

CDR-H1			CDR-H2			CDR-H3		
SEQ6	GYILSPNF	8	SEQ55	-IESDGGST-	8	SEQ98	-----AVLFFWWRQYGV-----	13
SEQ27	GYILSPNF	8	SEQ78	-IESDGGST-	8	SEQ121	-----AVLFFWWRQYGV-----	13
SEQ17	GRYSPNF	8	SEQ44	-IESDGGST-	8	SEQ95	-----VIDAWAKHIF-----NY	13
SEQ38	GRYSPNF	8	SEQ45	-IESDGGST-	8	SEQ92	-A-----GAWHSLTVPHY-----	13
SEQ7	GRYSPNF	8	SEQ47	-IESDGGST-	8	SEQ113	AA-----GAWHSLTVPHY-----	14
SEQ29	GRYSPNF	8	SEQ48	-IESDGGST-	8	SEQ109	-----AVFFWGLGGLAYPSDETHFYYSY	22
SEQ32	GRYSPNF	8	SEQ49	-IESDGGST-	8	SEQ108	AA---WPYVQWATFLAHNY-----	17
SEQ38	GRYSPNF	8	SEQ51	-IESDGGST-	8	SEQ101	-----RAZEDVWNNY-----	11
SEQ6	GRYSPNF	8	SEQ52	-IESDGGST-	8	SEQ123	-----AIEDVWNNY-----	10
SEQ15	GRYSPNF	8	SEQ56	-IESDGGST-	8	SEQ91	-A-----KNTAKSLXIAHNY-----	16
SEQ37	GRYSPNF	8	SEQ58	-IESDGGST-	8	SEQ99	-A-----KNTAL---TWNNY-----	13
SEQ18	GRYSPNF	8	SEQ59	-IESDGGST-	8	SEQ116	AA---KNTAL---TWNNY-----	14
SEQ22	GRYSPNF	8	SEQ60	-IESDGGST-	8	SEQ98	-ASIRGIT-----IEHNNKYV-----	16
SEQ16	GRYSPNF	8	SEQ62	-IESDGGST-	8	SEQ112	AAKIRGIT-----IEHNNKYV-----	17
SEQ11	GRYSPNF	8	SEQ64	-IESDGGST-	8	SEQ107	-AKIRGIT-----IEHNNKYV-----	16
SEQ34	GRYSPNF	8	SEQ65	-IESDGGST-	8	SEQ126	-AKIRGIT-----IEHNNKYV-----	16
SEQ3	GRYSPNF	8	SEQ67	-IESDGGST-	8	SEQ105	AVTETAIT-----LPOVCKHNY-----	17
SEQ31	GRYSPNF	8	SEQ68	-IESDGGST-	8	SEQ119	AVTETAIT-----LPOVCKHNY-----	17
SEQ31	GRYSPNF	8	SEQ69	-IESDGGST-	8	SEQ133	AAQAVNYV---VTDHCKHNY-----	17
SEQ36	GRYSPNF	8	SEQ70	-IESDGGST-	8	SEQ125	-AGVNYV---VTDHCKHNY-----	16
SEQ43	GRYSPNF	8	SEQ73	-IESDGGST-	8	SEQ107	AAQVSPFI---GNNHCKHNY-----	17
SEQ4	GRYSPNF	8	SEQ76	-IESDGGST-	8	SEQ87	-AAQVSPFI---GNNHCKHNY-----	16
SEQ30	GRYSPNF	8	SEQ80	-IESDGGST-	8	SEQ99	AAQVSPFI---GNNHCKHNY-----	17
SEQ8	GRYSPNF	8	SEQ82	-IESDGGST-	8	SEQ110	AAQVSPFI---GNNHCKHNY-----	17
SEQ30	GRYSPNF	8	SEQ83	-IESDGGST-	8	SEQ111	AAQVSPFI---GNNHCKHNY-----	17
SEQ14	GRYSPNF	8	SEQ84	-IESDGGST-	8	SEQ136	---AAQVSPFI---GNNHCKHNY-----	14
SEQ1	GRYSPNF	8	SEQ85	-IESDGGST-	8	SEQ104	-KVVVSPFI---GNNHCKHNY-----	20
SEQ24	GRYSPNF	8	SEQ84	EXNSYACTY	10	SEQ100	AAQVSPFI---GNNHCKHNY-----	14
SEQ5	GRYSPNF	8	SEQ77	EXNSYACTY	10	SEQ108	AAQVSPFI---GNNHCKHNY-----	14
SEQ2	GRYSPNF	8	SEQ43	EXNSYACTY	10	SEQ94	-VME-----NHIPLOFY-----	13
SEQ26	GRYSPNF	8	SEQ41	-IESDGGST-	8	SEQ126	-AAST-----TFELHNY-----Y-----	11
SEQ41	GRYSPNF	8	SEQ46	EXNSYACTY	10	SEQ93	---AAST-----TFELHNY-----	12
SEQ42	GRYSPNF	8	SEQ71	EXNSYACTY	10	SEQ115	---AAST-----TFELHNY-----	13
SEQ15	GRYSPNF	8	SEQ74	EXNSYACTY	10	SEQ113	-AAST-----TFELHNY-----	13
SEQ33	GRYSPNF	8	SEQ79	EXNSYACTY	10	SEQ124	---AAST-----TFELHNY-----	12
SEQ40	GRYSPNF	8	SEQ66	EXNSYACTY	10	SEQ89	---AAST-----TFELHNY-----	10
SEQ12	GRYSPNF	8	SEQ47	-IESDGGST-	8	SEQ114	-AAST-----TFELHNY-----	11
SEQ35	GRYSPNF	8	SEQ43	-IESDGGST-	8	SEQ117	-AAST-----TFELHNY-----	11
SEQ13	GRYSPNF	8	SEQ50	EXNSYACTY	10	SEQ122	---AAST-----TFELHNY-----	10
SEQ3	GRYSPNF	8	SEQ72	EXNSYACTY	10	SEQ128	---AAST-----TFELHNY-----	10
SEQ10	GRYSPNF	8	SEQ75	EXNSYACTY	10	SEQ95	-AAST-----TFELHNY-----	11
SEQ21	GRYSPNF	8	SEQ81	EXNSYACTY	10	SEQ97	-AAST-----TFELHNY-----	11
SEQ25	GRYSPNF	8	SEQ68	-IESDGGST-	8	SEQ120	---AAST-----TFELHNY-----	10

