

Heavy chain variable region (CDR is underlined)

Germline IGHV 1-69	QVQLVQSGAEVKKPGSSVKVSKASGGTFSSYAISWVRQAPGQGLEWMGRIIPILGIANYA <u>QKFQGRVTITADKSTSTAYMELSSLRSED</u> TAVYYCAR---GYSYGNFDYWGQGLTVTVSS
YN-002VH	QVQLVQSGAEV R KPGSSVKVSKASGGTFSSYAISWVRQAPGQGLEWMGRIIPILGIANYA <u>QKFQGRVTITADKSTSTAYMELSSLRSED</u> TAVYYCAR TMD GYSYGNFDYWGQGLTVTVSS
YN-003VH	QVQLVQSGAEVKKPGSSVKVSKASGGTFSSYAISWVRQAPGQGLEWMGRIIPILGIANYA <u>QKFQGRVTITADKSTSTAYMELSSLRSED</u> TAVYYCAR TMD GYSYGNFDYWGQGLTVTVSS

Light chain variable region (CDR is underlined)

Germline IGLV2-14	QSALTQPASVSGSPGQSITISCTGTSSDVGGYNYVSWYQQHPGKAPKLMIEVSNRPSGV SNRFGSGSKSGNTASLTISGLQAEDEADYYC SS YTSS----VFGGGTKLTVLG
YN-002VL	QSALTQPASVSGSPGQSITISCTGTSSDVGGYNYVSWYQQHPGKAPK L LIY GNS NRPSGV PDR FSGSGSKSG TS AS L AITGLQAEDEADYYC QSYDSSL SGS VFGGGTKLTVLG
YN-003VL	QSALTQPASVSGSPGQSITISCTGTSSDVGGYNYVSWYQQHPGKAPKLMIEY GNS NRPSGV SNRFGSGSKSGNTASLTISGLQAEDEADYYC QSYDSSL SGS VFGGGTKLTVLG

