ireland. People with a history of cardiovascular disease or who were receiving anti-platelet therapy or thromboembolic prophylaxis were excluded.

Statistical analysis

GraphPad Prism software (version 6.0) was used to apply a two-tailed student's unpaired t-test to compare sFc γ RIIa or sGPVI levels between different groups of known Fc γ RIIa genotype and a two-tailed Pearson rank was used to analyse correlations between sFc γ RIIa and sGPVI. P < 0.05 was considered to be statistically significant. ***P < 0.001; **P < 0.01; *P < 0.05; ns: not significant.

Results

sFcyRIIa measurement in plasma by ELISA

Evidence in the literature indicates that a soluble form of FcyRIIa (sFcyRIIa) exists in plasma [21]. To measure levels of sFcyRIIa in human plasma of healthy donors or patients with immune-related disease, an ELISA was established using antibodies against separate epitopes within the ectodomain. The ELISA output was reported in relative units, since purified or

Page 9 of 27

Platelets

recombinant sFcyRIIa proteins of known concentration were not available to enable calibrating the signal to a standard concentration curve. Serial dilution of sample produced a linear concentration-dependent signal. Serial dilution (from 1 in 2 to 1 in 64) of samples from either normal plasma or supernatant from washed platelets showed a linear relationship with relative light emission (Relative Unit, RLU) (Figure 1), with linear regression analysis for plasma (r^2 = 0.998) and supernatant sample (r^2 = 0.984) within an acceptable limit. sFcyRIIa levels in plasma from 24 healthy individuals was 0.743 ± 0.048 x 10⁶ RLU (Mean ± SE) (data not shown), with a normal range from 0.11 to 1.19 x 10⁶ RLU. The ELISA had a coefficient of variation of 3.8% and inter-assay variability of less than 10% indicating acceptable precision. Linear regression equations provided the best fit for the data.

Effect of FcyRIIa polymorphisms on plasma sFcyRIIa levels

In 24 healthy individuals with known genotype of FcγRIIa, the plasma sFcγRIIa level appeared to be dependent upon genotype (Figure 2A), however as three of the genotypes contained data from only one individual, conclusions from this cohort were limited [31]. Of note, the allele frequency of the W27 variant in this cohort was 0.3125, which was significantly higher than the minor allele frequency reported in global genetics reference databases (MAF 0.06 to 0.1) [32]. The reason for this was not clear but may be related to the heterogeneous ethnicity of the donor group. To improve the statistical power, in subsequent analyses, we grouped data based on occurrence of HH at position 131. Previous studies demonstrated that the amino acid at position 131 influenced binding affinity of FcγRIIa for IgG complexes [10]. FcγRIIa with a histidine (H) has a higher binding affinity to human IgG2 than FcγRIIa with an arginine (R) [4, 5]. However, in this study, the sFcγRIIa level was not significantly different in non-131H and 131H/H (Figure 2B), but was significantly lower in 27W than 27Q (Figure 2C), indicating that

sFcγRIIa release may be distinct from ligand binding and that FcγRIIa polymorphisms at position 27 and 131 might influence release of sFcγRIIa.

