

1 **A method to detect the binding of hyper-glycosylated fragment crystallizable**  
2 **(Fc) region of human IgG1 to glycan receptors**

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7 **Summary**

8 Engineering the fragment crystallizable (Fc) of human IgG1 can bring improved effector  
9 functions to monoclonal antibodies and Fc fusion-based medicines and vaccines. Such Fc-  
10 effector functions are largely controlled by post-translational modifications (PTMs) within  
11 the Fc, including the addition of glycans that introduce structural and functional  
12 heterogeneity to this class of therapeutic. Here we describe a detailed method to allow the  
13 detection of hyper-sialylated Fcs binding to glycan receptors that will facilitate the future  
14 development of new mAbs and Fc-fragment therapies and vaccines.

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16 **Key words:** IgG1, glycans, glycosylation, Fc receptors, antibody-dependent cell-mediated  
17 cytotoxicity (ADCC), antibody-dependent cell-mediated phagocytosis (ADCP), complement-  
18 dependent cytotoxicity (CDC), effector function, therapeutic antibodies.

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## 24 Introduction

25 Immunoglobulin G antibodies are glycosylated at asparagine 297 (Asn297) of the unique N-  
26 linked sequon located within the Fc (Czajkowsky et al. 2012; Lund et al. 1996). The Fc  
27 glycans attached at Asn297 are typically biantennary complex types, exhibiting high levels of  
28 fucosylation of the core GlcNAc residue, partial galactosylation and bisecting GlcNAc, and of  
29 these structures less than 20% are sialylated (Dalziel et al. 2014). The low levels of branching  
30 and terminal structures, such as sialic acid, are a consequence of constraints imposed on  
31 Asn297 glycan processing by the Fc protein backbone (Frank et al. 2014; Subedi et al. 2014).

32 The composition of glycans attached at Asn297 significantly affects Fc-mediated  
33 interactions with different receptors (Dalziel et al. 2014), and multiple lines of evidence  
34 have shown that glycosylation is critical to driving either the anti- or pro-inflammatory  
35 capability of IgG1 (Schwab and Nimmerjahn 2013). Glycosylation of Asn297 in the Fc is thus  
36 essential for interactions with type 1 receptors (Fc $\gamma$ ) and type 2 receptors (glycan  
37 dependent) and is also necessary for driving interactions with the complement cascade  
38 (Czajkowsky et al. 2012). Although protocols essentially are well described for detecting  
39 interactions with type I Fc $\gamma$  receptors, detailed protocols for the detection of such reagents  
40 to type 2 glycan receptors are less commonly described in the literature.

41 In humans, infusion of Fc fragments is sufficient to ameliorate idiopathic  
42 thrombocytopenic purpura (ITP) in children, demonstrating the therapeutic utility of the Fc  
43 *in vivo* (Debre et al. 1993). The anti-inflammatory property of the Fc is lost after  
44 deglycosylation of IgG1, and a small population of IgG1-bearing sialylated Fcs has been  
45 identified as making a significant contribution to the control of inflammation in animal  
46 models (Anthony et al. 2011; Anthony et al. 2008; Washburn et al. 2015). Higher levels of  
47 sialylation also lead to longer serum retention times (Li et al. 2006; Liu 2015), and studies in

48 humans and mice have shown that influx and efflux of IgG1 into the central nervous system  
49 (CNS) is glycan and sialic acid dependent (Finke et al. 2017; St-Amour et al. 2013; Zhang et  
50 al. 2004). Consequently, IgG1-Fc sialylation has emerged as an important but controversial  
51 concept for regulating anti-inflammatory activity of antibodies and Fc fragments (Schwab  
52 and Nimmerjahn 2013).

53           Methods to enhance the sialylation of the Fc have largely focused on modifications  
54 to Asn297 attached glycans (Fiebiger et al. 2015; Washburn et al. 2015). We have previously  
55 shown how Asn297-limited approaches can be overcome through the introduction of  
56 additional N-linked glycosylation sites into a limited number of exposed areas within the  
57 IgG1-Fc fragment (Blundell et al. 2017). For example, insertion of a N-terminal hinge-distal  
58 glycan site (N221) significantly increases sialylation, allowing engagement with glycan  
59 receptors not previously known to bind the Fc (Blundell et al. 2017). By adding a cysteine-  
60 silenced 18 amino-acid C-terminal extension containing an additional N-linked site (N563),  
61 further glycan complexity can also be brought to the IgG1 Fc (Fig. 1).

