



Extended Data Fig. 1. The conserved tail sequence of mIgG and the identification of the hIgG1-G396R variant

(A) Sequences of the conserved tail of membrane-bound IgG in different species. (B) The distribution of three genotypes, including hIgG1-G396R homozygotes, heterozygotes and WT, detected by TaqMan-probe based genotyping. (C) The aligned target sequences of WT, hIgG1-G396R heterozygote and homozygote. The C/T polymorphism are highlighted.