The mechanism of FcyRIIa release from platelets

Soluble forms of FcyRIIa have been detected in human serum or plasma [21-23, 33, 34]. Platelets express FcyRIIa and are a potential source of plasma sFcyRIIa by either secretion of a soluble form of the receptor lacking the transmembrane domain, or proteolytic shedding of the ectodomain from the platelet surface. Previous studies demonstrated that sGPVI is generated from platelet GPVI by metalloproteinase-mediated ectodomain shedding [16] and that engagement of platelet FcyRIIa also triggered shedding of GPVI [12]. GPVI and FcyRIIa are structurally analogous members of the immunoreceptor family, and both receptors bind calmodulin within the cytoplasmic domain, a common feature of metalloproteinase-dependent shedding. To test the hypothesis that engagement of FcyRIIa, induces metalloproteinasemediated ectodomain shedding and releases soluble fragment, human washed platelets were treated with aggregated-IgG (A-IgG) in the presence or absence of metalloproteinase inhibitor EDTA. Levels of sFcyRIIa or sGPVI were subsequently measured by ELISA. Treatment of washed platelets with A-IgG induced release of sFcyRIIa (Figure 3A), and sGPVI (Figure 3B) with a trend towards dose-dependency. This was consistent with previous studies showing engagement of FcyRIIa or GPVI by their respective ligands triggers both calpain-mediated intracellular cleavage of FcyRIIa and metalloproteinase-mediated ectodomain shedding of GPVI [12]. The release of sGPVI but not sFcyRIIa was statistically significant. Further, blockade of FcyRIIa by inclusion of IV.3 monoclonal antibody or inclusion of inhibitors of metalloproteinases, calpain or signaling molecules prevented GPVI shedding (Figure 3B), but did not inhibit the generation of sFcyRIIa. This strongly suggested that the mechanism of sFcyRIIa release was different from GPVI shedding (metalloproteinase-mediated). Taken

Platelets

together, these data demonstrate that the presence of sFcγRIIa could possibly be driven by a mechanism involving one or more unidentified sheddases, but not by an EDTA-inhibitable metalloproteinase.

FcyRIIa polymorphisms, FcyRIIa release and GPVI shedding

In addition to sharing similar structures, both FcyRIIa and GPVI/FcRy contain an immunoreceptor tyrosine-based activation motif (ITAM) within the cytoplasmic tail involved in signaling transduction. In human platelets treated with agonists acting at either FcyRIIa or GPVI in vitro, there is both intracellular calpain-mediated cleavage of FcyRIIa (removing the ITAM signaling sequence) and extracellular metalloproteinase-mediated ectodomain shedding of GPVI [12]. In order to examine the relationship between generation of sFcyRIIa and sGPVI *in vivo*, the respective ELISA assays were used to measure $sFc\gamma RIIa$ and sGPVI in the same plasma samples obtained from healthy individuals. The observed lack of correlation between sFcyRIIa and sGPVI (Figure 4A) is not only consistent with the lack of effect of FcyRIIa polymorphisms on GPVI shedding (Figure 4B), but also suggests that sFcyRIIa could be derived from other vascular cells, since FcyRIIa is also expressed and shed from neutrophils and monocytes/macrophages [23]. Our previous studies demonstrated that coagulation induces platelet GPVI shedding as sGPVI level was markedly elevated in serum. To assess whether coagulation influences FcyRIIa release, blood was collected in different anti-coagulants or clotting tube and sFcyRIIa level was measured by ELISA with sGPVI level as a positive control. We showed that unlike elevated serum sGPVI level, sFcyRIIa level in serum is comparable with plasma level in citrate, acid citrate dextrose or EDTA anti-coagulants (data not shown), indicating that unlike GPVI, coagulation did not affect FcyRIIa release. All these results displayed above revealed that different mechanisms are likely to be involved in release of FcyRIIa, compared with GPVI release.

FcyRIIa polymorphisms, release and platelet aggregation

14A2, a murine monoclonal antibody against CD151, induces platelet activation and aggregation via engagement of FcyRIIa [27]. Platelet aggregation in response to 14A2 contributed to identifying functional differences associated with FcyRIIa polymorphisms [35]. In the present study, 14A2 at a threshold concentration (1.0 µg/ml,) was used to investigate platelet aggregation and polymorphisms of FcyRIIa. Based on the time taken to achieve 50% aggregation, platelet aggregation to 14A2 was divided into quartiles (+++, ++, +, ---). Among 24 healthy individuals with known genotype of FcyRIIa (Table 1), platelet aggregation in response to 14A2 was heterogeneous, consistent with published studies [26, 35]. Standard appropriately powered statistical evaluation approaches could not be used to evaluate genotype and propensity to aggregate, due to three genotypes containing data from a single individual. However, in healthy individuals with HH at position 131 of FcyRIIa, platelet aggregation was completely impaired regardless of the polymorphism at 27, possibly due to the lower binding affinity to murine IgG (14A2) for FcyRIIa with HH [10]. However, non-responsiveness to 14A2 was also found in individuals with HR, indicating that other factors, such as copy number or surface density of FcyRIIa, or genetic variation of the protein tyrosine phosphatase CD148 that regulates ITAM receptor-mediated platelet activation could also contribute to the function of FcyRIIa [36]. The plasma level of sFcyRIIa under resting conditions was also heterogeneous in individuals with different platelet responses to 14A2.