62           We provide a detailed protocol for the detection of glycan modified IgG1-Fc  
63 fragments containing one, two or three additional N-linked glycans to glycan receptors by  
64 enzyme-linked immunosorbent assays (ELISA).

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72 **Materials**

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74 Coating buffer: 0.05 M carbonate-bicarbonate solution (Sigma C3041), pH 9.6.

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76 Incubation and wash buffer: TMS (20 mM Tris-HCl, 150 mM NaCl, 2 mM CaCl<sub>2</sub>, 2 mM MgCl<sub>2</sub>).

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78 Blocking buffer: TMS solution containing 5% bovine serum albumin (Sigma A7030).

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80 Developing solution: Sigmafast p-Nitrophenyl phosphate dissolved in Tris-buffer (Sigma  
81 N2770).

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83 Glycan receptors: lyophilized histidine-tagged recombinant Siglec-1 (R&D Systems, 5197-SL-  
84 050) or Siglec-4/MAG (Sinobiologicals, 13186-H08H) were reconstituted to 100 µg/ml with  
85 ddH<sub>2</sub>O.

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87 Detecting antibodies: Affinity-purified alkaline-phosphatase conjugated F(ab')<sub>2</sub> fragment  
88 goat anti-human IgG1 Fcγ-fragment specific (Jackson Immuno Research 109-056-008, or  
89 Invitrogen H10108).

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91 Specialist equipment: Microplate reader capable of measuring absorbance at 405 nm  
92 wavelength, microplate washer, data analysis and graphing software.

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96 **Methods**

97 Nunc microtiter plates are coated down with recombinant glycan receptors<sup>1</sup>, typically 100 µl  
98 per well at 2-10 µg/ml in carbonate-bicarbonate coating buffer pH9.6, and incubated  
99 overnight at 4°C.

100 The following day plates are washed five times with excess TSM incubation buffer (1-  
101 5 min incubation between washes), prior to blocking for 2 hrs in 150 µl per well of TMS  
102 buffer containing 5% bovine serum albumin. Plates are then washed as before. 100 µl of  
103 varying concentrations of Fc fragments in TSM incubation buffer are added<sup>2</sup> in duplicate  
104 wells. We typically titrate the Fc fragments by doubling dilution down the plate from 50 to 0  
105 µg/ml. Receptors are allowed to bind the Fc fragments overnight at 4°C.

106 The following day plates are washed five times with excess TSM incubation buffer (1-  
107 5 min incubation between washes), prior to the addition of 100 µl per well of alkaline-  
108 phosphatase conjugated F(ab')<sub>2</sub> goat anti-human IgG1 Fcγ fragment-specific detection  
109 antibody diluted 1 in 500 in TMS buffer. Glycosylated Fc fragments that have bound to the  
110 glycan receptors are allowed to bind the conjugated antibody for 1 h at room temperature  
111 on a rocking platform.

112 Plates were washed as above and developed for 5-15 minutes with 100 µl/well of  
113 developing solution. The plates were read at 405nm wavelength using a LT-4500 microplate  
114 absorbance reader (Labtech), and the data plotted with GraphPad Prism (Fig. 1).

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120 **Notes**

121 1. We have found that this protocol is amenable to the study of many other glycan  
122 receptors. However, from our experience we advise determining background binding  
123 of the detecting antibody to each glycan receptor first, as the Fab domains within  
124 the alkaline phosphatase-conjugated detecting antibody can themselves be  
125 glycosylated (van de Bovenkamp et al. 2016). For example, we have seen direct  
126 binding of both the Jackson and Invitrogen F(ab')<sub>2</sub> detecting reagents to Siglec-5,  
127 CLEC-1B and DC-SIGNR (Blundell et al. 2017). Preliminary removal of F(ab')<sub>2</sub>  
128 associated glycans with glycosidases e.g. PNGase F (New England Biologicals) may be  
129 required for the study of certain receptors.