FIG. 1

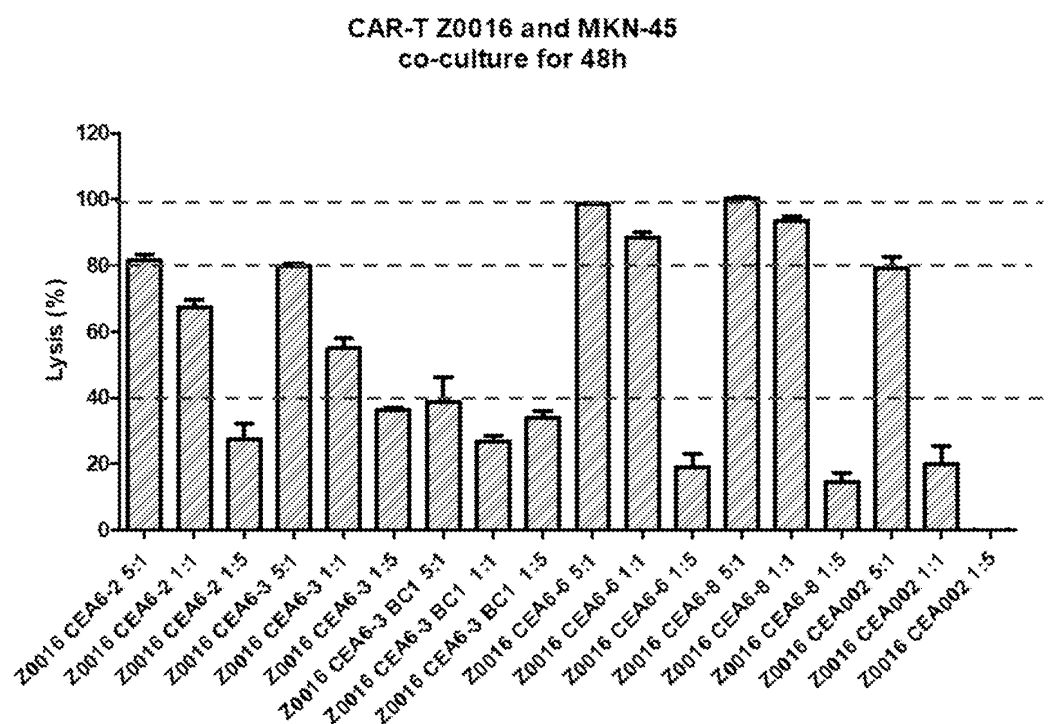


FIG. 2A

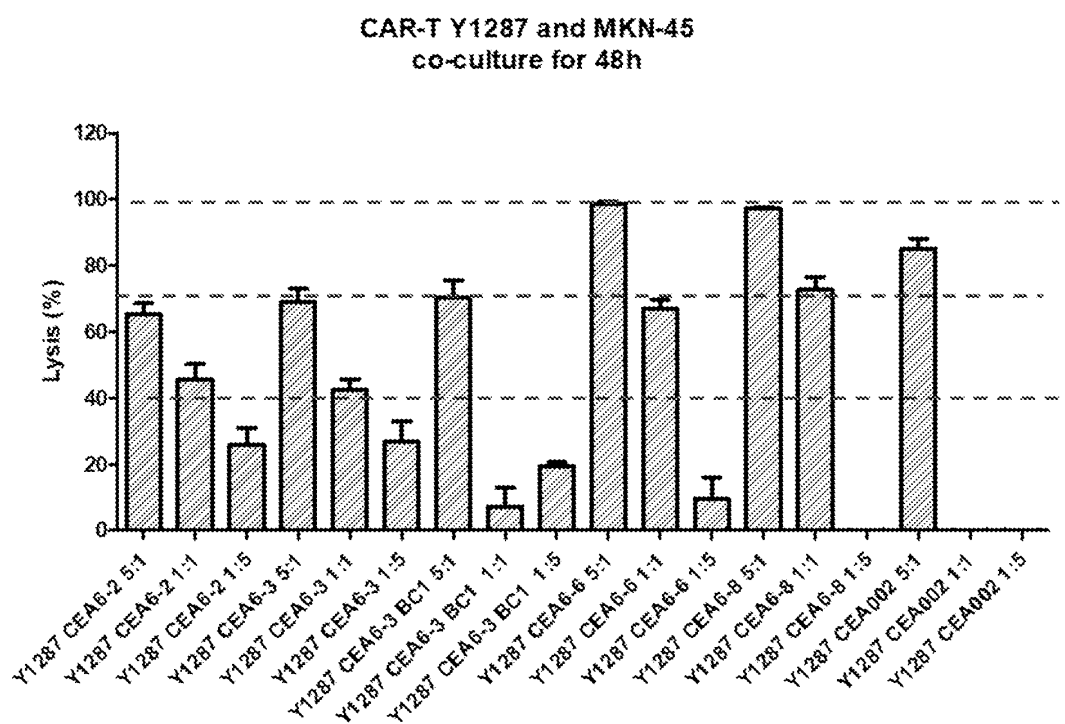


