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## Distinct Fc $\gamma$ receptors mediate the effect of Serum Amyloid P on neutrophil adhesion and fibrocyte differentiation

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### Abstract

The plasma protein Serum Amyloid P (SAP) reduces neutrophil adhesion, inhibits the differentiation of monocytes into fibroblast-like cells called fibrocytes, and promotes phagocytosis of cell debris by macrophages. Together, these effects of SAP reduce key aspects of inflammation and fibrosis, and SAP injections improve lung function in pulmonary fibrosis patients. SAP functions are mediated in part by Fc $\gamma$  receptors, but the contribution of each Fc $\gamma$  receptor is not fully understood. We found that amino acids Q55 and E126 in human SAP affect human fibrocyte differentiation and SAP binding to Fc $\gamma$ RI. E126, K130 and Q128 affect neutrophil adhesion and SAP affinity for Fc $\gamma$ RIIa. Q128 also affects phagocytosis by macrophages and SAP affinity for Fc $\gamma$ RI. All the identified functionally significant amino acids in SAP form a binding site that is distinct from the previously described SAP-Fc $\gamma$ RIIa binding site. Blocking Fc $\gamma$ RI with an IgG blocking antibody reduces the SAP effect on fibrocyte differentiation, and ligating Fc $\gamma$ RIIa with antibodies reduces neutrophil adhesion. Together, these results suggest that SAP binds to Fc $\gamma$ RI on monocytes to inhibit fibrocyte differentiation, and binds to Fc $\gamma$ RIIa on neutrophils to reduce neutrophil adhesion.

### Introduction

Aberrant scar tissue formation is the hallmark of fibrosing diseases such as end-stage kidney disease, liver cirrhosis, pulmonary fibrosis, and congestive heart disease (1–3). The inappropriate scar tissue in fibrosis ultimately leads to organ failure and/or death. Fibrosing diseases are associated with 45% of deaths in the US, but despite their high prevalence, there are no FDA-approved therapies (1, 4).

Serum Amyloid P component (SAP) is a pentameric protein that belongs to the pentraxin family of evolutionarily conserved proteins. Pentraxins also include C-reactive protein (CRP) and the long pentraxin PTX-3 (5). SAP, CRP, and PTX-3 all have regulatory roles in the immune system (6–8). Injections of SAP inhibit inflammation and fibrosis in mouse models of pulmonary fibrosis, ischemic cardiac fibrosis, and renal fibrosis (9–12), and in a phase 1b clinical trial, SAP injections appear to improve lung function in pulmonary fibrosis patients (13).

At the onset of tissue damage and inflammation, neutrophils are recruited to the tissue in response to chemokines such as CXCL2 and CXCL8 to remove pathogens and/or cell debris via phagocytosis (14). This migration and activation of neutrophils is tightly regulated by factors expressed and secreted by endothelial cells, macrophages and other cell types (14). When this regulation is compromised, the elevated influx of neutrophils and recruitment of other immune cells by activated neutrophils can cause severe organ damage and fibrosis (14–16). SAP binds neutrophils to inhibit their spreading and adhesion to components of extracellular matrix and endothelial cells (12, 17). Injections of SAP decrease the infiltration of neutrophils into the lungs following bleomycin insult in mice (12). However, the mechanism for this function is not well understood.

Following neutrophil migration into the inflammation site, CD14<sup>+</sup> monocytes enter and differentiate into macrophages and fibrocytes (3). Fibrocytes are CD45<sup>+</sup> collagen I<sup>+</sup> fibroblast-like cells that share characteristics of both hematopoietic and stromal cells (18). Fibrocytes are found in healing dermal wounds and some fibrotic lesions, and secrete collagen and enzymes which modify the extracellular matrix (3, 9, 10, 19, 20). SAP inhibits fibrocyte differentiation partly through a group of receptors called Fc $\gamma$  receptors (11, 21–24). These receptors bind IgG and consist of Fc $\gamma$ RI, Fc $\gamma$ RIIa, Fc $\gamma$ RIIb, Fc $\gamma$ RIIIa, and Fc $\gamma$ RIIIb (25). We have previously shown that Fc $\gamma$ RI is one of the receptors responsible for the effect of SAP on fibrocyte differentiation in both humans and mice (21). SAP also binds the IgA receptor Fc $\alpha$ RI (26).

In addition to modifying neutrophil adhesion and monocyte differentiation, SAP can also enhance phagocytosis of cell debris by professional phagocytes such as macrophages (24, 27). The SAP pentamer forms a flat disk, and binds to bacteria and cell debris on one surface, and to Fc $\gamma$  receptors on the other surface, to promote phagocytosis by cells (24). Previous studies have implicated Fc $\gamma$ RI as the key receptor for SAP-induced phagocytosis, but the precise role of each Fc $\gamma$  receptor in this process is unclear (24, 27).

In this report, we examined how SAP interacts with Fc $\gamma$  receptors to regulate different aspects of the immune system. We found that SAP inhibits fibrocyte differentiation and promotes phagocytosis by macrophages through Fc $\gamma$ RI, while it reduces neutrophil adhesion via Fc $\gamma$ RIIa. Using site-directed mutagenesis we determined that although the same site on SAP affects monocytes, macrophages, and neutrophils, it is possible to affect specific SAP functions without altering the other functions in an appreciable way. In addition, we identified a novel Fc $\gamma$  receptor binding site that is distinct from the site previously identified in a co-crystal structure of SAP and Fc $\gamma$ RIIa (23).

## Materials and methods

### PBMC and neutrophil isolation, cell culture, fibrocyte and macrophage differentiation

Human peripheral blood was collected into heparin tubes (BD Bioscience, San Jose, CA) from healthy adult volunteers who gave written consent and with specific approval from the Texas A&M University human subjects Institutional Review Board. Peripheral blood mononuclear cells (PBMC) were isolated from the blood using Ficoll-Paque Plus (GE Healthcare Biosciences, Piscataway, NJ), as described previously (28). PBMCs were

cultured in Fibrolife (LifeLine Cell Technology, Walkersville, MD) defined serum-free medium (SFM) in the presence or absence of SAP variants as previously described (21). To determine the contribution of each Fc $\gamma$  receptor to the SAP effect on fibrocyte differentiation, we incubated PBMCs with 5  $\mu$ g/ml of F(ab')<sub>2</sub> fragments of either anti-Fc $\gamma$ RI antibody clone 10.1 (mouse IgG1, Ancell, Bayport, MN), anti-Fc $\gamma$ RII antibody clone 7.3 (mouse IgG1, Ancell), anti-Fc $\gamma$ RIII antibody clone 3G8 (mouse IgG1, Ancell), or a mouse IgG1 isotype control (Ancell) in the presence and absence of SAP. Fibrocytes were identified and counted based on their elongated spindle-shaped morphology in five different 900  $\mu$ m-diameter fields of view per well (28–30). PBMCs were incubated overnight in RPMI-1640 (Lonza, Allendale, NJ) with 10% fetal bovine serum (Caisson Laboratories, North Logan, UT) to generate macrophages, as described previously (24). Neutrophils were isolated from blood using Lympholyte-poly (Cedarlane Laboratories, Hornby, BC, Canada) following the manufacturer's protocol and resuspended in 2% BSA (fraction V, A3059; Sigma-Aldrich) in RPMI-1640 (12, 31). HEK293 cells (Life Technologies, Grand Island, NY) were cultured in Freestyle (Life Technologies) medium following the manufacturer's protocol. K562 cells (ATCC, Manassas, VA) were grown in RPMI-1640 with 10% fetal bovine serum (Caisson).