130 2. This protocol works equally well for Fc fragments and Fab'<sub>2</sub> fragments (both  
131 available from Jackson Immuno Research) that have previously been digested with  
132 commercially available glycosidases e.g. PNGase F, Endo H or neuraminidase. For  
133 example, Siglec-5 binding above is lost after neuraminidase treatment.

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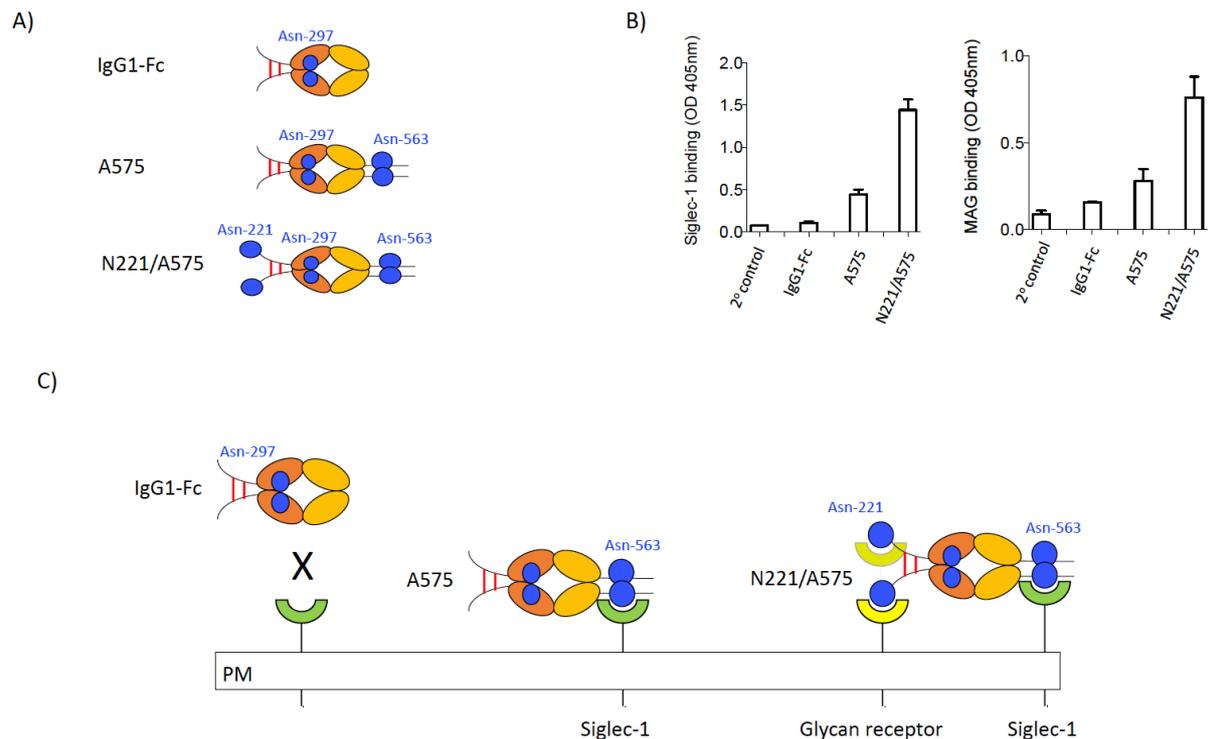
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218 **Fig. 1. Detection of hyper-sialylated Fc-fragment binding to glycan receptors by ELISA.**

219 (A) Fc fragments containing one, two or three N-linked attachment sites together with  
 220 methods for determining their glycan composition by HILIC-UPLC analysis have been  
 221 described previously (Blundell et al. 2017).

222 (B) Fc fragments containing three N-linked glycans (N221/A575) bind more strongly to both  
 223 Siglec-1 and Siglec-4 (myelin-associated glycoprotein), than Fc fragments only containing two  
 224 (A575) or one N-linked sugar (IgG1 Fc control). Error bars represent standard deviations  
 225 around the mean value, n = 3 independent experiments.

226 (C) In contrast with the Asn297 glycan, which is largely buried within the Fc cavity, Asn221  
 227 and Asn563 are located at the N- and C-terminal tips of the Fc respectively and, as our data  
 228 show, would therefore be more accessible for post-translational modifications by glycan-  
 229 modifying enzymes that permit A575 and N221/A575 to bind more strongly to sialic-acid  
 230 dependent receptors.