Fig. 1

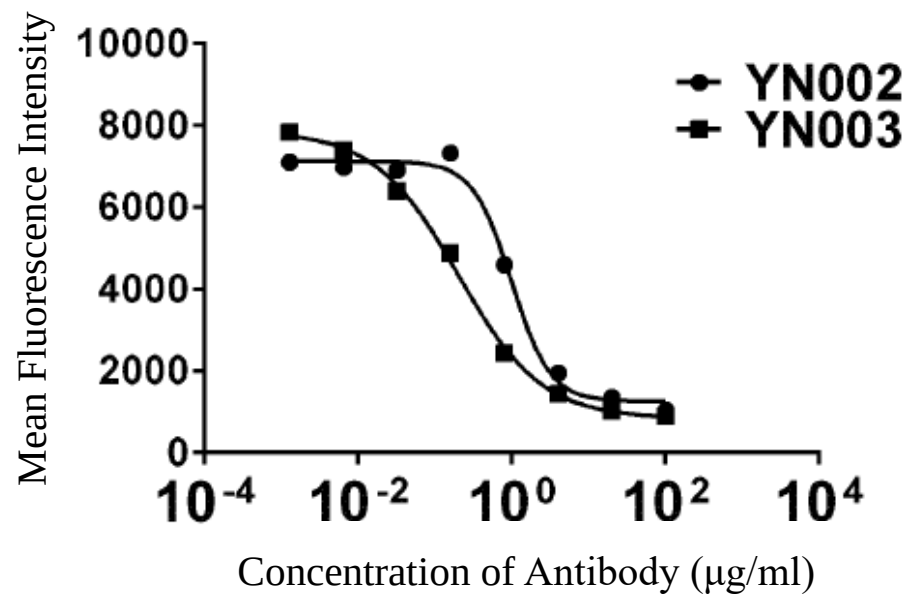


Fig. 2

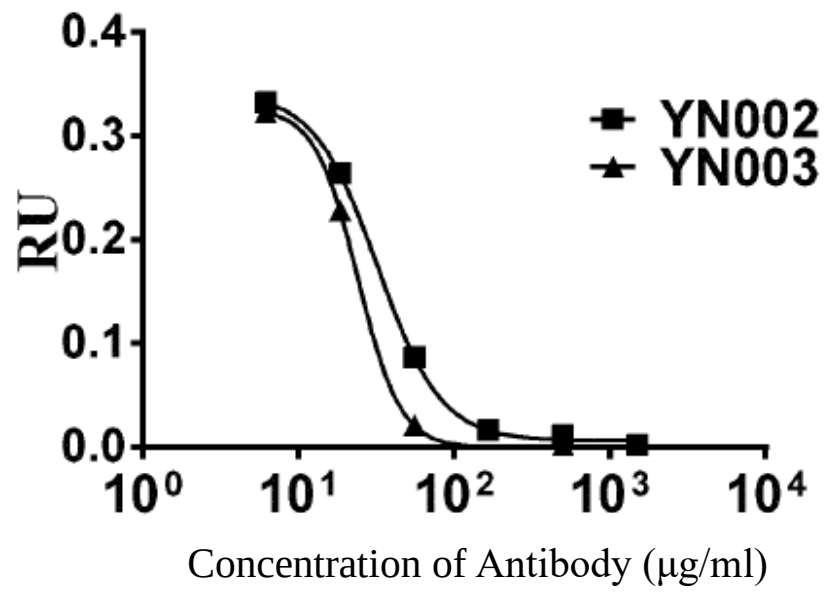


Fig. 3

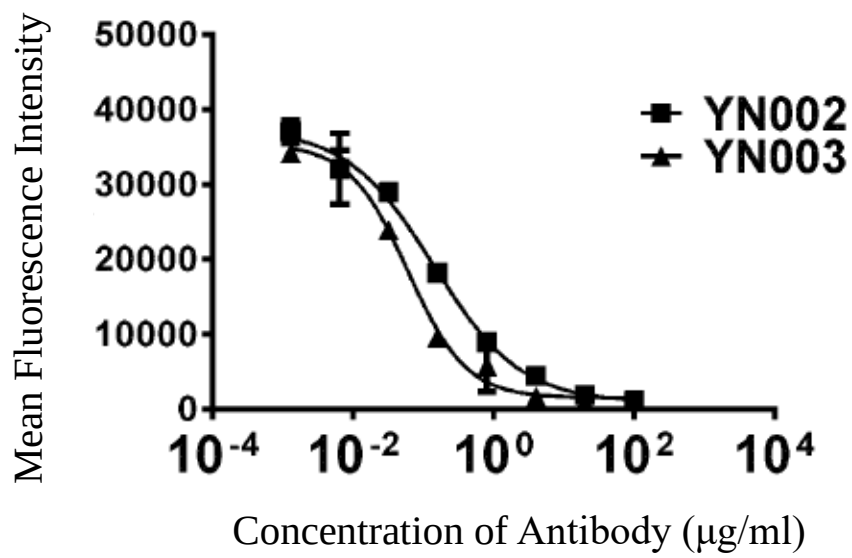


Fig. 4

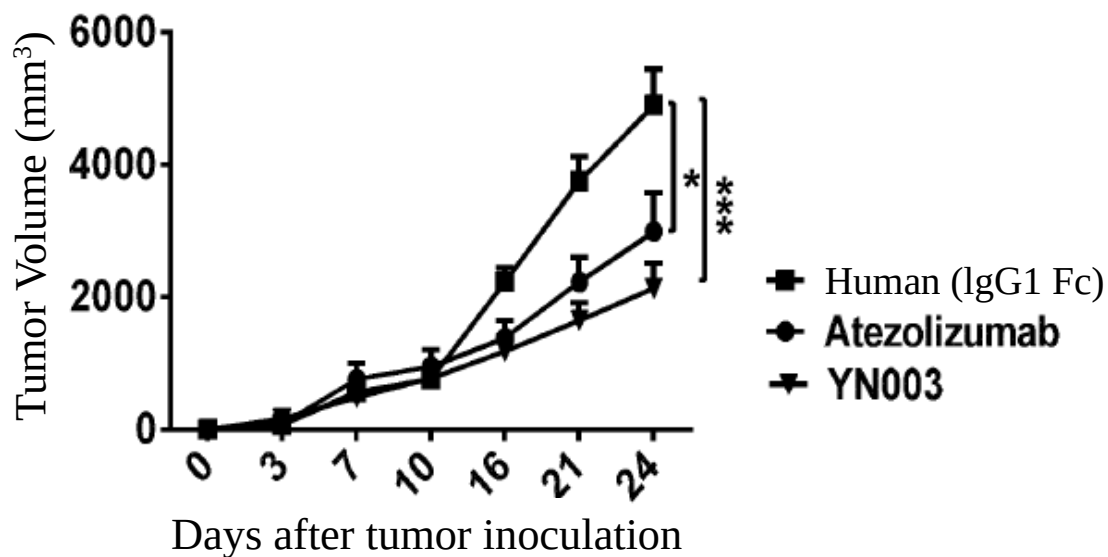


Fig. 5

Heavy chain variable region (CDR is underlined)

Germline IGHV 3-23	QVQLVESGGGLVQPGGSLRLSCAASGFTFSSYAMSWVRQAPGKGLEWVSAISGSGGSTYYAD SVKGRFTISRDN SKNTLYLQMNSLRAEDTAVYYCAKTNWG--DAFDIWGQGTMTVSS
YN-005VH	QVQLVQSGGGLVQPGGSLRLSCAASGFTFSSYAMSWVRQAPGKGLEWVSAISGSGGSTYYAD SVKGRFTISRDN SKNTLYLQMNSLRAEDTAVYYCAKTNWGPDAFDIWGQGTMTVSS
YN-006VH	QVQLVESGGGLVQPGGSLRLSCAASGFTFSSYAMSWVRQAPGKGLEWVSAISGSGGSTYYAD SVKGRFTISRDN SKNTLYLQMNSLRAEDTAVYYCAKTNWGPDAFDIWGQGTMTVSS

Light chain variable region (CDR is underlined)

Germline IGLV1-44	QSVLTQPPSASGTPGQRVTISCSGSSSNIGSNTVNWYQQLPGTAPKLLIYSNNQRPSGVPDR FSGSKSGTSASLAISGLQSEDEADYYCAAWDDSLNGYVFGTGKLTVLG
YN-005VL	QSVLTQPPSASGTPGQRVTISCSGTS SD IGSYSVNWYQQLPGTAPKLLIYSNNQRPSGVPDR FSGSKSGTSASLAISGLQSEDEADYYCAAWDDSLNGYVFGTGKLTVLG
YN-006VL	QSVLTQPPSASGTPGQRVTISCSGTS SD IGSYSVNWYQQLPGTAPKLLIYSNNQRPSGVPDR FSGSKSGTSASLAISGLQSEDEADYYCAAWDDSLNGYVFGTGKLTVLG

Fig. 6

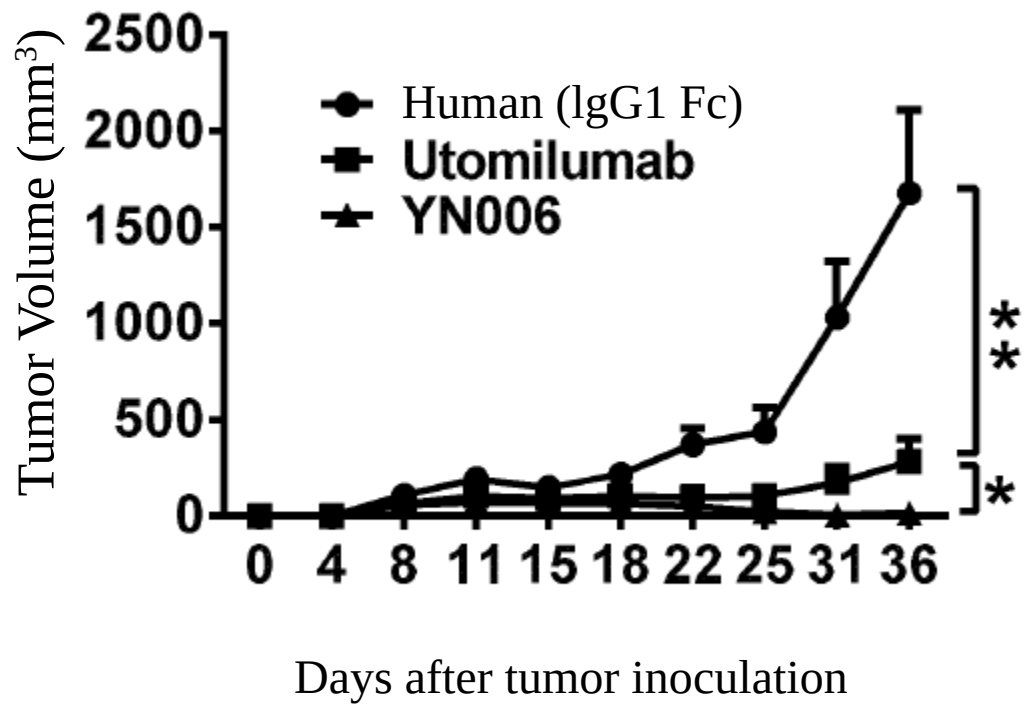


Fig. 7

Heavy chain variable region (CDR is underlined)