### **SAP variant expression, purification, size exclusion chromatography, and labeling**

Starting with the previously described SAP expression vector (21), SAP variants were generated with a QuikChange II Site-Directed Mutagenesis Kit (Agilent Technologies, Santa Clara, CA) following the manufacturer's protocol, and the DNA sequences of the constructs were verified. SAP variants were expressed in HEK293 cells as described previously and then purified by affinity purification (21). Briefly, cell supernatants from the SAP-expressing HEK293 cells were clarified by centrifugation at 300  $\times$  g. 1 M CaCl<sub>2</sub> was added to the supernatant to a final concentration of 2 mM, and the cell supernatant was then mixed with 1 ml of a 50% slurry of Sepharose Fast Flow (GE Healthcare BioSciences, Piscataway, NJ, USA) in 20 mM Tris, 140 mM NaCl, 2 mM CaCl<sub>2</sub>, pH 7.4 for 1 h. The Sepharose beads were collected and washed 3 times with 15 ml wash buffer (20 mM Tris, 300 mM NaCl, 2 mM CaCl<sub>2</sub>, pH 7.4). Bound protein was eluted overnight at 4°C with 20 mM Tris, 140 mM NaCl, 50 mM EDTA, pH 7.4. The eluted protein was then buffer exchanged into 20 mM sodium phosphate buffer pH 7.4 (21). The purified SAP was assayed by size exclusion chromatography using a Superose 12 (GE Healthcare Life Sciences, Piscataway, NJ) column on an AKTA chromatography system (GE Healthcare Life Sciences) as previously described (21). Purified SAP was labeled using Alexa Fluor 647-NHS (Life Technologies) following the manufacturer's protocol.

### **Neutrophil adhesion assay, macrophage phagocytosis assay, and SAP binding to Zymosan A**

Neutrophils were incubated with 80 nM (10  $\mu$ g/ml) of wildtype (WT) or mutated SAP, and their binding to human plasma fibronectin (Trevigen, Gaithersburg, MD) was assessed as previously described (12). To determine the effect of Fc $\gamma$  receptor ligation on neutrophil adhesion, neutrophils were incubated with 5  $\mu$ g/ml of either anti-Fc $\gamma$ RI antibody clone 10.1 (mouse IgG1, eBiosciences), anti-Fc $\gamma$ RII antibody clone Clkm-5 (mouse IgG1, Millipore, Billerica, MA), anti-Fc $\gamma$ RII antibody clone FUN-2 (mouse IgG1, Biologend), anti-Fc $\gamma$ RII

antibody clone AT10 (mouse IgG1, Abcam, Cambridge, MA), anti-Fc $\gamma$ RIII antibody clone 3G8 (mouse IgG1, Biolegend), or a mouse IgG1 isotype control (Biolegend). Phagocytosis of FITC-conjugated Zymosan A bio-particles (Life Technologies) by macrophages was assayed as described previously (24). To measure the binding of SAP to Zymosan A bio-particles, we first quenched the fluorescence of FITC-conjugated Zymosan A with 2% Trypan blue in PBS for 20 minutes and then incubated the quenched Zymosan A bio-particles with 240 nM (30  $\mu$ g/ml) of WT SAP or mutant SAP in 20 mM Tris, 140 mM NaCl, 2 mM CaCl<sub>2</sub> for 1 hour. The bio-particles were then washed, and bound SAP was detected by staining with anti-SAP antibody clone 5.4A (Millipore) and goat anti-mouse Alexa Fluor-647 (Life Technologies) on an Accuri C6 flow cytometer (BD Bioscience).

### SAP affinity assays and receptor expression

Fc $\gamma$ RI and Fc $\epsilon$  common  $\gamma$ -chain (Fc $\gamma$ ) cDNA (PSI:Biological-materials repository, Tempe, AZ) (32, 33) were ligated into the pCMV6-AC-His vector (OriGene, Rockville, MD) and then transfected into HEK293 cells using jetPRIME (Polyplus, New York, NY) following the manufacturer's protocol. Fc $\gamma$ RIIIb plasmid was obtained from the PSI:Biological-materials repository and transfected into HEK293 cells. K562 cells, which express Fc $\gamma$ RIIa, were used to measure the affinity of SAP for Fc $\gamma$ RIIa (34). The binding of fluorescently labeled WT SAP or mutant SAP to cells was then measured as previously described using an Accuri C6 flow cytometer (24). When measuring SAP binding to HEK293 cells expressing Fc $\gamma$ RI or Fc $\gamma$ RIIIb, mock transfected HEK293 cells were used to estimate the non-specific binding. K562, HEK293, Fc $\gamma$ RI<sup>+</sup> HEK293, and Fc $\gamma$ RIIa<sup>+</sup> HEK293 cells were stained for Fc $\gamma$ RI (Clone 10.1, eBiosciences), Fc $\gamma$ RII (Clone FUN-2, Biolegend), and Fc $\gamma$ RIII (Clone 3G8, Biolegend) to determine the expression of the indicated receptor by flow cytometry (29). Leukocytes stained for CD3 (Biolegend), CD14 (Biolegend), CD15 (Biolegend), CD19 (Biolegend), CD45 (Biolegend), Fc $\gamma$ RI (Clone 10.1, eBiosciences), Fc $\gamma$ RII (Clone FUN-2, Biolegend), and Fc $\gamma$ RIII (Clone 3G8, Biolegend) were assayed by flow cytometry to determine the presence of different immune cell populations as previously described (29, 30, 35).

### Statistical analysis

Data was analyzed by ANOVA (with Dunnett's post test) or t-test when appropriate using Prism software (GraphPad software, San Diego, CA). Data were fit to the appropriate model of binding as determined by F-tests. Normality was tested using Shapiro-Wilk and D'Agostino-Pearson omnibus tests when applicable.

## Results

### Identification of SAP amino acids that affect neutrophil adhesion

We previously made site-directed mutations of human SAP at amino acids that interact with human Fc $\gamma$ RIIa in a SAP-Fc $\gamma$ RIIa co-crystal structure (23), and observed that changes to these amino acids had no significant effect on the ability of SAP to inhibit fibrocyte differentiation (21). To better understand the interaction of SAP with Fc $\gamma$  receptors, we compared the amino acid sequence of human SAP to the related pentraxin CRP. SAP and CRP have 51% sequence identity and similar crystal structures, but have different affinities

for Fc $\gamma$  receptors and different roles in the immune system (6, 11, 23, 27). Thus, the sequence differences can be used to identify structurally and functionally significant amino acids. Of the amino acids that were different between SAP and CRP, we mutated only the ones that were exposed on the surface of SAP (23). E153, which is at the interface between SAP monomers, was also mutated in an attempt to destabilize the pentameric protein and introduce functional defects. We then expressed all the generated SAP variants in HEK293 cells. All the SAP variants eluted at 10–12 ml from a Superose 12 size exclusion chromatography column, indicating the absence of aggregates larger than pentamers, and the absence of free monomers (Table SI and Fig 1A). We subsequently tested the ability of these variants to decrease neutrophil adhesion, inhibit fibrocyte differentiation, and promote phagocytosis. In addition, we examined the binding of these variants to Fc $\gamma$  receptors.

We first examined the ability of our SAP variants to reduce neutrophil adhesion to human fibronectin. We screened 29 SAP variants for their ability to reduce neutrophil adhesion and then based on our preliminary data focused on 13 of the examined variants (data not shown). These 13 SAP variants were screened at 80 nM (10 $\mu$ g/mL) (Fig. 1B). We chose this concentration because it was close to the IC<sub>50</sub> (67  $\pm$  7 nM) of SAP for reducing neutrophil adhesion (Fig. 1C) and hence allowed us to detect both increases and decreases in the SAP effect on neutrophils. Following our screen, we observed that SAP variants E126A and Q128A had increased inhibitory effect on neutrophil adhesion compared to WT SAP (Fig. 1B). SAP variant K130V conversely had a significantly reduced inhibitory effect on neutrophil adhesion (Fig. 1B).

