

FIGURE 1

mAb1 anti-FIX(a) arm CDR- and variable domain sequences and full length sequences								
		SEQ ID NO		SEQ ID NO		SEQ ID NO	SEQ ID NO	
VH	EVQLVESGGGLVQPGRSRLRSCAASGFTFHDYAMHWVRQAPGKGLEWVSGISWRGDIGGYKSVKGRFTISRDNAKNSLYQMNSLRAEDTALYYCVKSYGSGSFYNAFDSWGAGTLVTS	6	DYAMH	3	GISWRGDIGGYV KSVKG	4	SYGSGSFYN AFDS	5
VL	DIQMTASPSTLSASVGDRTVITCRASQSISSWLAWYQQKPGKAPKFLIYKASKLERGTPSRFSGSGSGTEFSLTISSLPDDFATYYCLEYSSYIRTFGGGT KVEIK	11	RASQSISSWL A	8	KASKLER	9	LEYSSYIRT	10
HC	EVQLVESGGGLVQPGRSRLRSCAASGFTFHDYAMHWVRQAPGKGLEWVSGISWRGDIGGYKSVKGRFTISRDNAKNSLYQMNSLRAEDTALYYCVKSYGSGSFYNAFDSWGAGTLVTSSASTKGPSVFPLAPCSRSTSESTAALGCLVKDYFPEPTVSWNSGALTSGVHTFPAVLQSSGLYSLSSVWTPSSSLGTIKTYTCNVDHKPSNTKYDKRVEISKYGPCCPAPAEFLGGPSVFLFPPKPKDITLMISRTPEVTCVWDVDSQEDPEVQFNWYVDGVEVHNAKTKPREEQFNSTYRVSVLTVLHQDWLNGKEYKCKVSNKGLPSSIEKTIKAKGQPREPQAVYTLPPSQEEMTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTTPVLDSGSGFFLYSRLTYDKSRWQEGNVFSCSVMEALHNHYTQKSLSLGLG							7
LC	DIQMTASPSTLSASVGDRTVITCRASQSISSWLAWYQQKPGKAPKFLIYKASKLERGTPSRFSGSGSGTEFSLTISSLPDDFATYYCLEYSSYIRTFGGGT KVEIKRTVAAPSVFIFPPSDEQLKSGTASVVCCLNNFYPREAKVQWKVDNALQSGNSQESVTEQDSKDSYISLSTLTLSKADYEKHKVYACEVTHQGLSSPVTKSFNRGEC							12

"Figure 1 continued"

mAb1 anti-FX(a) arm CDR- and variable domain sequences and full length sequences						
	SEQ ID NO	SEQ ID NO	SEQ ID NO	SEQ ID NO	SEQ ID NO	SEQ ID NO
VH	EVQLVQSGAEVKKPGESLRISCKGSGYFSTSWIVWVRQMPGKGLVWVGQGLTVTVSSASTKGPSTSESTAAALGCLVKDYFPEPTV LQWSSLKASDTAMYYCARLHYNSEEFDVWGQGLTVTVSSASTKGPSTSESTAAALGCLVKDYFPEPTV SWNSGALTSQVHTFPVAVLQSSGLYSLSSVTVPSSSLGKTKYTCNVDHKPSNTKVDKRVESKYGPPCPAPEFLGGP SVFLFPKPKDGLPSSIEKTIKAKGQPREPQVYTLPPSQEEMTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTTPV EYCKVSNKGLPSSIEKTIKAKGQPREPQVYTLPPSQEEMTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTTPV LDSDGSFLLYSKLTVDKSRWQEAGNPFSCVMHEALHNHYTQKSLSLGLG	16	13	14	15	
VL	EIVLTQSPGTLSPGERATLSCRAS QSVSSSYLAWSYQQKPGQAPRLIYG QSSRTRGIPDRFSGSGSGTDFLTIS RLEPEDFAVYYCQQFGDSQLFTFGQ GTKLEIK	21	18	19	20	
HC	EVQLVQSGAEVKKPGESLRISCKGSGYFSTSWIVWVRQMPGKGLVWVGQGLTVTVSSASTKGPSTSESTAAALGCLVKDYFPEPTV LQWSSLKASDTAMYYCARLHYNSEEFDVWGQGLTVTVSSASTKGPSTSESTAAALGCLVKDYFPEPTV SWNSGALTSQVHTFPVAVLQSSGLYSLSSVTVPSSSLGKTKYTCNVDHKPSNTKVDKRVESKYGPPCPAPEFLGGP SVFLFPKPKDGLPSSIEKTIKAKGQPREPQVYTLPPSQEEMTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTTPV EYCKVSNKGLPSSIEKTIKAKGQPREPQVYTLPPSQEEMTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTTPV LDSDGSFLLYSKLTVDKSRWQEAGNPFSCVMHEALHNHYTQKSLSLGLG	16	13	14	15	
LC	EIVLTQSPGTLSPGERATLSCRASQSVSSSYLAWSYQQKPGQAPRLIYGQSSRTRGIPDRFSGSGSGTDFLTISRLEP EDFAVYYCQQFGDSQLFTFGQGTKLEIKRTVAAPSVFIFPPSDEQLKSGTASVWCCLLNFFYPREAKVQWKVDNALQSGNS QESVTEQDSKDSYSTLSSTLTLSKADYEKHKVYACEVTHQGLSSPVTKSFNRGEC	21	18	19	20	22

