

Supporting Information for

Dissection of complement and Fc-receptor mediated pathomechanisms of autoantibodies to myelin oligodendrocyte glycoprotein.

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This PDF file includes:

Figures S1 Tables S1

Fig. S1.



Supplementary Figure 1: Antigen-recognition, C1q- and FcγR-binding of mutated MOG-specific mAbs.

The antibody r8-18C5 with human IgG1 Fc part and all 13 Fc mutant antibodies show comparable binding to MOG in a cell based flow cytometry assay (A). IgG binding to

MOG was normalized to binding of the r8-18C5 (delta MFI value). The assay was repeated twice for each antibody (**B**) C1q binding was analyzed by ELISA. Specifically, ED-hMOG was coated first before the addition of the indicated mAbs. C1q was added and its binding was quantified using an anti-C1q Ab. Binding of the mutant Abs was normalized to the binding of the antibody r8-18C5 and the assay was repeated three times for each antibody. (**C-F**) FcγR binding was investigated by ELISA. Anti-FLAG antibodies were coated on an ELISA plate, which bound to the soluble FcγR (FcγRI, FcγRIIb, FcγRIII, FcγRIV). We quantified the binding of antibody-MOG complexes (immune complexes) to the soluble FcγR. Binding of the mutant Abs was normalized to the binding of the antibody r8-18C5. FcyRI and FcyRIV were read 40 min after reaction started and FcyRIIb, FcyRIII were read 300 min after reaction started. The assay was repeated three times for the antibodies that were used *in vivo* (HK3, TAP, SAI and r8-18C5), while it was performed twice for all other antibodies. Data is shown as mean with SEM.

Variant (name used in our manuscript)	Introduced mutation	Previously observed effect with other mAbs	Observed effect after MOG-binding
E345K	E345K	enhanced C1q binding/ CDC ¹ enhanced FcyR binding/ ADCC ^{1,2}	intact CDC intact FcyR binding
E430G	E430G	enhanced C1q binding/CDC ¹ enhanced FcyR binding/ ADCC ²	intact CDC intact FcyR binding
LL	L234A/L235A	reduced C1q binding/ CDC ^{3,4} reduced FcyR binding/ ADCC ^{3,4}	reduced C1q reduced FcyR binding
LLP	L234A/L235A/P329G	reduced C1q binding/CDC ⁵ reduced FcyR binding/ ADCC ⁵	reduced C1q reduced FcyR binding
TAP	T299L/A330S/P331S	reduced C1q binding/CDC ⁶⁻⁸ not defined for FcyR binding/ ADCC	reduced C1q reduced FcyR binding
T299L	T299L	not defined for C1q binding/CDC reduced FcyR binding/ ADCC ⁹	reduced C1q reduced FcyR binding
KE	K326W/E333S	enhanced C1q binding/CDC ^{10,11} reduced FcyR binding/ ADCC ¹⁰	intact CDC intact FcyR binding
KQ	K320E/Q386R	enhanced C1q binding/CDC ⁹ reduced FcyR binding/ ADCC ⁹	intact CDC intact FcyR binding
TKQ	T299L/K320E/Q386R	enhanced C1q binding/CDC ⁹ intact FcyR binding/ ADCC ⁹	reduced C1q reduced FcyR binding
K322A	K322A	reduced C1q binding/CDC ¹² intact FcyR binding/ ADCC ⁴	reduced C1q intact FcyR binding
AP	A330S/P331S	reduced C1q binding/CDC ⁶ reduced FcyR binding/ ADCC ¹³	reduced C1q intact FcyR binding
SAI	S239D/A330L/I332E	reduced C1q binding/CDC ^{7,10} enhanced FcyR binding/ ADCC ^{7,10,14}	reduced C1q intact FcyR binding
I253D	1253D	reduced C1q binding/CDC ^{12,15} not defined for FcyR binding/ ADCC	reduced CDC intact FcyR binding

Table S1. Fc mutants of the MOG mAb r8-18C5 used in this study

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