#### Identification of SAP amino acids that affect fibrocyte differentiation

Since SAP inhibits fibrocyte differentiation, we also screened the 29 SAP variants for their ability to inhibit the differentiation of monocytes into fibrocytes (Table SI) (21, 28, 29, 36). Using PBMCs from a variety of donors, we observed 1200 to 3100 fibrocytes per 10<sup>5</sup> PBMCs. Because of this variability, fibrocyte counts were normalized to the no-SAP control, as described previously (21, 28, 37). WT SAP inhibited fibrocyte differentiation with an IC<sub>50</sub> of 2.9  $\pm$  0.3 nM, similar to previously published data (21, 36). 25 out of the 29 variants tested did not significantly alter the ability of SAP to inhibit fibrocyte differentiation (Table SI). Compared to wildtype SAP, variants Q55A and K130V were more effective at inhibiting fibrocyte differentiation whereas variants E153A and E126A had reduced activity (Fig. 2 and Table SI). In addition, we observed significant changes in the Hill coefficient of SAP variants V68A and Q128A compared to WT SAP (Table SI). This change could be due to alterations in SAP variant binding to Fc $\gamma$  receptors and/or how these variants activate the receptors.

#### Identification of SAP amino acids that affect phagocytosis

SAP enhances phagocytosis of pathogens and cell debris through Fc $\gamma$  receptors (24, 27). However, the exact receptor and amino acids involved are unknown. We first screened the 29 SAP variants for their ability to enhance phagocytosis and then based on our preliminary data focused on 13 variants (data not shown). The 13 SAP variants were screened at 240 nM (30 $\mu$ g/mL, physiological concentration in the human plasma) for their ability to promote phagocytosis of Zymosan A bio-particles by macrophages (Fig 3A). 11 of the SAP variants

examined did not significantly alter the ability of SAP to enhance phagocytosis (Fig 3A). Compared to WT SAP, SAP variants Q128A and E153A had significantly reduced ability to promote phagocytosis by macrophages (Fig 3A). We then measured the binding of WT, Q128A and E153A SAP to Zymosan A bio-particles to determine if these SAP variants had deficiencies in binding the bio-particles. Compared to WT SAP, we found no statistically significant differences in the binding of Q128A or E153A to Zymosan A (Fig 3B). Together, this indicates that SAP variants Q128A and E153A have defects in binding and/or activating Fc $\gamma$  receptors to promote phagocytosis of Zymosan A.

### **SAP binds to endogenous Fc $\gamma$ RI and Fc $\gamma$ RIIa on immune cells**

Much of the work done on SAP binding to Fc $\gamma$  receptors has focused on SAP binding to recombinant Fc $\gamma$  receptors or to receptors expressed on non-human cells such as COS-7 and NIH-3T3 (11, 23, 24). This is problematic because the affinity of Fc $\gamma$  receptors for their ligands is sensitive to the receptor glycosylation state and the presence of intracellular signaling proteins (38–41). Therefore to identify functionally significant receptors in the SAP response, we examined the binding of SAP to endogenous Fc $\gamma$  receptors on human immune cells and to receptors expressed on the human-derived cell line HEK293. We fluorescently labeled WT SAP (SAP-f) and then measured the binding to different peripheral blood cell populations as identified by their flow characteristics and receptor expression (Fig. S1). We tested the activity of SAP-f on neutrophils, monocytes, and macrophages and observed no functional defects compared to unlabeled SAP (Fig. S2). When SAP-f was incubated with leukocytes, we observed no binding to the lymphocyte population (Fig 4A). Since B cells (~5 % of lymphocytes) express Fc $\gamma$ RIIb (25), and NK cells (~5–10% of lymphocytes) express Fc $\gamma$ RIIIa (25), this suggests that SAP does not bind to these receptors under our experimental conditions. However, SAP-f did bind to monocytes and neutrophils (Fig. 4A). Monocytes express Fc $\gamma$ RI, Fc $\gamma$ RIIa, and some Fc $\gamma$ RIIIa (Fig S1) (12, 30). This indicates that SAP could be binding to any or all of the Fc $\gamma$  receptors on monocytes. As NK cells express Fc $\gamma$ RIIIa, and we did not detect binding of SAP-f to NK cells, this suggests that SAP binds to Fc $\gamma$ RI and/or Fc $\gamma$ RIIa on monocytes. Neutrophils express Fc $\gamma$ RIIa and Fc $\gamma$ RIIIb (Fig S1) (12, 30). When we incubated SAP-f with Fc $\gamma$ RIIIb<sup>+</sup> HEK293 cells, we did not detect any appreciable binding although the receptor was functional as determined by human IgG binding (Fig. 4B and C and Fig. S3). This then suggests that SAP binds to Fc $\gamma$ RIIa on neutrophils. We observed more SAP binding to monocytes than to neutrophils, and we hypothesize that this is likely due to the presence of Fc $\gamma$ RI on monocytes (Fig. 4A). Together, our data indicates that SAP-f binds to endogenous Fc $\gamma$ RI and Fc $\gamma$ RIIa on monocytes and neutrophils but not to Fc $\gamma$ RIIb, Fc $\gamma$ RIIIa, and Fc $\gamma$ RIIIb (Fig. 4).

### **SAP binds to Fc $\gamma$ RI and Fc $\gamma$ RIIa on HEK293 cells**

Following our initial binding assays using human immune cells, we investigated the binding of our SAP variants to Fc $\gamma$ RI and Fc $\gamma$ RIIa. Of the 29 SAP variants, we chose 6 that had altered functions as measured by neutrophil adhesion assays, fibrocyte differentiation assays, and macrophage phagocytosis assays. The 6 SAP variants were fluorescently labeled and then incubated with K562 cells to measure the binding to Fc $\gamma$ RIIa. The only known receptor that binds SAP on the surface of K562 cells is Fc $\gamma$ RIIa ((34) and Fig S3). WT SAP

bound to Fc $\gamma$ RIIa with a  $K_d$  of  $19.7 \pm 3.4$  nM (Fig 5A and Table 1). Previous measurements of the  $K_d$  for SAP binding to Fc $\gamma$ RIIa range from 0.29 nM to 1.4  $\mu$ M (11, 23, 24). As previously described, these inconsistencies are most likely caused by the method of receptor expression and how the  $K_d$  was estimated (38). Of the 6 SAP variants tested, compared to WT SAP, E126A, Q128A, and K130V had significant differences in their binding to Fc $\gamma$ RIIa (Fig 5A and Table I). These changes in affinity correlate with the ability of these SAP variants to reduce neutrophil adhesion. Variants E126A and Q128A have a higher affinity for Fc $\gamma$ RIIa and have an increased inhibitory effect on neutrophil adhesion (Table I and Fig. 1). Conversely, variant K130V has a decreased affinity for Fc $\gamma$ RIIa and has a reduced inhibitory effect on neutrophil adhesion to fibronectin (Table I and Fig. 1). In addition, variant E126A has a Hill coefficient of  $3.3 \pm 0.5$ , indicating cooperativity in SAP E126A-Fc $\gamma$ RIIa binding. This cooperativity is absent from the other SAP variants as their Hill coefficient is not significantly different from the  $1.2 \pm 0.2$  we measured for WT SAP. One possible explanation for this increased Hill coefficient is self-aggregation of SAP E126A following binding to cells. This would then manifest as an increase in the maximal binding ( $B_{max}$ ) of SAP E126A to cells. However, the SAP E126A  $B_{max}$  was  $69.7 \pm 10.8$  % of WT SAP (mean  $\pm$  SEM, *p* not significant by *t*-test), indicating that SAP E126A was not aggregating on the surface of cells.