Germline IGHV1-69	QVQLVQSGAEVKKPGSSVKVSKKASGGTFSYAIISWVEQAPGQGLEWMGRIPILGIANVYAKFKQGRVTITADKSTSTAYMELSSLRSEDTAVYYCAR---
YN003VH	QVQLVQSGAEVKKPGSSVKVSKKASGGTFSYAIISWVEQAPGQGLEWMGRIPILGIANVYAKFKQGRVTITADKSTSTAYMELSSLRSEDTAVYYCAR <u>TMDGYSYGNFDYWGQGTLLTVSS</u>

Light chain variable region (CDR is underlined)

Germline IGLV2-14	QSALTQPPASVSGSPGQSITISCTGTSSDVGGYNVSWYQQHPGKAPKLMIEVSNRPSGVSNRFSGSKSGNTASLTISGLQAEDEADYYCSSYTSS-----VFGGGTKLTVLG
YN003VL	QSALTQPPASVSGSPGQSITISCTGTSSDVGGYNVSWYQQHPGKAPKLMIEVSNRPSGVSNRFSGSKSGNTASLTISGLQAEDEADYYC <u>QSYDSSLSSGS</u> VFGGGTKLTVLG

Fig. 8

Heavy chain variable region (CDR is underlined)

Germline IGHV1-69	QVQLVQSGAEVKKPGSSVKVSKCKASGGTFSSYAISWVRQAPGGQGLEWMGRIIPILGIANYAQKFKQGRVTTITADKSTSTAYMELSSLRSEDTAVYYCAR---
YN003VH	QVQLVQSGAEVKKPGSSVKVSKCKASGGTFSSYAISWVRQAPGGQGLEWMGRIIPILGIANYAQKFKQGRVTTITADKSTSTAYMELSSLRSEDTAVYYCAR MD GYSYGNFDYWGQGLTLTVSS
YN035VH	QVQLVQSGAEVKKPGSSVKVSKCKAS RGPFSTYA ISWVRQAPGGQGLEWMGRIIPILGIANYAQKFKQGRVTTITADKSTSTAYMELSSLRSEDTAVYYCAR MEGYGF GNFDYWGQGLTLTVSS
YN036VH	QVQLVQSGAEVKKPGSSVKVSKCKAS RGPFSTYA ISWVRQAPGGQGLEWMGRIIPILGIANYAQKFKQGRVTTITADKSTSTAYMELSSLRSEDTAVYYCAR TMGEYSYGN YDYWGQGLTLTVSS
YN037VH	QVQLVQSGAEVKKPGSSVKVSKCKAS RGPFSTYA ISWVRQAPGGQGLEWMGRIIPILGIANYAQKFKQGRVTTITADKSTSTAYMELSSLRSEDTAVYYCAR TMGEYSYGN YDYWGQGLTLTVSS
YN038VH	QVQLVQSGAEVKKPGSSVKVSKCKAS RGPFSTYA ISWVRQAPGGQGLEWMGRIIPILGIANYAQKFKQGRVTTITADKSTSTAYMELSSLRSEDTAVYYCAR TMGEYSYGN YDYWGQGLTLTVSS
YN039VH	QVQLVQSGAEVKKPGSSVKVSKCKAS RGPFSTYA ISWVRQAPGGQGLEWMGRIIPILGIANYAQKFKQGRVTTITADKSTSTAYMELSSLRSEDTAVYYCAR TMGEYSYGN YDYWGQGLTLTVSS

Fig. 9

line IGLV2-14

YN003VL	QSAITQPASVSGSPGQSIITISCTGTSDDVGGYNYVSWYQQHPGKAPKLMIVENSNRPSGVSNRFGSKSGNTASLTISGLQAEADAYVCSSTSS---VFGGGTKLTVLG
YN003VL	QSAITQPASVSGSPGQSIITISCTGTSDDVGGYNYVSWYQQHPGKAPKLMIVENSNRPSGVSNRFGSKSGNTASLTISGLQAEADAYVCSYDSSLSGSVFGGGTKLTVLG
YN035VL	QSAITQPASVSGSPGQSIITISCTGTVSEVGGYNEVSWYQQHPGKAPKLMIVENSNRPSGVSNRFGSKSGNTASLTISGLQAEADAYVCSYDSSLSGSVFGGGTKLTVLG
YN036VL	QSAITQPASVSGSPGQSIITISCTGTVSEVGGYNEVSWYQQHPGKAPKLMIVENSNRPSGVSNRFGSKSGNTASLTISGLQAEADAYVCSYDSSLSGSVFGGGTKLTVLG
YN037VL	QSAITQPASVSGSPGQSIITISCTGTSVVGGYREVSWYQQHPGKAPKLMIVENSNRPSGVSNRFGSKSGNTASLTISGLQAEADAYVCSYDSSLSGSVFGGGTKLTVLG
YN038VL	QSAITQPASVSGSPGQSIITISCTGTRSDVGGYNEVSWYQQHPGKAPKLMIVENSNRPSGVSNRFGSKSGNTASLTISGLQAEADAYVCSYDSSLSGSVFGGGTKLTVLG
YN039VL	QSAITQPASVSGSPGQSIITISCTGTVSDVGGYNEVSWYQQHPGKAPKLMIVENSNRPSGVSNRFGSKSGNTASLTISGLQAEADAYVCSYDSSLSGSVFGGGTKLTVLG