FIGURE 2

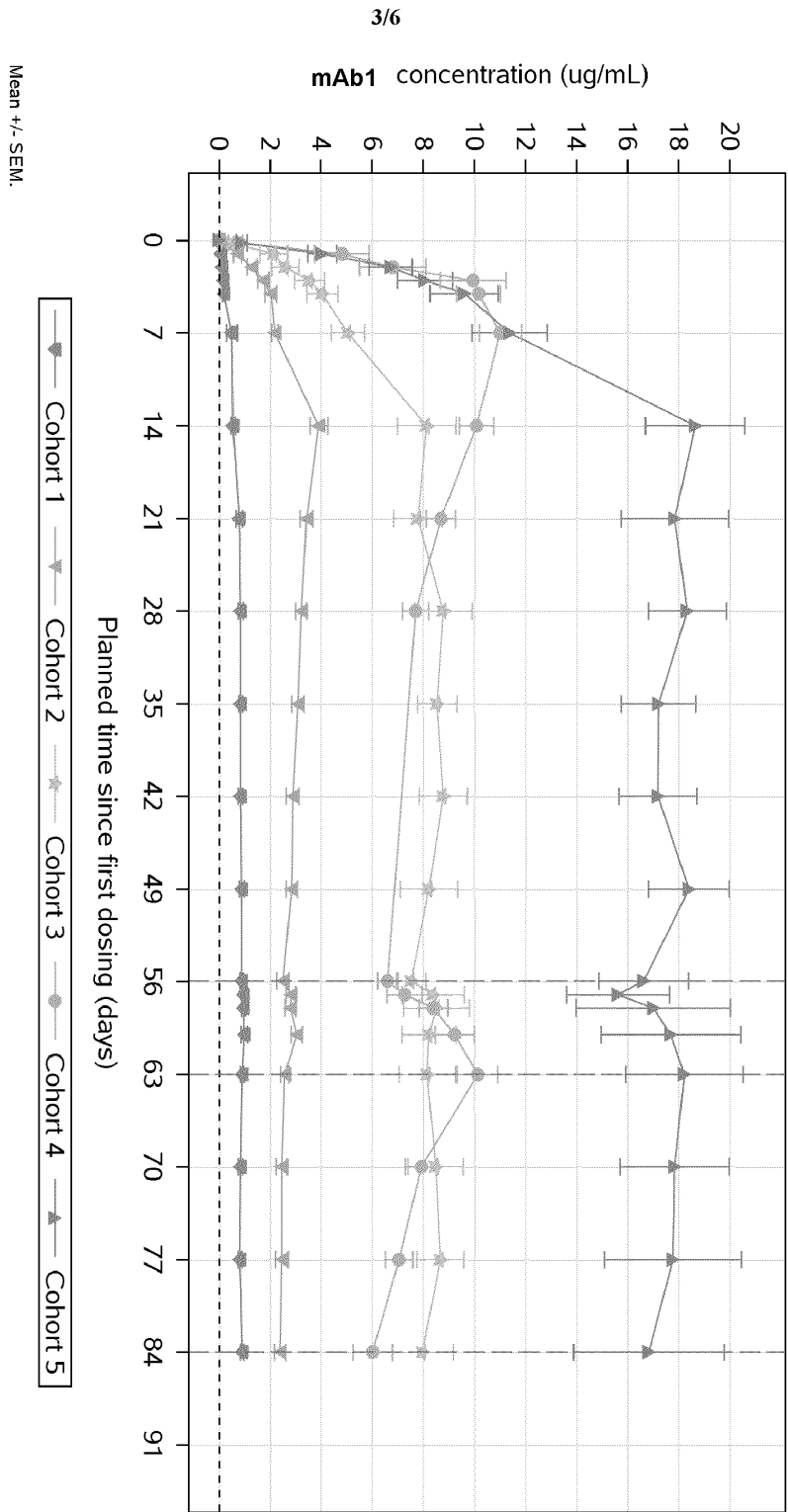