To measure the binding of our 6 SAP variants to Fc $\gamma$ RI, we used Fc $\gamma$ RI<sup>+</sup> HEK293 cells (Fig S3). The Fc $\gamma$ RI<sup>+</sup> HEK293 cells were co-transfected with FcR $\gamma$ , as this intracellular protein is necessary for Fc $\gamma$ RI localization to the cell membrane (42). The mock transfected cells were used to estimate the non-specific binding (Fig. 6). WT SAP bound to Fc $\gamma$ RI with a  $K_d$  of  $4.6 \pm 0.8$  nM and a Hill coefficient of  $2.1 \pm 0.6$  (Table I). This affinity matches the previously published  $K_d$  of 4.3 nM (11). The Hill coefficient of greater than 1 for the SAP-Fc $\gamma$ RI interaction suggests cooperativity, and could indicate Fc $\gamma$ RI receptor-receptor interactions. Compared to WT SAP, variants Q55A, E126A, and Q128A had significant changes in their affinity for Fc $\gamma$ RI (Fig 5B and Table I). However, we did not observe any significant changes in the  $B_{max}$  values of Q55A ( $105 \pm 36.7$  % of WT SAP), E126A ( $123.9 \pm 45.2$ ), and E128A ( $110.9 \pm 14.4$ ) when binding to Fc $\gamma$ RI. All the changes in affinity observed for Q55A, E126A, E128A correlate with the ability of these variants to inhibit fibrocyte differentiation. Variant Q55A has a higher affinity for Fc $\gamma$ RI and increased inhibitory effect on fibrocyte differentiation (Table I and Table SI). SAP variant E126A has decreased affinity for Fc $\gamma$ RI and has reduced inhibitory effect on fibrocyte differentiation (Table I and Table SI). The decrease in affinity of Q128A variant for Fc $\gamma$ RI does not alter the  $IC_{50}$  of SAP for fibrocyte differentiation but it does abrogate enhancement of phagocytosis by macrophages (Fig 3A and Table SI). Additionally, the decrease in affinity of SAP Q128A for Fc $\gamma$ RI significantly increases the Hill coefficient from  $1.8 \pm 0.1$  in WT SAP to  $2.7 \pm 0.3$  in our fibrocyte differentiation assay (Table SI). This indicates that although the  $IC_{50}$  is not altered by this mutation, Q128 still plays a role in inhibition of fibrocyte differentiation by SAP. Together, our data indicates that SAP binds to Fc $\gamma$ RIIa on neutrophils to reduce neutrophil adhesion and to Fc $\gamma$ RI on monocytes to inhibit fibrocyte differentiation.

### **Fc $\gamma$ RIIa and Fc $\gamma$ RIIIb ligation reduces neutrophil adhesion**

Our analysis of SAP variant binding to Fc $\gamma$ RI and Fc $\gamma$ RIIa indicates the significance of Fc $\gamma$ RIIa in SAP reduction of neutrophil adhesion. To corroborate this finding, we investigated the effect of antibody-mediated ligation of Fc $\gamma$ RI, Fc $\gamma$ RIIa, and Fc $\gamma$ RIIIb on neutrophil adhesion. Mouse IgG1 and the anti-Fc $\gamma$ RI antibody clone 10.1 had no effect on neutrophil adhesion (Fig. 7). However, 2 of 3 anti-Fc $\gamma$ RIIa antibodies and an anti-Fc $\gamma$ RIIIb antibody tested decreased neutrophil adhesion (Fig. 7). The anti-Fc $\gamma$ RII antibody clone AT10 is a blocking antibody and reduces IgG binding to Fc $\gamma$ RII, but in our assay it had no effect on neutrophil adhesion. Together, this indicates that SAP binds Fc $\gamma$ RIIa to decrease neutrophil adhesion, and that Fc $\gamma$ RIIIb ligation by antibodies can also reduce neutrophil adhesion to fibronectin.

### **Blocking Fc $\gamma$ RI using antibodies abrogates SAP inhibition of fibrocyte differentiation**

SAP variant binding to Fc $\gamma$ RI and the functional data on fibrocyte differentiation suggests a critical role for Fc $\gamma$ RI in mediating the SAP effect on fibrocytes. To test this hypothesis, we incubated PBMCs with F(ab')<sub>2</sub> fragments of anti-Fc $\gamma$ RI, anti-Fc $\gamma$ RII, and anti-Fc $\gamma$ RIII antibodies in the presence or absence of WT SAP. The anti-Fc $\gamma$ RII and anti-Fc $\gamma$ RIII antibodies used in this experiment do not discriminate between the different Fc $\gamma$ RII or Fc $\gamma$ RIII isoforms (43). However, they do block IgG binding to all Fc $\gamma$ RII or Fc $\gamma$ RIII receptors (43). Mouse IgG1, anti-Fc $\gamma$ RII, and anti-Fc $\gamma$ RIII antibodies had no effect on the inhibitory effect of SAP on fibrocyte differentiation (Fig. 8). However, the F(ab')<sub>2</sub> fragment of anti-Fc $\gamma$ RI antibody clone 10.1 reduced the ability of SAP to inhibit fibrocyte differentiation (Fig. 8). This indicates that SAP binds to Fc $\gamma$ RI to inhibit fibrocyte differentiation.

### **Identification of a novel Fc $\gamma$ receptor binding site on SAP**

Following our functional assays, we mapped all the mutated amino acids onto the SAP structure (Fig 9A). Excluding amino acid E153, all the functionally significant amino acids form a distinct binding site on the surface of SAP (Fig 9B and 9C). This novel binding site is different from the previously identified Fc $\gamma$ RIIa binding site (Fig 9B and (24)). The position of this novel binding site on SAP may allow for the binding of multiple Fc $\gamma$  receptors.

## **Discussion**

The pentraxin Serum Amyloid P (SAP) is an anti-fibrotic agent that inhibits aberrant scar tissue formation by regulating neutrophils, monocytes, and macrophages (11, 12, 24, 44, 45). All SAP functions appear to be mediated partly through Fc $\gamma$  receptors (11, 21, 46). As there are multiple Fc $\gamma$  receptors on neutrophils, monocytes, and macrophages, we determined how each receptor contributed to different SAP function. We found through site-directed mutagenesis that SAP binds to Fc $\gamma$ RI on monocytes to inhibit fibrocyte differentiation and to Fc $\gamma$ RIIa on neutrophils to reduce adhesion to fibronectin. In addition, we identified a novel Fc $\gamma$  receptor binding site on SAP.

Mutations in SAP that affect binding to Fc $\gamma$ RIIa significantly change the ability of SAP to reduce neutrophil adhesion to fibronectin. Similar to SAP, ligating Fc $\gamma$ RIIa by anti-Fc $\gamma$ RII



antibodies decreases neutrophil adhesion. This suggests that SAP binds to Fc $\gamma$ RIIa to decrease cell adhesion. The activation of Fc $\gamma$ RIIa results in the phosphorylation of the immunoreceptor tyrosine-based activation motif (ITAM) in the cytosolic region of this receptor (47). ITAM phosphorylation is implicated in inside-out signaling and regulation of adhesion molecules (47). This then suggests that Fc $\gamma$ RIIa activation can reduce adhesion of neutrophils to fibronectin by regulating adhesion molecules on neutrophils. Additionally, ligating Fc $\gamma$ RIIIb by an anti-Fc $\gamma$ RIII antibody reduces neutrophil adhesion to fibronectin, suggesting that ligands of Fc $\gamma$ RIIIb such as immunoglobulin could also regulate neutrophil adhesion to fibronectin.

Mutations in SAP that affect binding to Fc $\gamma$ RI significantly alter the ability of SAP to inhibit fibrocyte differentiation. In addition, blocking Fc $\gamma$ RI with an IgG blocking antibody reduces the SAP effect on fibrocyte differentiation. Together, this suggests that although there are multiple Fc $\gamma$  receptors on monocytes, SAP activates Fc $\gamma$ RI to inhibit fibrocyte differentiation. This is in agreement with our previous results where we observed that siRNA knock down of Fc $\gamma$ RI in humans results in decreased inhibitory effect of SAP on fibrocyte differentiation (21), and that cross-linking Fc $\gamma$ RI with antibodies can mimic the inhibitory effect of SAP on fibrocyte differentiation (48). Here, we have identified a Fc $\gamma$ RI binding site on each SAP monomer, suggesting that SAP can cross-link multiple Fc $\gamma$  receptors (Fig 9). Together, this suggests a role for Fc $\gamma$ RI cross-linking in SAP inhibition of fibrocyte differentiation.