Fig.10

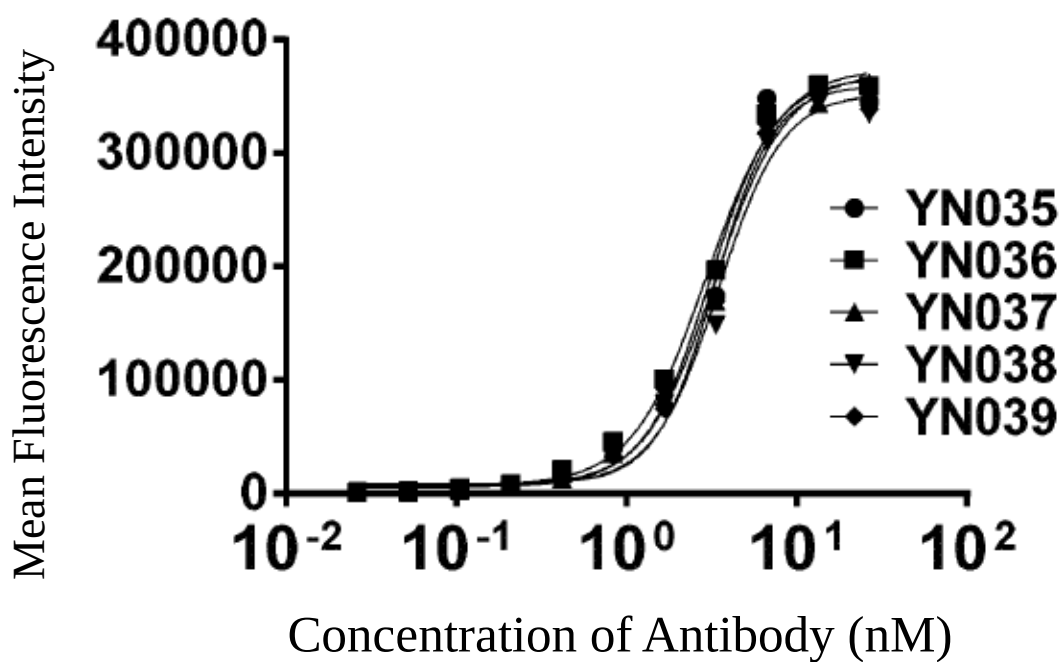


Fig. 11

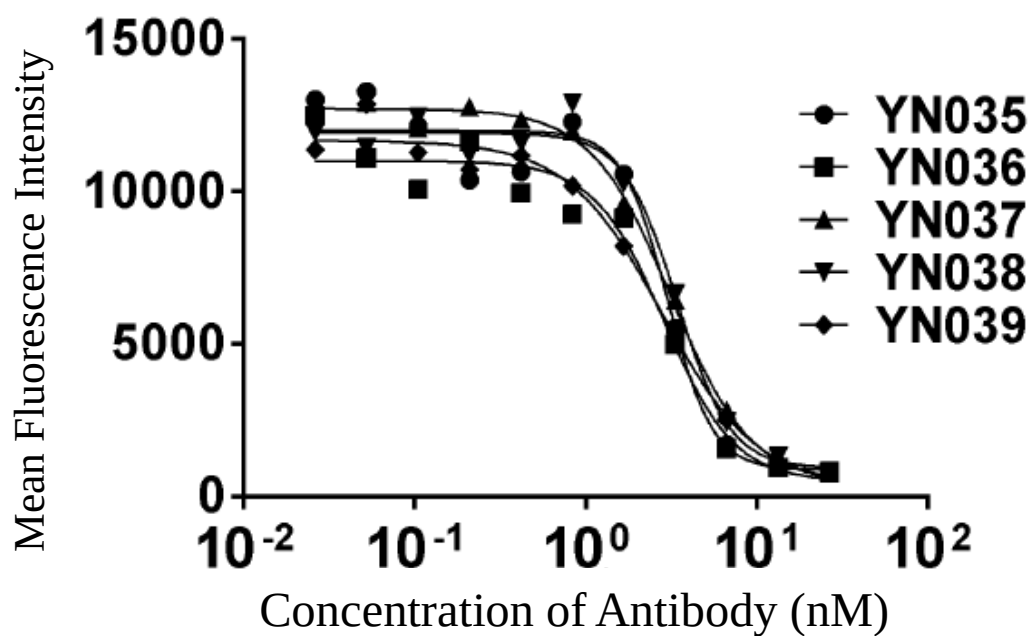


Fig. 12

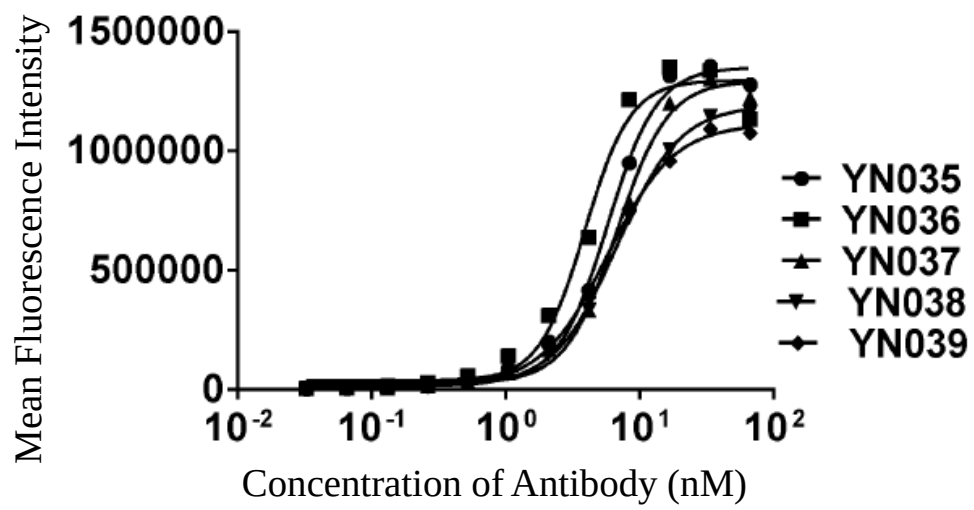