FIG. 2B

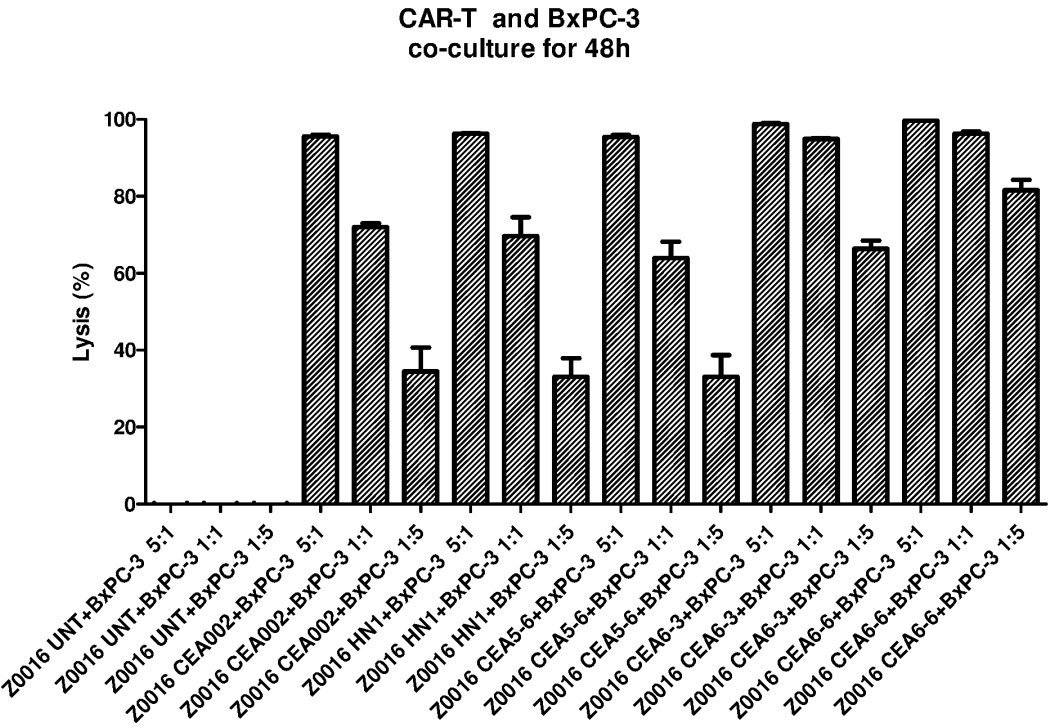


FIG. 2C