FIGURE 3

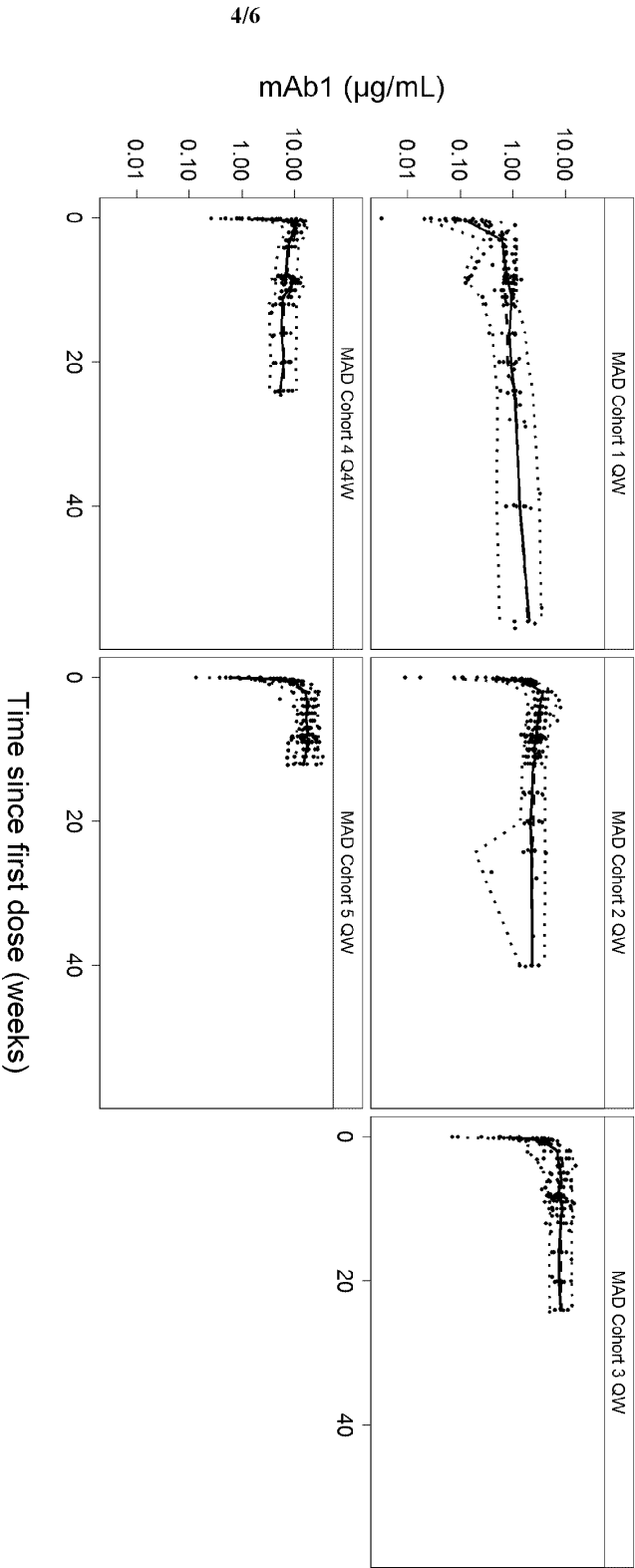


FIGURE 4