Fig. 13

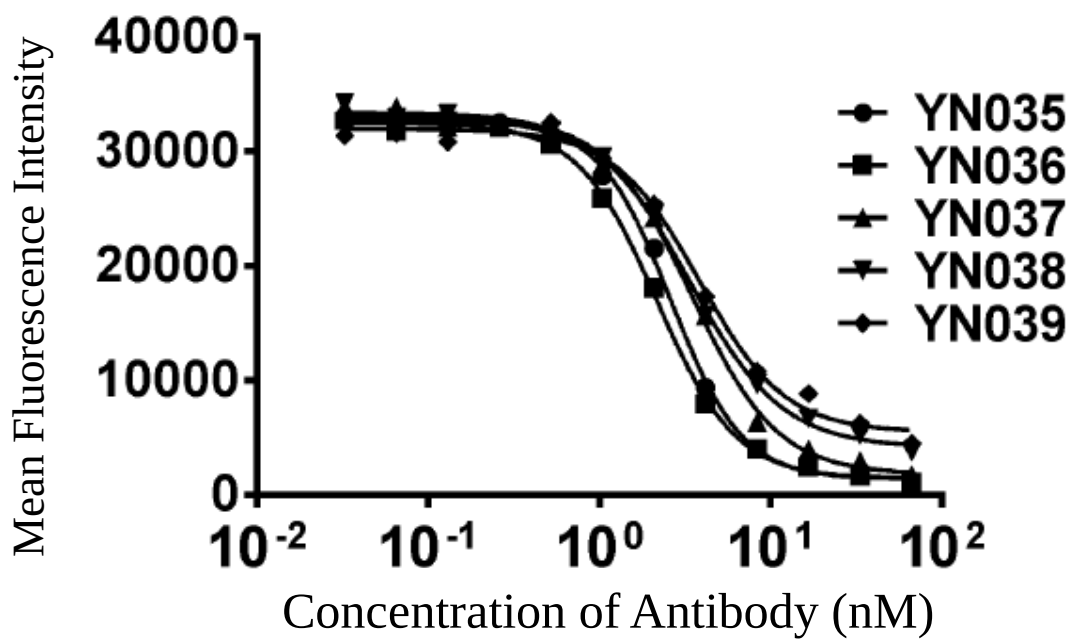


Fig. 14

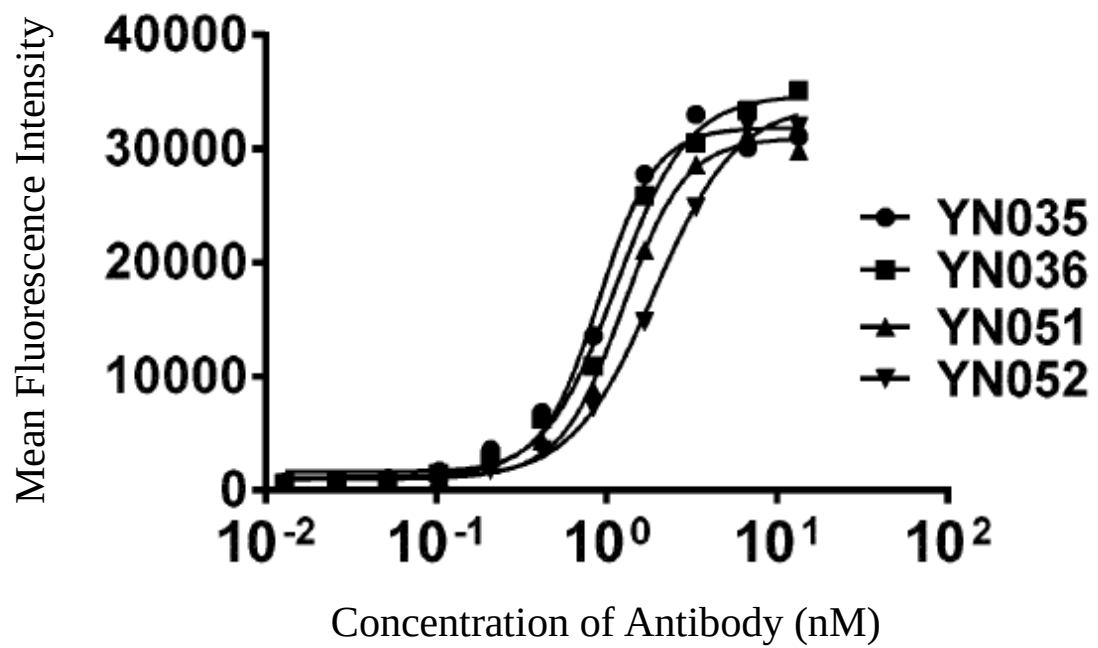


Fig. 15

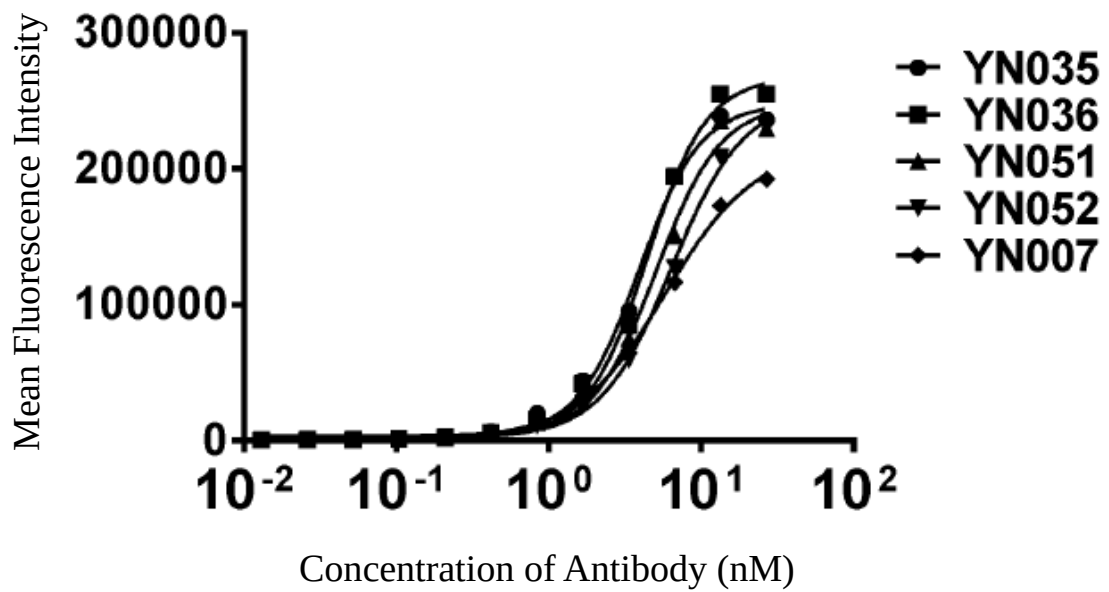


