

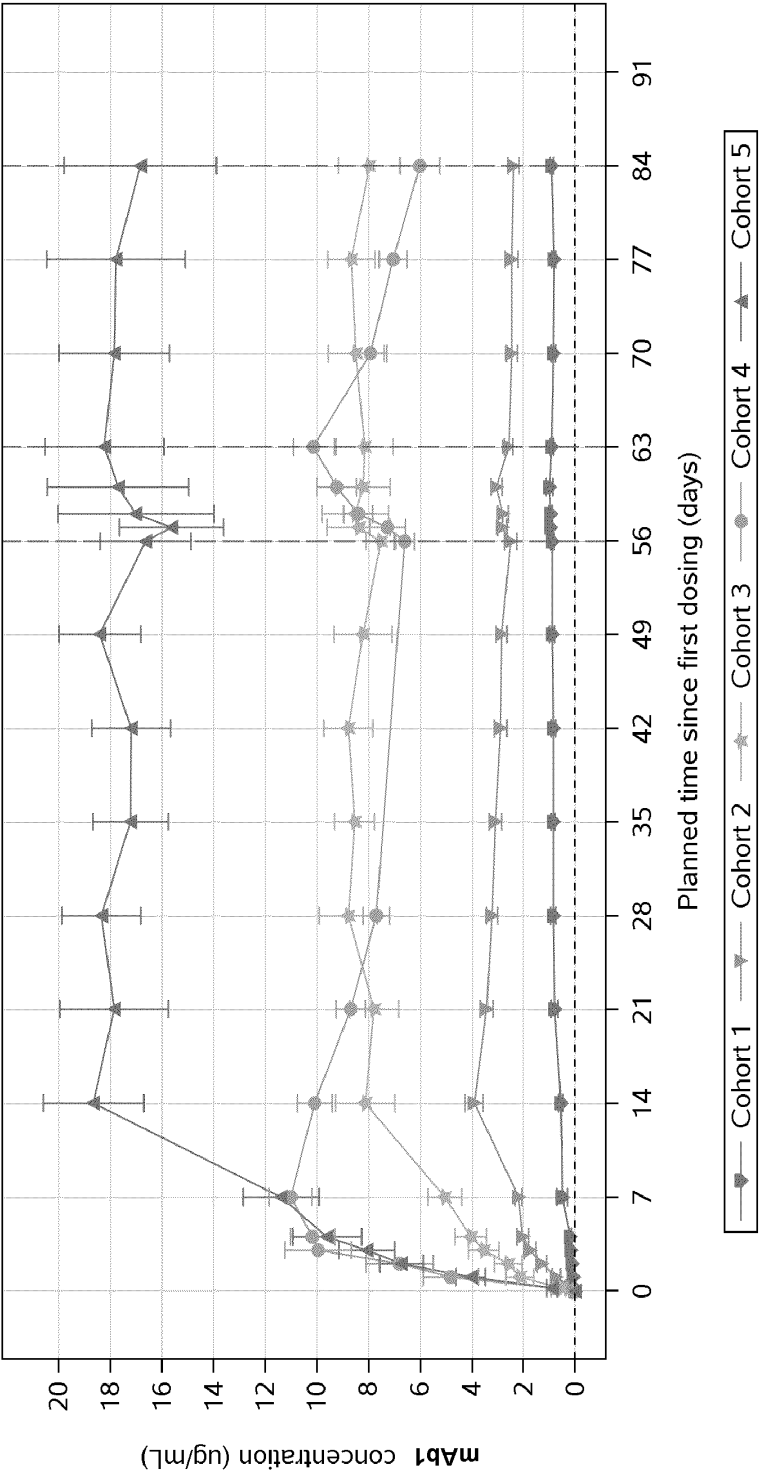
FIGURE 1

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	EVQLVESGGGLVQPGRSLRLSCAA SGFTFHDYAMHWVRQVPGKGLEW VSGISWRGDIGGYVKSVKGRFTISR DNAKNSLYLQMNSLRAEDTALYYCV KSYGSGSFYNAFDSWGQGLTVTS S	6	DYAMH	3	GISWRGDIGGYV KSVKG	SYGSGSFYN AFDS	5
VL	DIQMTQSPSTLSASVGDRVTITCRAS QSISSWLAWYQQKPGKAPKFLYKA SKLERGTSPRFSGSGSGTEFSLTISS LQPDDEFATYYCLEYSSYIRTFGQGT KVEIK	11	RASQSISSWL A	8	KASKLER	LEYSSYIRT	10
HC	EVQLVESGGGLVQPGRSLRLSCAA SGFTFHDYAMHWVRQVPGKGLEW VSGISWRGDIGGYVKSVKGRFTISR DNAKNSLYLQMNSLRAEDTALYYCV KSYGSGSFYNAFDSWGQGLTVTS S	6	DYAMH	3	GISWRGDIGGYV KSVKG	SYGSGSFYN AFDS	5
LC	DIQMTQSPSTLSASVGDRVTITCRAS QSISSWLAWYQQKPGKAPKFLYKA SKLERGTSPRFSGSGSGTEFSLTISS LQPDDEFATYYCLEYSSYIRTFGQGT KVEIK	11	RASQSISSWL A	8	KASKLER	LEYSSYIRT	10

"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
VH	EVQLVQSGAEVKKPGESLRISCKGSGYFSTSWIVWVRQMPGKGLEWMGIDPDSFTSYSPSFQGHVTISADK SISTAYLQWSSLKASDTAMYYCARL HYNSEEFDVWGQGLTVTVSS	16	TSWIV	13	MIDPDSFTSYSP SFQG	14	LHYNSEEFD V	15
VL	EIVLTQSPGTLSPGERATLSCRASQSVSSSYLAWYQQKPGQAPRLIYGQSSRTRGIPDRFSGSGSGTDFLTISRLEPEDFAVYYCQFGDSQLFTFGQ GTKLEIK	21	RASQSVSSSYLA	18	GQSSRTR	19	QQFGDSQLF T	20
HC	EVQLVQSGAEVKKPGESLRISCKGSGYFSTSWIVWVRQMPGKGLEWMGIDPDSFTSYSPSFQGHVTISADKSI STAYLQWSSLKASDTAMYYCARLHYNSEEFDVWGQGLTVTVSSASTKGPSTSESTAAALGCLVKDYFPEPTV SWNSGALTSGVHTFPAVLQSSGLYSLSSVTVPPSSSLGKTKYTCNVDHKPSNTKVDKRVESKYGPPCPPCPAPEFLGGP SVFLFPPKPKDTLMISRTPEVTCWVDVSEQEDPEVQFNWYVDGVEVHNAKTKPREEQFNSTYRVWSVLTVLHQQDWLNGK EYKCKVSNKGLPSSIEKTIKAKGQPREPQVYTLPPSQEEMTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTTPV LDSDGSFLLYSKLTVDKSRWQEAGNPFSCVMHEALHNHYTQKSLSLGLG							17
LC	EIVLTQSPGTLSPGERATLSCRASQSVSSSYLAWYQQKPGQAPRLIYGQSSRTRGIPDRFSGSGSGTDFLTISRLEP EDFAVYYCQFGDSQLFTFGQGTKLEIKRTVAAPSVFIFPPSDEQLKSGTASVWCCLLNFFYPREAKVQWKVDNALQSGNS QESVTEQDSKDSITYLSSTLTLSKADYEKHKVYACEVTHQGLSSPVTKSFNRGEC							22

FIGURE 2



Mean +/- SEM.

FIGURE 3