Analysis of sFcyRIIa in immune-related disease

There is increasing evidence showing the critical role of FcRs in auto-immune disease induced by immune complexes [1, 23, 37, 38], specifically the importance of activating FcγRIIa in the pathogenesis of RA [39]. Available plasma samples from RA patients, previously used to measure levels of sGPVI, provided an opportunity to apply the new ELISA to measure Page 13 of 27

Platelets

sFcγRIIa levels in the same samples. As seen in Figure 5A, the sFcγRIIa level in RA patients was significantly higher than that of healthy controls, indicating FcγRIIa may be a potential marker in the pathogenesis of RA, and consistent with previous studies [39]. Plasma sGPVI levels were also significantly higher in RA patients when compared with controls (Figure 5B), revealing platelets might also play a role in this disease. There was no correlation between sFcγRIIa and sGPVI in RA patients (Figure 5C), consistent with data obtained from healthy individuals. These results support the possibility that the release mechanisms for these two receptors are different, further supporting the *in vitro* studies of shedding mechanisms shown in Figure 3. Unfortunately, genotyping of the RA patients was not possible, and detailed clinical information on the progression and outcomes of disease in different individuals could not be obtained. Nevertheless, findings presented here provide pilot data which would justify expanded future studies of platelet activation and plasma sFcγRIIa analysis in RA using the ELISA and reference data described in this study.

Discussion

FcyRIIa is the only IgG-Fc receptor with low affinity expressed on platelets and megakaryocytes. Three RNA splice variants of the *FCGR2A* gene have been described with FcyRIIa1 the most widely studied. FcyRIIa2 is secreted or released from human epidermal Langerhans cells [40], neutrophils [23, 34] or thrombin-activated human platelets or a megakaryocytic cell line [22]. Subsequent studies demonstrated that FcyRIIa2 lacking a hydrophobic segment of the transmembrane exonic sequence was present in human serum [21]. FcyRIIa3, is identical to FcyRIIa1 with the addition of a 19-amino acid insertion in the juxtamembrane region of the cytoplasmic tail. However, to date the levels of sFcyRIIa present in normal plasma or in immune-related disease states have not been widely analysed in terms of

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Platelets

normal variability between individuals or how levels change in immune or other disease states. In this study, an ELISA was developed for quantifying relative levels of sFcγRIIa in human plasma, to analyse the mechanism of sFcγRIIa release from platelets, evaluate the effect of sFcγRIIa genotype on function, and increase understanding of sFcγRIIa as a biomarker and/or role in the pathology of immune-related platelet dysfunction.