SAP appears to promote phagocytosis of bio-particles such as Zymosan A through Fc $\gamma$ RI (Table I and (27)). However, not all SAP variants with alteration in Fc $\gamma$ RI binding have defects in phagocytosis. For instance, variant E126A has a ~ 10 fold reduction in affinity for Fc $\gamma$ RI and defects in inhibiting fibrocyte differentiation, but has no deficiencies in promoting phagocytosis. This suggests that a SAP opsonized bio-particle activates Fc $\gamma$ RI to promote phagocytosis in a manner that is distinct from how SAP activates Fc $\gamma$ RI to inhibit fibrocyte formation. This is supported by the fact that Fc $\gamma$ RI mediated phagocytosis is Syk-kinase dependent but inhibition of fibrocyte differentiation by SAP is Syk-kinase independent (46, 49–51). It is also possible that SAP binding to Fc $\gamma$ RI is sufficient to promote phagocytosis irrespective of changes in SAP-Fc $\gamma$ RI affinity. Alternatively, it is feasible that Q128A and E153A modulate Zymosan A phagocytosis by altering macrophage activation. However, SAP opsonized Zymosan A particles were incubated with macrophages for a short time (60 minutes), which would not allow for significant alteration in macrophage activation and phenotype.

In surface plasmon resonance experiments, SAP binds to all of the Fc $\gamma$  receptors (11, 23). However, we observed that SAP only binds to endogenous Fc $\gamma$ RI and Fc $\gamma$ RIIa on immune cells. This inconsistency can be explained by the differences in the glycosylation state of the receptors and/or the lack of some intracellular signaling components (38). Fc $\gamma$ RIIIa is a highly glycosylated receptor in humans. Modifying Fc $\gamma$ RIIIa glycosylation changes its affinity for IgG and perhaps SAP (38). Fc $\gamma$ RI and Fc $\gamma$ RIIIa in humans interact with an intracellular protein called Fc $\epsilon$  common  $\gamma$ -chain (FcR $\gamma$ ) (38, 42). The absence of FcR $\gamma$  alters the affinity of Fc $\gamma$ RI and Fc $\gamma$ RIIIa for IgG in humans (38, 42). This can potentially alter SAP binding to Fc $\gamma$ RI and Fc $\gamma$ RIIIa. Together, this suggests that the SAP affinity for Fc $\gamma$

receptors is dependent on the modification of these Fc receptors and the interactions they make prior to binding SAP.

Our findings indicate that it is possible to mimic specific SAP functions by targeting particular Fc $\gamma$  receptors (Fig 10). For instance, activation of Fc $\gamma$ RIIa by antibodies or small molecules could be used to decrease neutrophil adhesion and hence reduce neutrophil accumulation in lungs of patients suffering from acute respiratory distress syndrome or cystic fibrosis. Similarly, blocking SAP binding to Fc $\gamma$ RI might promote fibrocyte differentiation and wound healing. In addition, our results suggest that altering the SAP sequence could improve its ability to inhibit fibrocyte differentiation and/or reduce neutrophil adhesion. This could lead to the development of a more potent SAP anti-fibrotic.

## Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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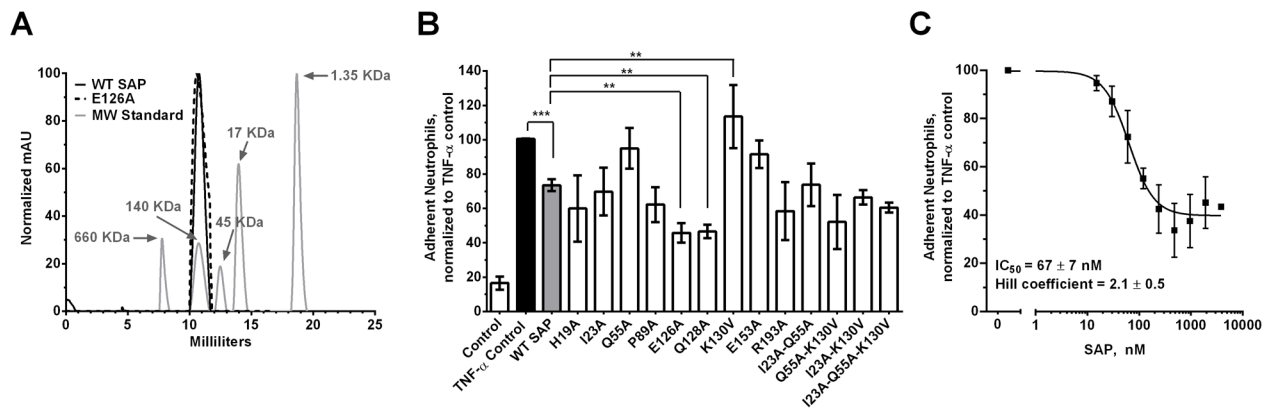
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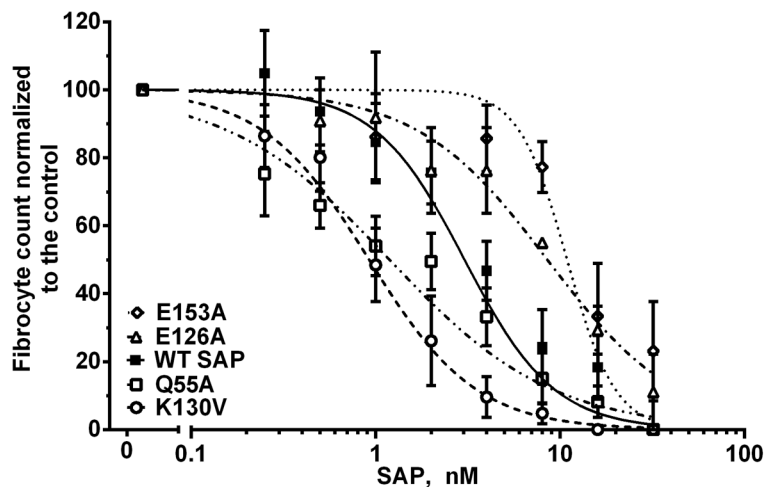
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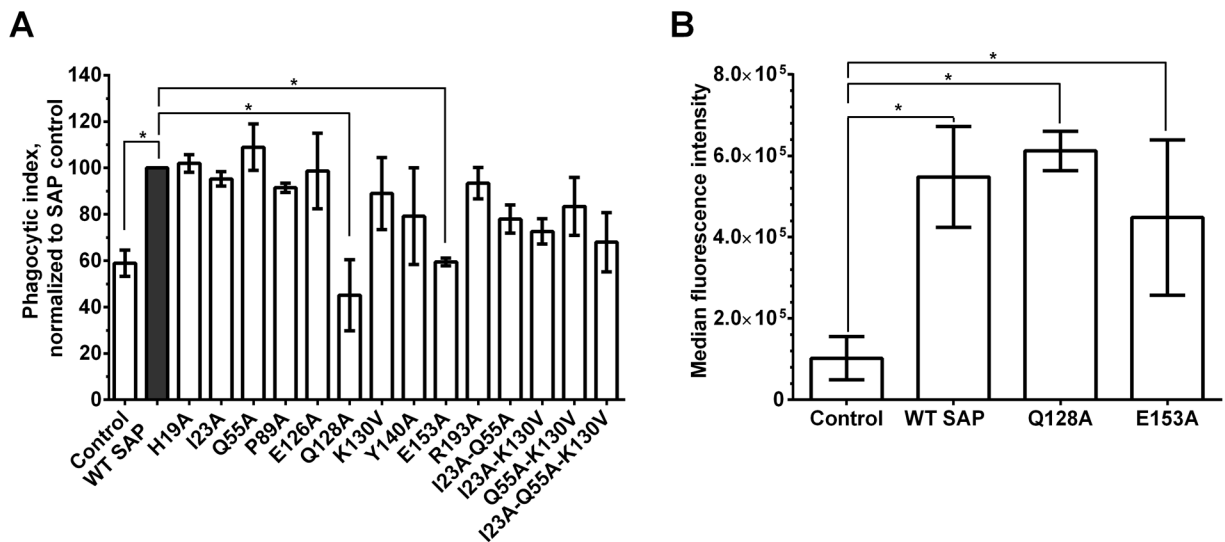
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**Figure 1. Some SAP variants have an altered effect on neutrophil adhesion to human fibronectin**  
**A)** A representative plot indicating that both WT SAP and K87A variant are pentameric. **B)** Human neutrophils were incubated with 80 nM (10  $\mu$ g/ml) of the indicated SAP variants. Following the initial incubation with SAP, the neutrophils were transferred to a fibronectin coated plate and were then activated by the addition of TNF- $\alpha$ . After 30 minutes, the non-adherent neutrophils were removed and the remaining cells were stained and counted. **C)** Human neutrophils were incubated with increasing concentrations of WT SAP to estimate the IC<sub>50</sub> of SAP for reducing neutrophil adhesion. The data were fit to a sigmoidal dose response curve with variable Hill coefficient. Values are adherent neutrophils normalized to the TNF- $\alpha$  control  $\pm$  SEM, n=3–7. \*\* represents p<0.01 and \*\*\* represents <0.001 by t-test.