Fig. 16

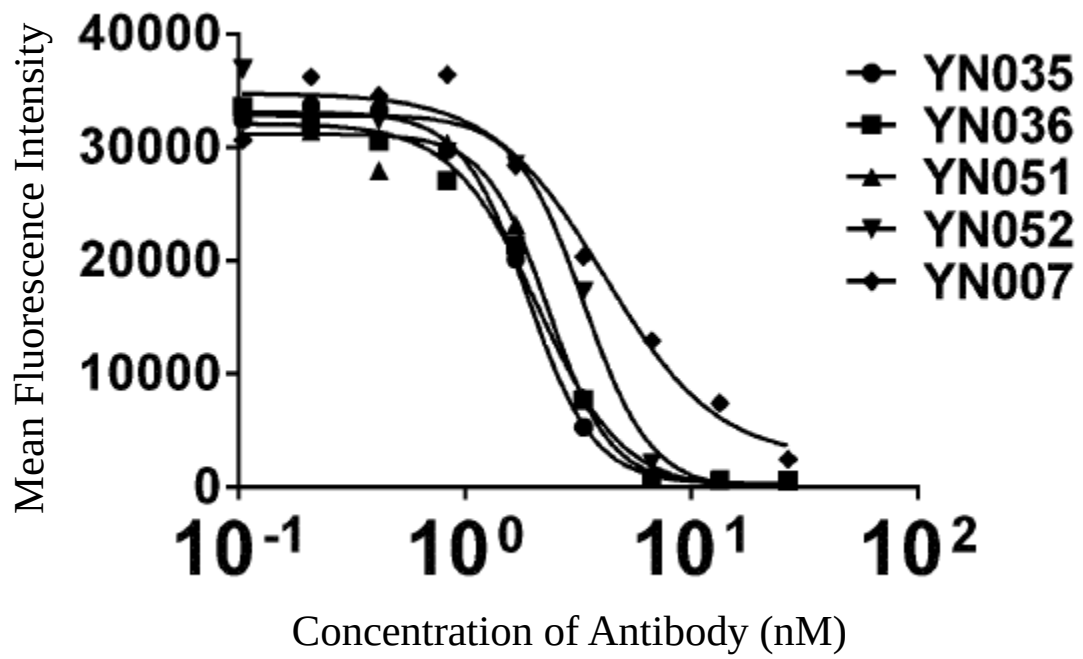


Fig. 17

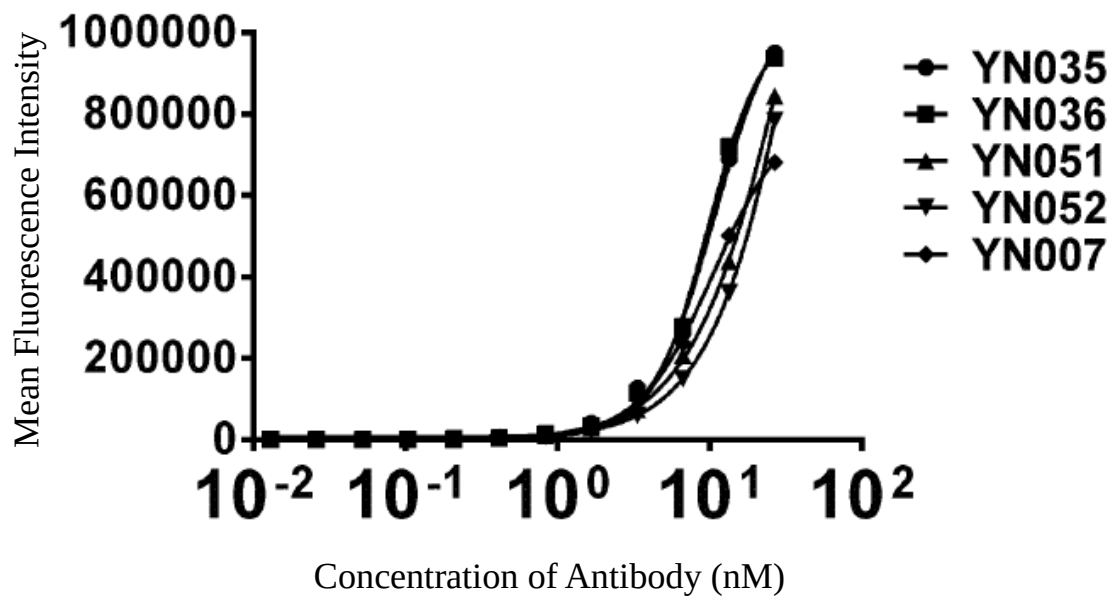


Fig.18