Our previous in vitro study showed engagement of either GPVI or FcyRIIa on platelet by their respective ligands induced both extracellular metalloproteinase-mediated ectodomain shedding of GPVI and intracellular calpain-mediated proteolysis of FcyRIIa [12]. When platelets were treated with A-IgG, release of GPVI (sGPVI) and FcyRIIa (sFcyRIIa) was detected by ELISA in a dose-dependent manner. From our study, several lines of evidence suggest that the release of sFcyRIIa from human platelets is regulated by a different mechanism to that involved in shedding of GPVI. First, A-IgG-induced release of sFcyRIIa from platelets is not inhibited by metalloproteinase inhibitors or by inhibitors of signaling pathways involved in GPVI shedding, consistent with a recent study showing metalloproteinase inhibitor GM6001 and E64d did not affect FcyRIIa shedding from neutrophils after addition of TLR agonists. In this case, TLR agonist-induced FcyRIIa shedding was shown to be dependent on serine proteases, including the pro-protein convertase furin [23]. Second, coagulation induced pronounced shedding of GPVI [41], whereas plasma and serum levels of sFcyRIIa were comparable, consistent with a previous study showing serum and plasma sFcyRIIa levels were similar [21]. Third, there is no apparent correlation between levels of sGPVI and sFcyRIIa in normal healthy plasma, which might have been expected to correlate if common triggers or pathways were involved in regulating their release from platelets. Although in this regard, unlike sGPVI which is only derived from platelets, sFcyRIIa may be derived from platelets or other cell types.

Page 15 of 27

Platelets

Previous studies demonstrated that the FcγRIIa polymorphism (H131R) was associated with some autoimmune diseases due to different IgG binding affinity [6, 7, 9] and another polymorphism at position 27 (W/Q) did not affect receptor function or IgG binding and was not linked with the polymorphism at position 131 [10]. Regarding the plasma sFcγRIIa level in individuals with different genotype, we showed the sFcγRIIa level was not significantly different between H131H and non-H131H groups, but is significantly lower in the 27W group, indicating sFcγRIIa release may not depend on ligand binding. Our work indicates that the genotype at positions 27 and 131 influence receptor stability of FcγRIIa in healthy individuals. These findings provide a potential mechanistic basis for altered responsiveness of platelets to antiplatelet antibodies, and may provide a suitable rationale justifying future studies to evaluate the link between genotype and immune-based platelet defects.

Consistent with previous studies showing FcyRIIa-R131 binds to murine IgG with higher binding affinity than FcyRIIa-H131 [6, 7] and that the amino acid at position 27 did not affect receptor function [10], platelet aggregation response to the threshold concentration of 14A2, a murine monoclonal antibody against CD151, was completely impaired in individuals with H131H, regardless of the amino acid at polymorphic position 27. However, impaired platelet aggregation was also observed in some individuals with H131R, suggesting apart from polymorphism, other factors, such as surface expression of FcyRIIa, genetic variation of the protein tyrosine phosphatase CD148 could also be involved in FcyRIIa-dependent platelet aggregation [36].

Finally, an ELISA for measuring sFcγRIIa in human plasma should ultimately enable the levels of sFcγRIIa to be evaluated in diseases involving elevated immune complexes. In this study, preliminary analysis of sFcγRIIa in healthy donors and patients with arthritis has shown that plasma levels of sFcγRIIa were significantly higher in RA patients when compared

Platelets

with control (healthy donors), indicating Fc γ RIIa was involved in the pathogenesis of RA, consistent with previous studies. sFc γ RIIa level measured by ELISA may reflect active immune-based arthritis and be predictive of active inflammation. In addition, plasma sGPVI level was also elevated in patients with RA when compared with control, implying platelet and/or collagen receptor GPVI might also be involved in RA, consistent with inflammation amplified by platelet in arthritis via collagen/GPVI-dependent microparticle generation [42]. The elevated sGPVI level might be due to GPVI-dependent platelet activation induced by collagen binding in RA. Unfortunately there was no genomic information available for this RA cohort to evaluate the Fc γ RIIa polymorphisms. It is worth noting however that other studies have identified an association between the FCGR2A R allele and RA in Europeans (p=0.020), but not in East Asians populations [43], or in a Japanese cohort [44].

In conclusion, it appears that soluble $Fc\gamma RIIa$ release results from secretion, possibly dependent on ligand engagement of $Fc\gamma RIIa$, rather than ectodomain shedding, consistent with $Fc\gamma RIIa$ release from activated human platelets *in vitro*. Recombinant soluble $Fc\gamma RIIa$ has been previously shown to inhibit Fc-mediated human platelet aggregation in a dose-dependent manner [22]. Therefore, soluble $Fc\gamma RIIa$ released from platelets could be a down-regulation mechanism of platelet function activated *via* Fc binding in autoimmune disease.