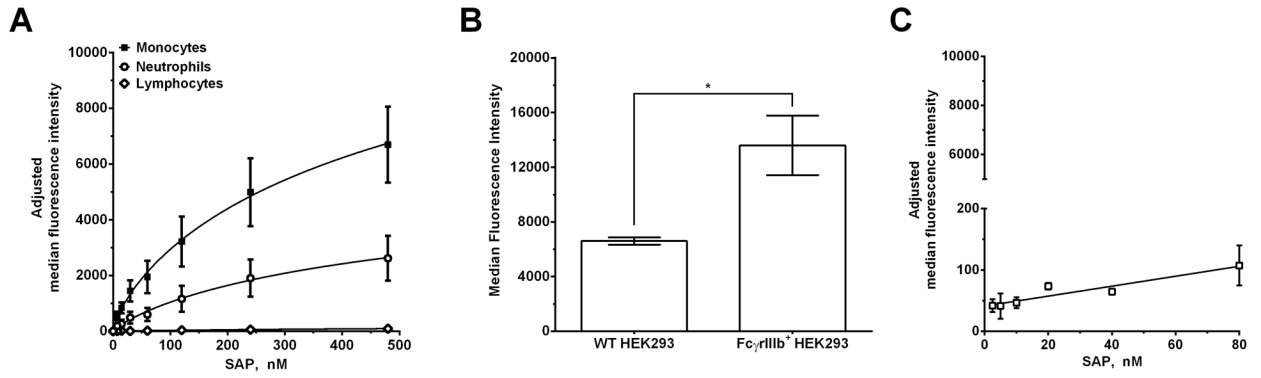


**Figure 2. Some SAP variants have an altered effect on human fibrocyte differentiation**  
 PBMCs were incubated for 5 days in the presence of WT SAP or SAP variants. Compared to WT SAP, the SAP variants K130V and Q55A were more effective inhibitors of fibrocyte differentiation, whereas E126A and E153A had reduced activity. Values are fibrocyte count normalized to the no-SAP control  $\pm$  SEM,  $n=3-5$ . The data were fit to sigmoidal dose response curves with variable Hill coefficients. The absence of error bars indicates that the error was smaller than the plot symbol.



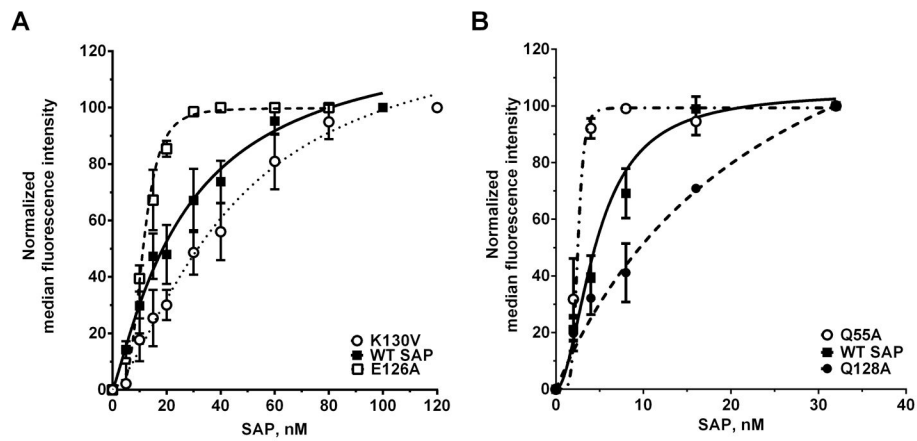
**Figure 3. Amino acids Q128 and E153 on SAP are necessary for SAP-mediated phagocytosis**  
**A)** FITC-labeled Zymosan A bio-particles were incubated in the presence of WT SAP or SAP variants and then added to monocyte-derived macrophages. After 1 hour, the free bio-particles were removed and the number of phagocytized particles was counted. The phagocytic index was estimated as the number of bio-particles engulfed by 100 macrophages. Values are means of the phagocytic index normalized to the WT SAP  $\pm$  SEM,  $n=3-6$ . **B)** The binding of WT, Q128A, and E153A SAP to Zymosan A was detected using an anti-SAP antibody and flow cytometry. Values are mean  $\pm$  SEM,  $n=3$ . \* represents  $p<0.05$  by t-test.





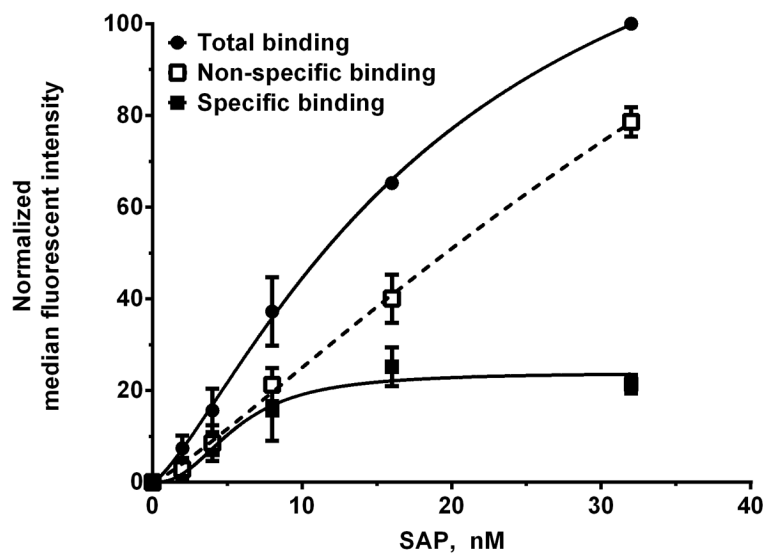
**Figure 4. Alexa Fluor 647-labeled SAP (SAP-f) binds to monocytes and neutrophils**

**A)** SAP-f was incubated with isolated leukocytes and then subjected to flow cytometry. Neutrophils, monocytes, and lymphocytes were identified based on forward scatter and side scatter as in Figure S2. Autofluorescence values were subtracted from the total binding values. Curves are fits of the resulting data to models of one-site binding with variable Hill coefficient. **B)** WT HEK293 cells and HEK293 cells expressing Fc $\gamma$ RIIIb were incubated with 20  $\mu$ g/ml of human IgG-Alexa Fluor 488 and then subjected to flow cytometry to determine if the expressed receptor is functional and therefore binds IgG. Values are mean  $\pm$  SEM, n=3. \* represents p<0.05 by t-test. **C)** HEK293 cells expressing Fc $\gamma$ RIIIb were incubated with SAP-f and then subjected to flow cytometry to measure SAP binding. Mock transfected cells were used to estimate the non-specific binding, which was then subtracted from the total binding. Values are adjusted median fluorescence intensity  $\pm$  SEM, n=3. The data were fit to a line. The absence of error bars indicates that the error was smaller than the plot symbol.