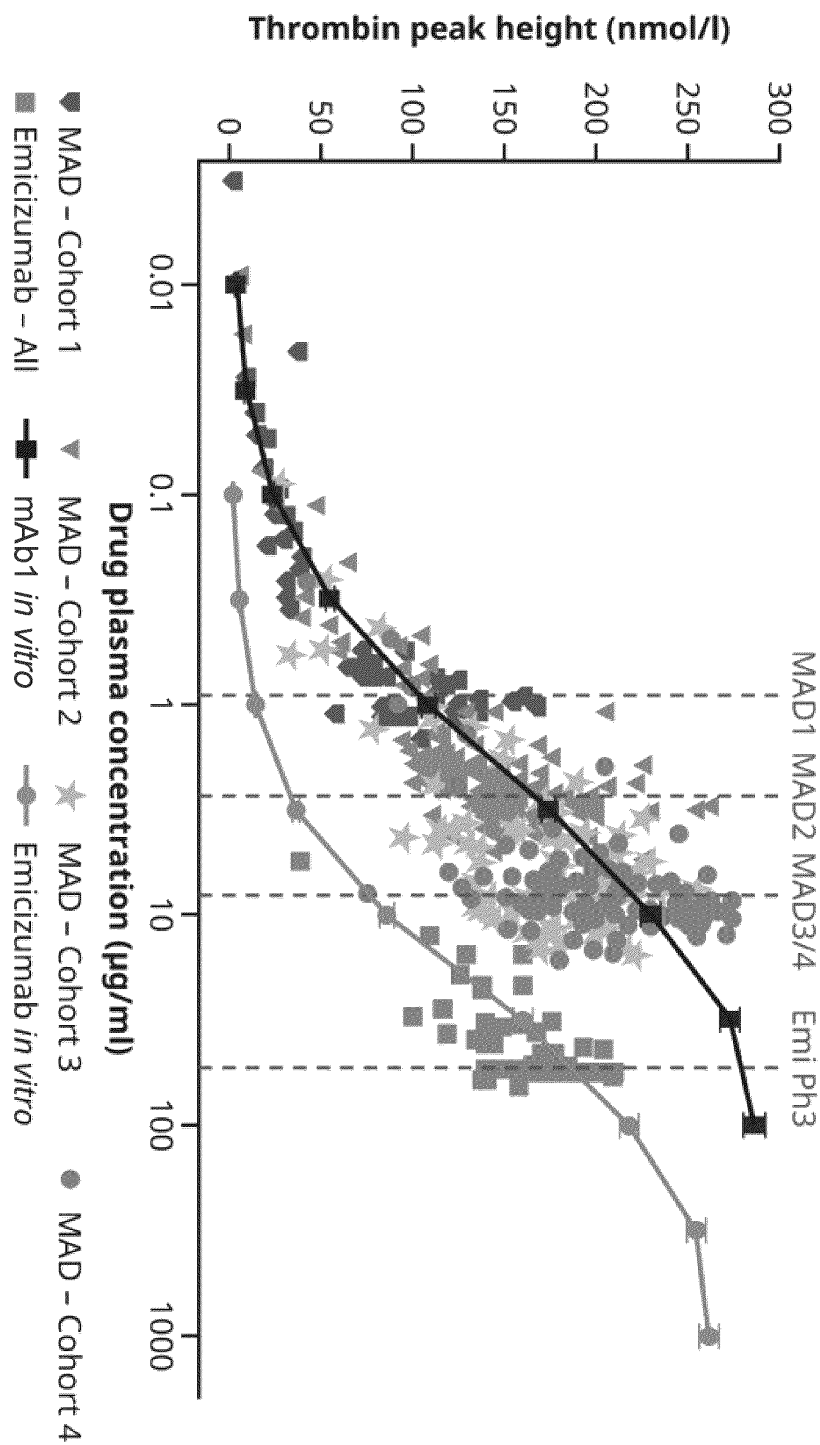


FIGURE 5