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Author Contributions

JQ performed research and interpreted data; ED and BW developed key reagents and interpreted data; DK, GMM, PMH and KX provided key patient samples and reagents; RKA and EEG designed the research and interpreted the data; all authors co-wrote the manuscript.

Conflicts of Interest

All authors have no conflict of interest to declare.

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Figure legends

Figure 1. Measurement of sFcγRIIa by ELISA assay. Serial dilution of plasma (A) or supernatant (S/N) from washed platelets (B) was used to measure the soluble level of FcγRIIa by ELISA as described in Methods.

Figure 2. sFcyRIIa levels in healthy individuals with different FcyRIIa genotype. Plasma or genomic DNA from 24 healthy individuals was used to measure sFcyRIIa level by ELISA and FcyRIIa genotype by DNA sequencing. (A) Six different alleles of FcyRIIa were identified. Plasma from healthy individuals with (B) non-HH and HH genotype or (C) 27Q and 27W genotypes of FcyRIIa were used for sFcyRIIa measurement. Unpaired student t-test was used to compare sFcyRIIa level with different genotypes. **P<0.01; ns: not significant.

Figure 3. sFcyRIIa and sGPVI release from platelets. Washed platelets from healthy donors were treated with A-IgG in the presence or absence of EDTA (10 mM), IV.3 (10 μ g/ml), GM6001 (100 μ M), BAY-61 (5 μ M), E64d (10 μ M) (all final concentrations) for 3 h at room temperature and supernatant was isolated to measure soluble forms of (A) FcyRIIa and (B) GPVI by ELISA. The bars represent mean \pm SEM for triplicate data points. Significant differences from the untreated sample were evaluated using 1-way ANOVA with Dunnett's multiple comparisons test. *p<0.05; **p<0.01; ***p<0.005.

Figure 4. sGPVI and sFcyRIIa level in healthy individuals. (A) Plasma obtained from 27 healthy individuals was used to evaluate any relationship between sGPVI and sFcyRIIa levels by ELISA. Plasma sGPVI level in healthy individuals with (B) non-HH and HH genotype or (C) non 27W/131H and 27W/131H genotype of FcyRIIa. Unpaired student t-test was used to compare sFcyRIIa level with different genotypes. ns: not significant.

Figure 5. Levels of sFcyRIIa and sGPVI in plasma from RA patients or healthy donors.

Plasma from RA patients or healthy donors (control) was used to measure (A) sFcyRIIa or (B) sGPVI by ELISA. (C) A correlative analysis of sFcyRIIa and sGPVI in RA patients. ***p < 0.001; **p < 0.01.

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Platelets

Table 1 Summary of FcyRIIa genotypes in healthy individuals. FcyRIIa genotype at position 27 and 131, platelet aggregation in response to 14A2 and sFcyRIIa levels in plasma from 24 healthy donors. The ranges of times taken to achieve 50% aggregation are < 2 min (+++), 2-3min (++) and \geq 3min (+) respectively. --- indicated no platelet aggregation in response to 1 µg/ml 14A2 after 20 min. All data were obtained using PRP and were reproducible on different days (n=2-3).

	FcyRIIa	genotype	<u> </u>	Aggregation (1.0 μg/ml 14A2)				Plasma soluble
n	27	131		+++	++	+		(RLU x 10 ⁶) (range)
3	QQ	HH	7				3	0.89-1.19
1	QW	HH					1	0.61
1	WW	HH					1	0.31
12	QQ	HR		1	4	2	5	0.49-1.19
6	QW	HR			4	2		0.32-0.80
1	QQ	RR			1			0.11
							2,	







Platelets



Figure 3