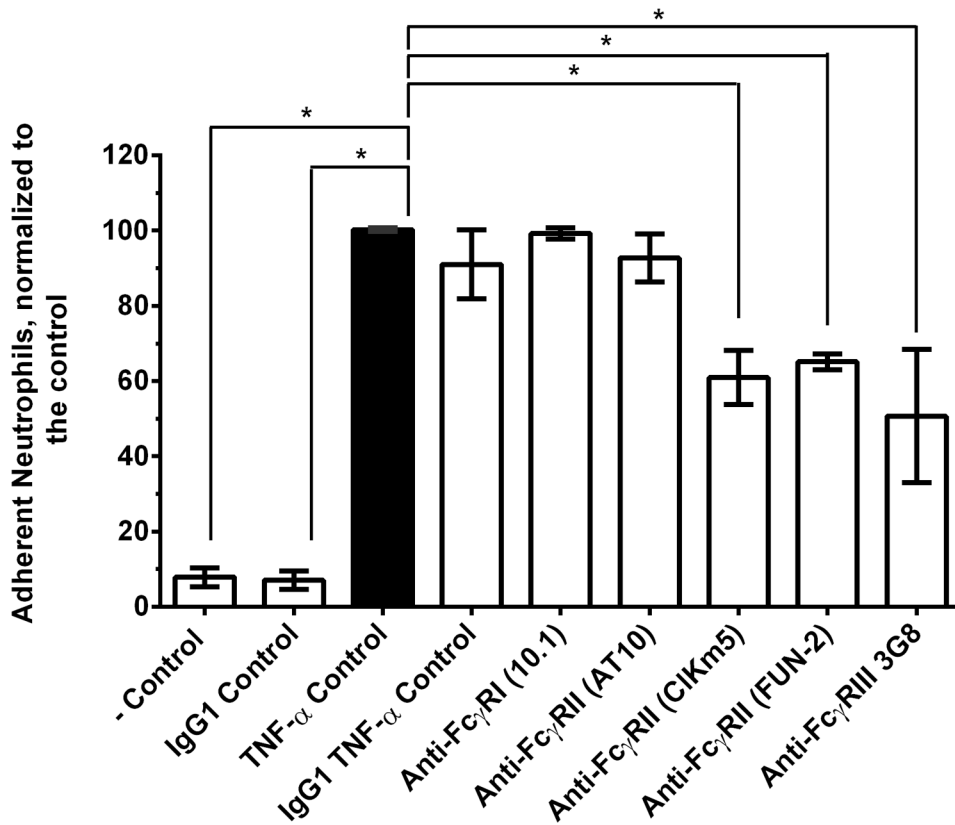


**Figure 5. SAP variant binding to Fc $\gamma$ RIIa and Fc $\gamma$ RI**

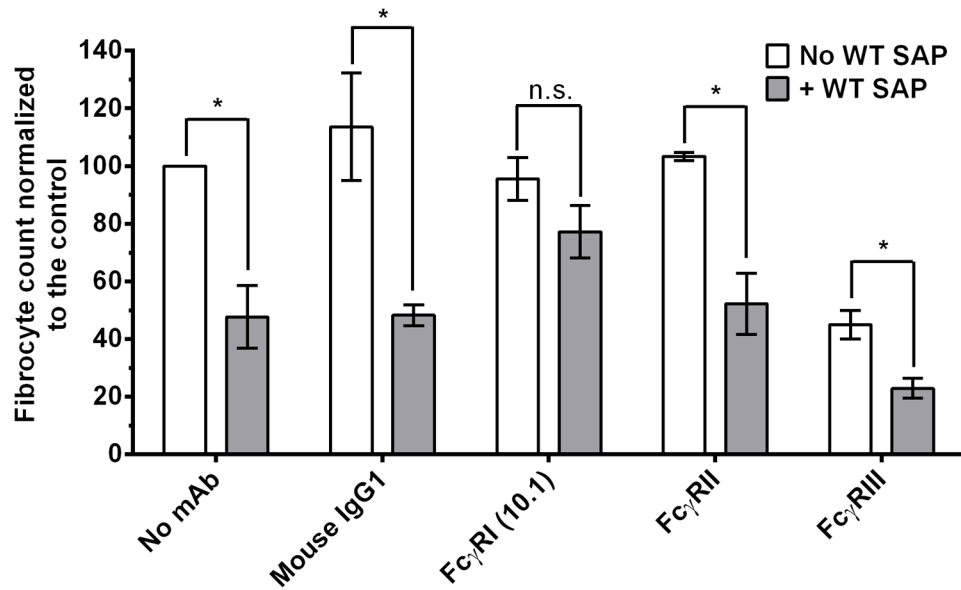
**A)** K562 cells, which express Fc $\gamma$ RIIa, were incubated with fluorescently-labeled SAP variants. The cells were then washed and the binding of the labeled SAP to the cells was measured by flow cytometry. **B)** HEK293 cells expressing Fc $\gamma$ RI were incubated with fluorescently-labeled SAP variants and then binding was measured by flow cytometry. Mock transfected cells were used to estimate the non-specific binding. Median fluorescence intensity values were normalized to the intensity value of the highest SAP concentration. Values are normalized mean  $\pm$  SEM,  $n=3-5$ . Curves are fits to models of one-site binding with variable Hill coefficient. The absence of error bars indicates that the error was smaller than the plot symbol.



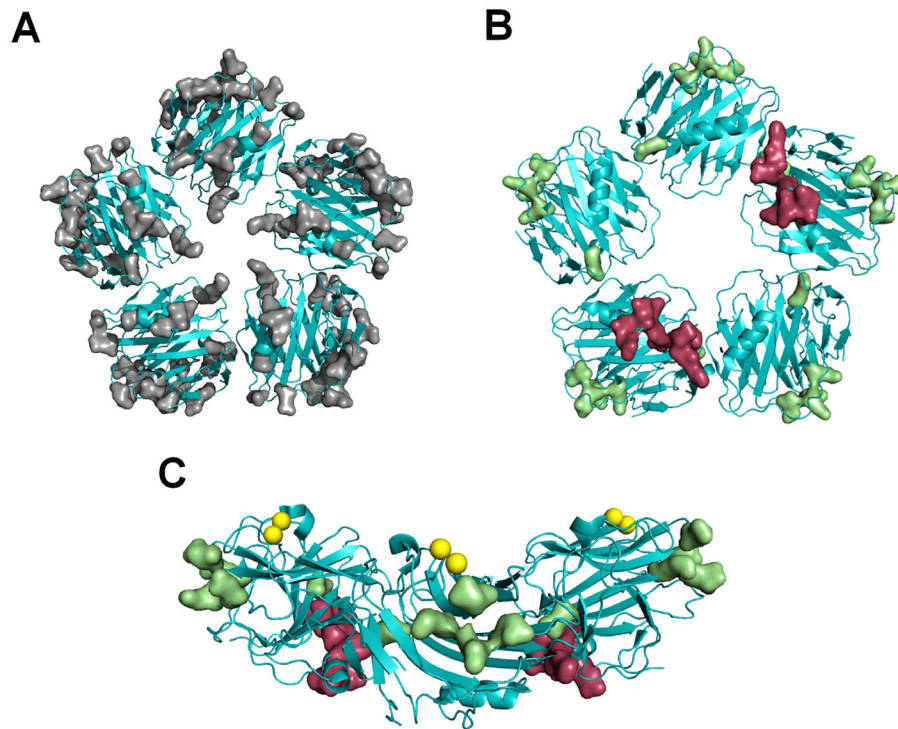
**Figure 6. Estimating the specific binding of WT SAP to Fc $\gamma$ RI**  
 HEK293 cells expressing Fc $\gamma$ RI were incubated with Alexa Fluor 647-labeled WT SAP (SAP-f) as in **Table 1**. The cells were then washed and the binding of SAP-f to the cells was measured by flow cytometry. The specific binding was estimated by subtracting the binding of SAP-f to mock transfected cells from Fc $\gamma$ RI expressing cells. The values were normalized to the median fluorescence intensity of SAP total binding at 32 nM. Values are mean  $\pm$  SEM, n=3. The absence of error bars indicates that the error was smaller than the plot symbol.