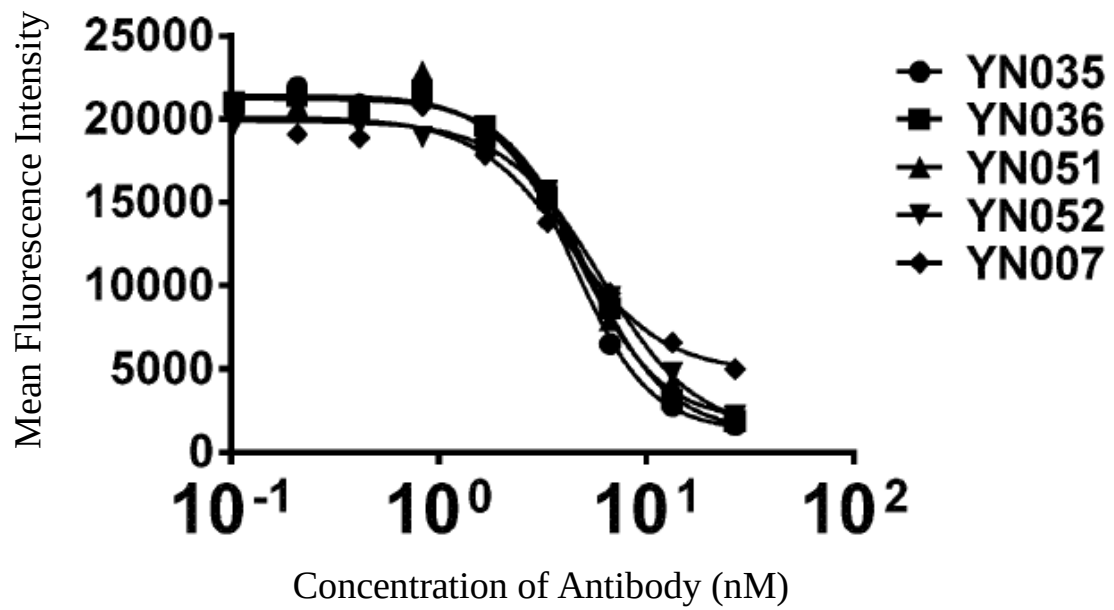


Fig. 19

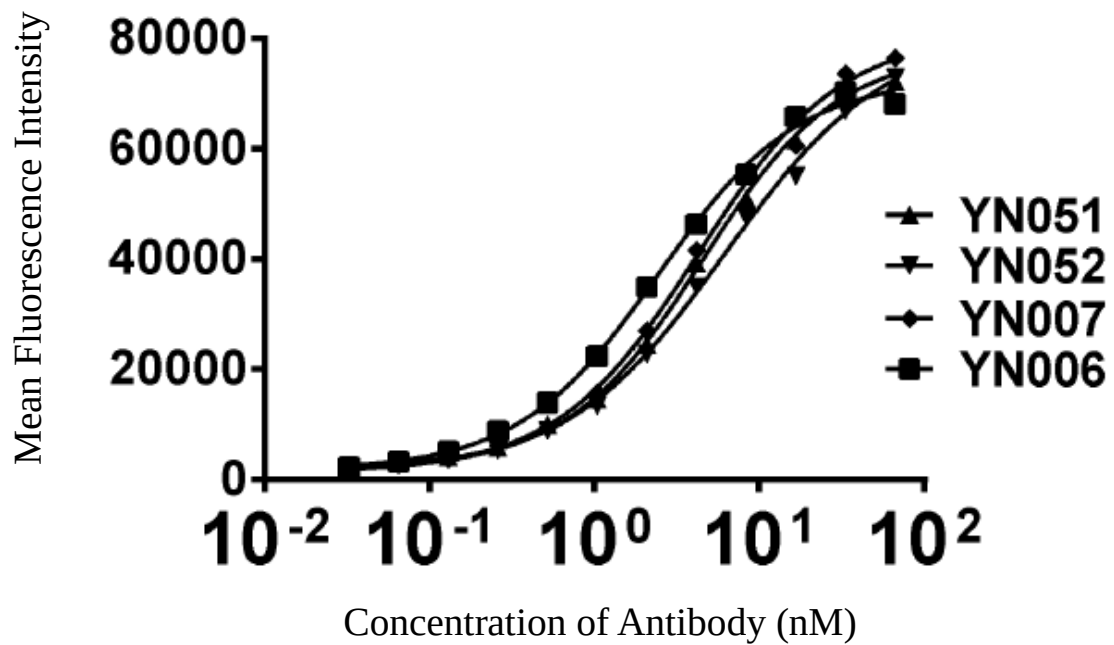


Fig. 20

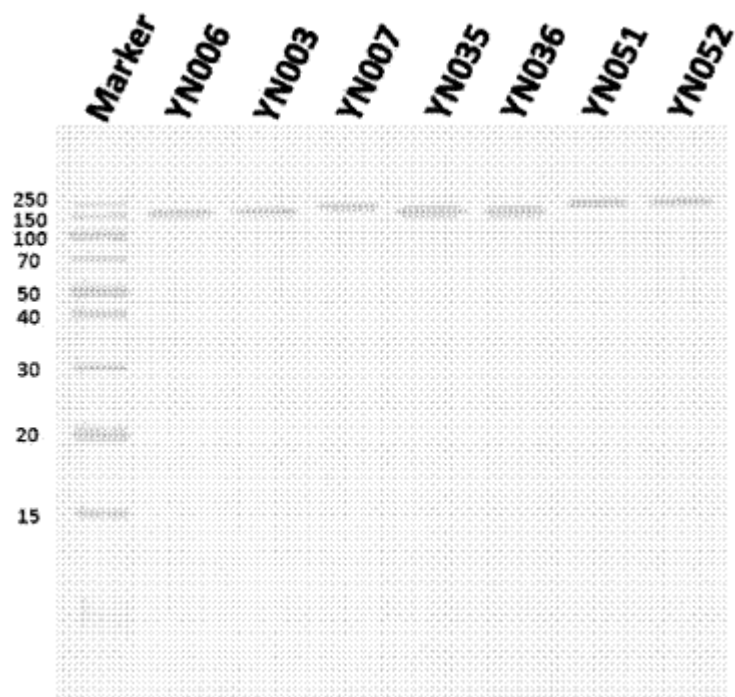


Fig.21a

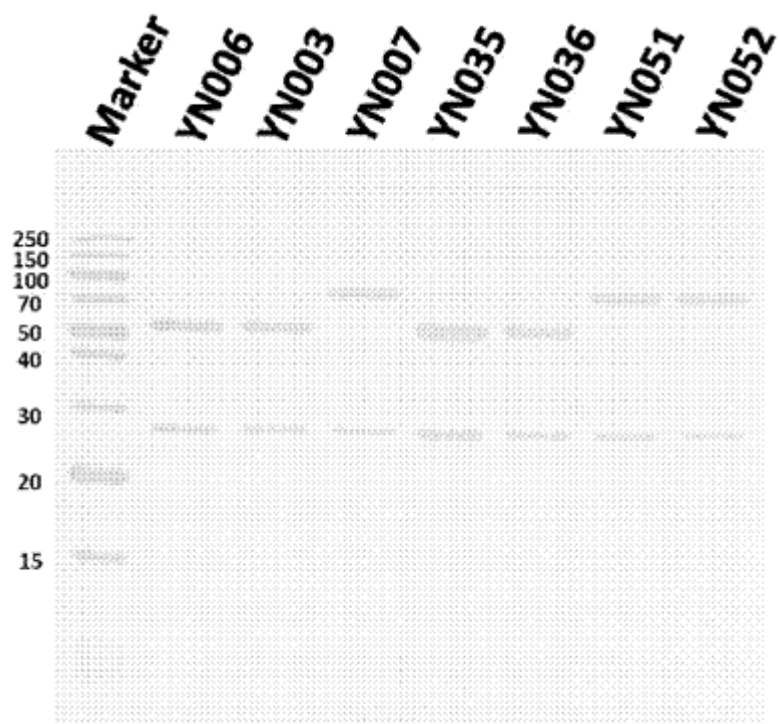


Fig.21b

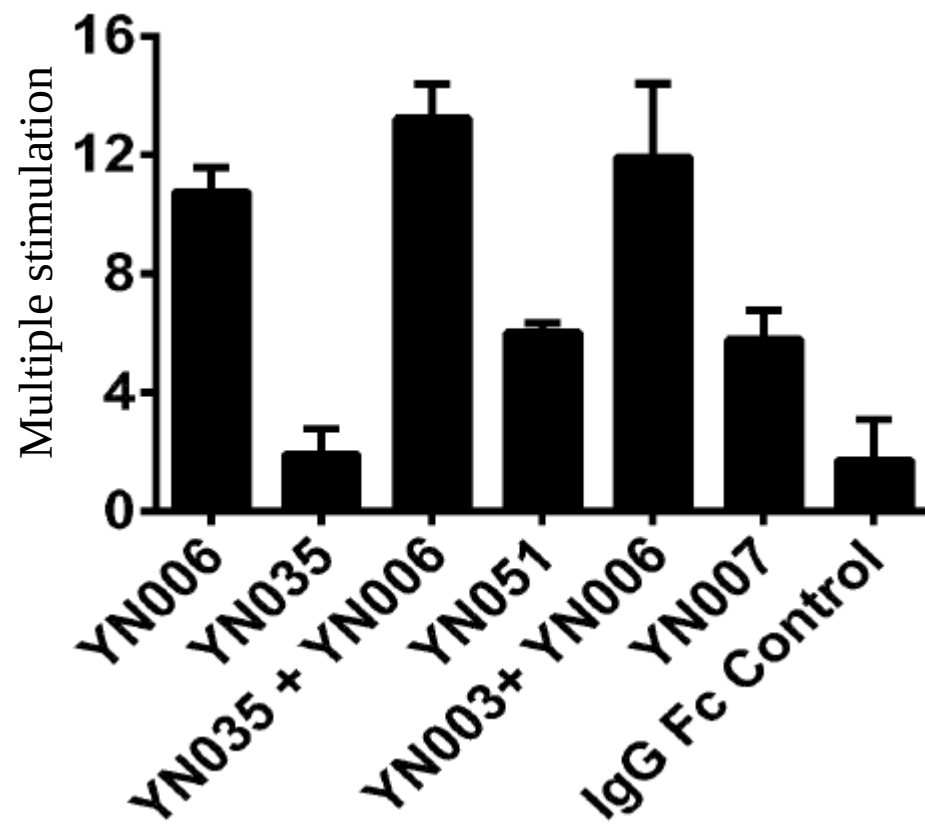


Fig. 22

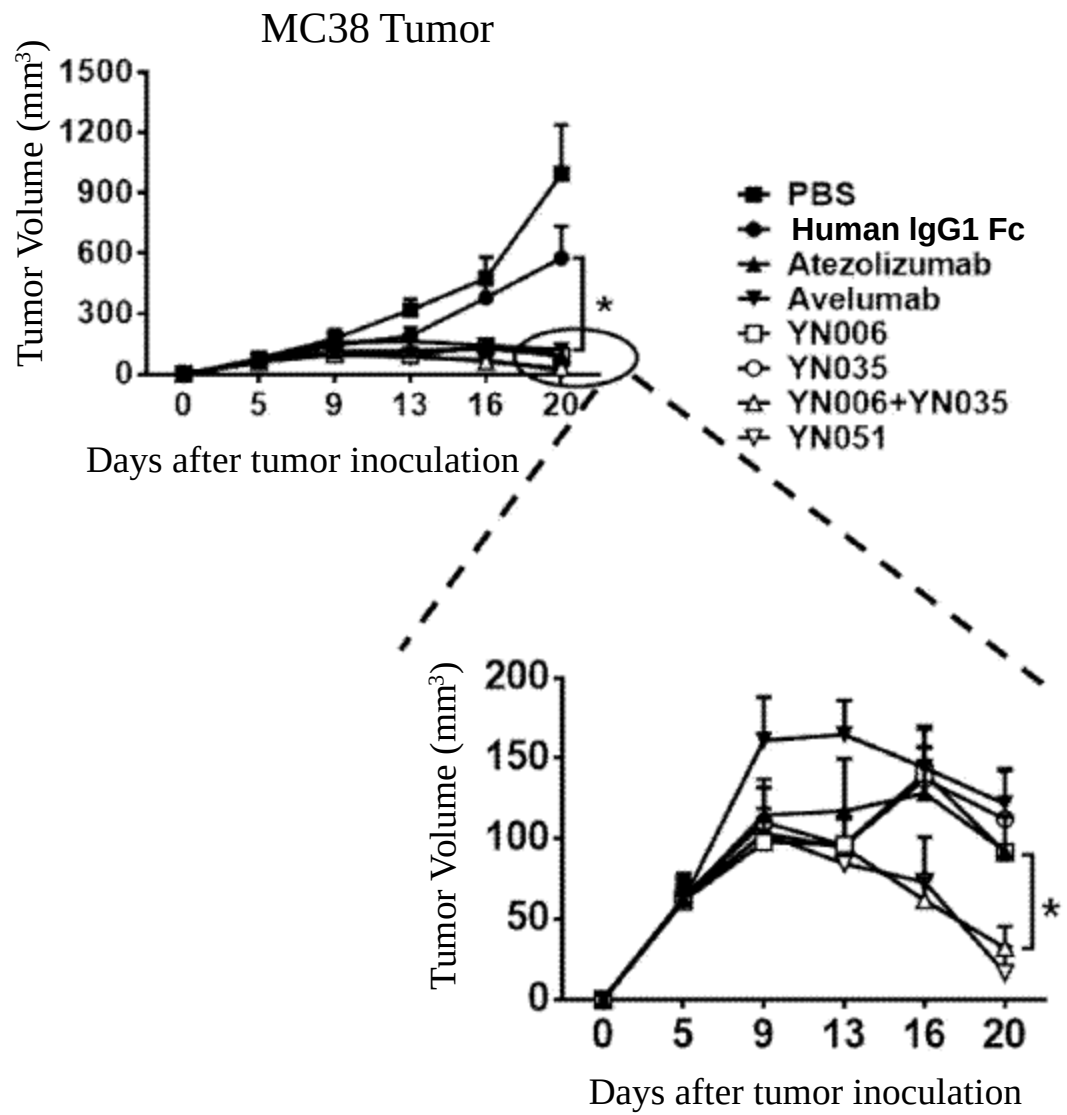


Fig. 23