**Figure 7. Ligating Fc $\gamma$ RIIa and Fc $\gamma$ RIIIb by antibodies decreases neutrophil adhesion**  
 Neutrophils were incubated with the indicated anti-Fc $\gamma$  receptor antibodies to assess the effect of Fc $\gamma$  receptor ligation on neutrophil adhesion to fibronectin as described in Figure 1. Values are adherent neutrophils normalized to the TNF- $\alpha$  control, mean  $\pm$  SEM, n=3. \* indicates p<0.05 by t-test.

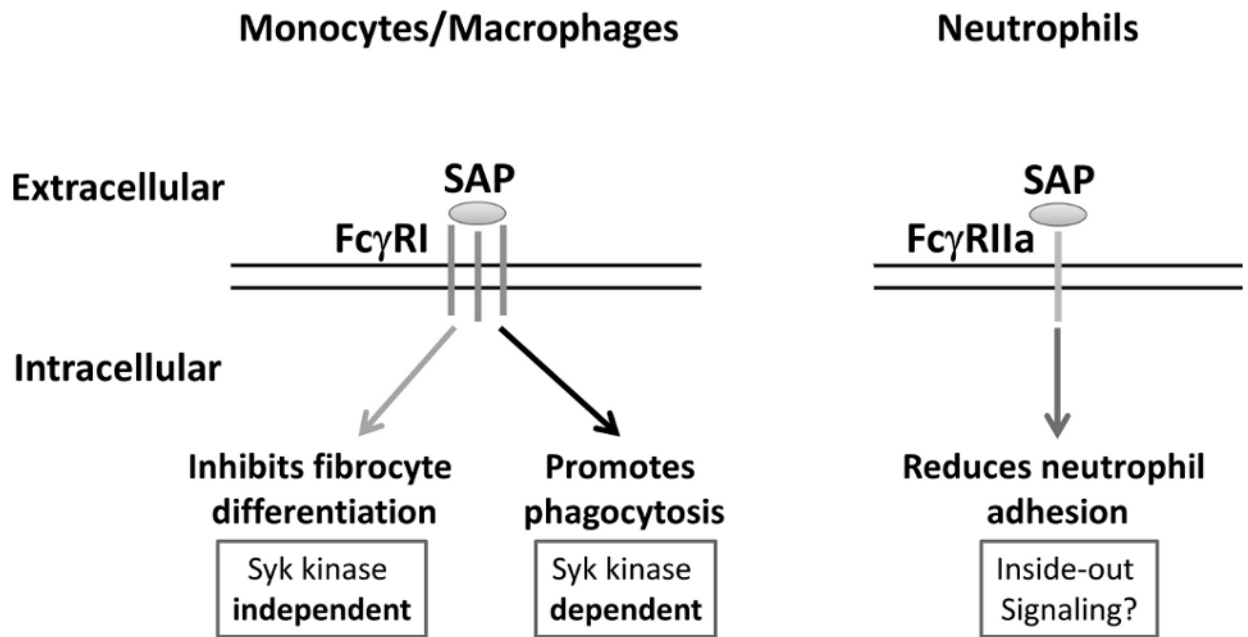


**Figure 8. Fc $\gamma$ RI blocking antibodies reduce the effect of SAP on fibrocyte differentiation**  
 PBMCs were incubated with F(ab')<sub>2</sub> fragments of anti-Fc $\gamma$  receptor antibodies in the presence or absence of WT SAP to determine their effect on SAP-mediated inhibition of fibrocyte differentiation. Values are fibrocyte count normalized to the control  $\pm$  SEM, n=4. \* represent p <0.05 by t-test; n.s. indicates not significant.



**Figure 9. Identification of a novel Fc $\gamma$  receptor binding site on SAP**

**A)** Mutated amino acid residues are indicated by molecular surface representation (gray) on the SAP structure. **B)** The functionally significant amino acid residues (green) are distinct from the previously identified Fc $\gamma$ RIIa binding site (red). **C)** When E153 is excluded; the remaining functionally significant amino acid residues form a distinct binding site. The yellow spheres represent the two calciums bound to SAP.



**Figure 10. A model of the SAP effect on monocytes and neutrophils**

SAP cross-links Fc $\gamma$ RI to inhibit fibrocyte differentiation in a Syk kinase independent manner. SAP also binds Fc $\gamma$ RI to promote phagocytosis of zymosan A by macrophages. However, this pathway is Syk kinase dependent. SAP binds to Fc $\gamma$ RIIa which then phosphorylates the ITAM domain of Fc $\gamma$ RIIa. This then regulates adhesion molecules on the surface of neutrophils and decrease neutrophil adhesion.

**Table I**  
**Binding of SAP variants to Fc $\gamma$ RI and Fc $\gamma$ RIIa**

HEK293 cells expressing Fc $\gamma$ RI were incubated with Alexa Fluor 647-labeled SAP variants. The cells were then washed and the binding of the labeled SAP to the cells was measured by flow cytometry. Mock transfected cells were used to estimate the non-specific binding. K562 cells were used to measure the binding of SAP variants to Fc $\gamma$ RIIa.

SAP Variants	Fc $\gamma$ RI K <sub>d</sub> (nM $\pm$ SEM)	Hill coefficient	Fc $\gamma$ RIIa K <sub>d</sub> (nM $\pm$ SEM)	Hill coefficient
WT SAP	4.6 $\pm$ 0.8	2.1 $\pm$ 0.6	19.7 $\pm$ 3.4	1.2 $\pm$ 0.2
I23G	9.8 $\pm$ 5.8	1.1 $\pm$ 0.4	25.2 $\pm$ 9.8	1.5 $\pm$ 0.4
Q55A	2.6 $\pm$ 0.1 ***	6.7 $\pm$ 0.2 ***	22.3 $\pm$ 3.3	1.7 $\pm$ 0.2
E126A	36.9 $\pm$ 2.8***	0.9 $\pm$ 0.1	11.9 $\pm$ 0.6 *	3.3 $\pm$ 0.5 *
Q128A	24.7 $\pm$ 6.0 *	0.9 $\pm$ 0.1*	11.9 $\pm$ 2.1 *	2.3 $\pm$ 0.6
K130V	4.5 $\pm$ 0.7	3.7 $\pm$ 1.9	43.7 $\pm$ 8.7 *	1.4 $\pm$ 0.4
E153A	5.3 $\pm$ 0.6	3.2 $\pm$ 0.9	27.8 $\pm$ 1.4	4.7 $\pm$ 0.9

Values are mean  $\pm$  SEM, n=3–6.

\* indicates  $p < 0.05$  and \*\*\* indicates  $p < 0.001$  by t-test when compared to the corresponding wildtype (WT) control.