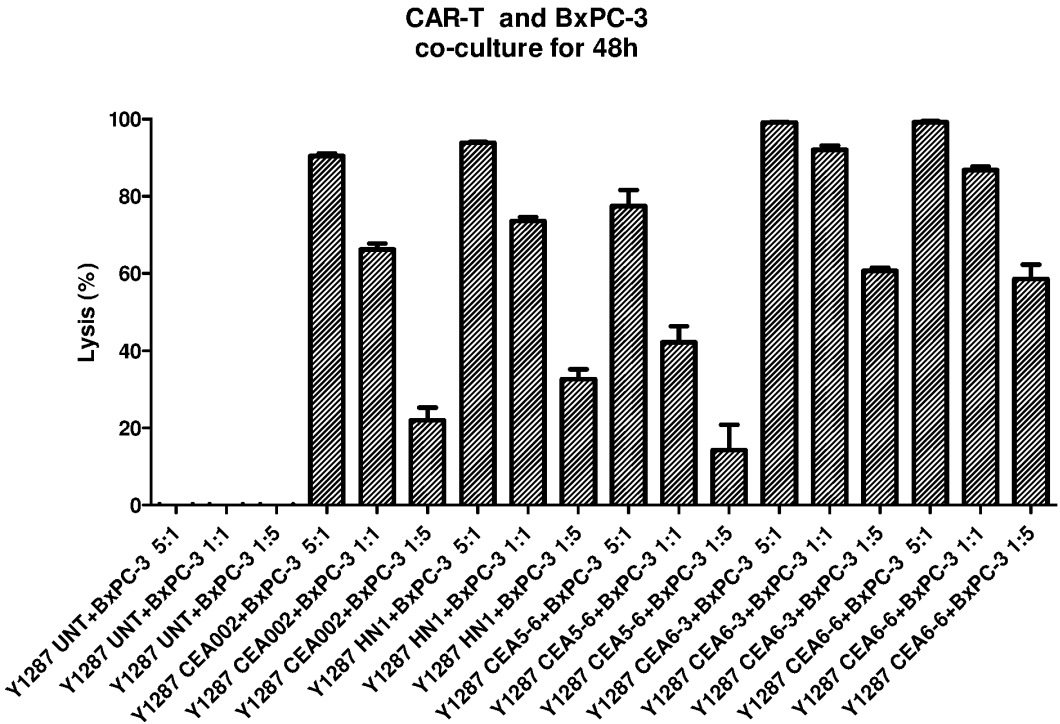


FIG. 2D

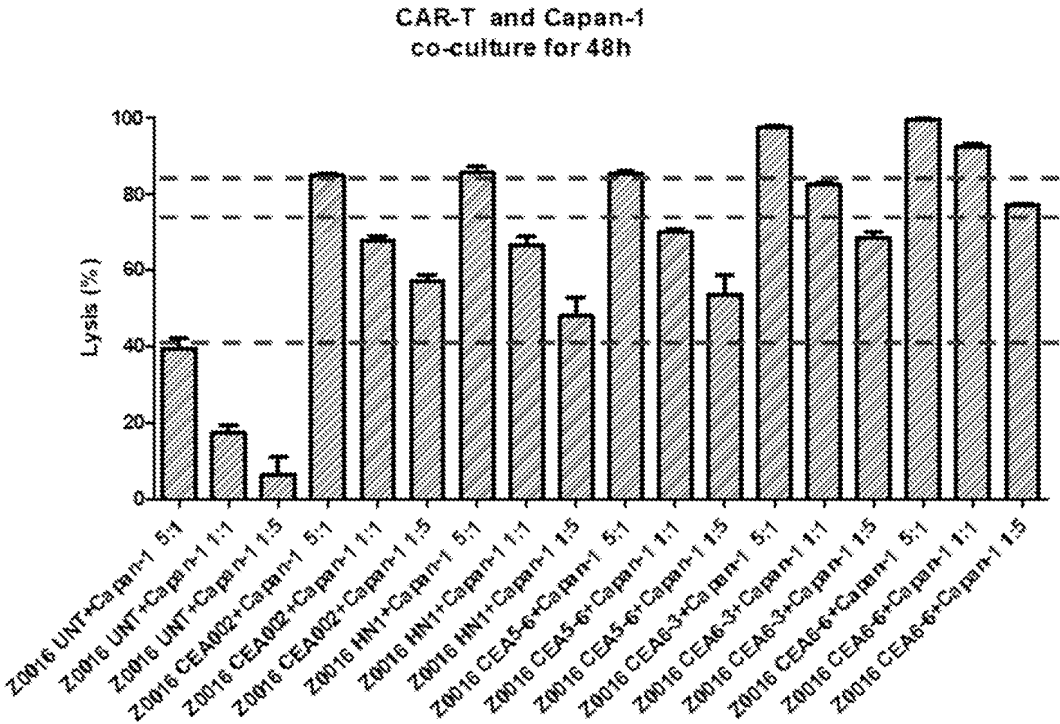


FIG. 2E

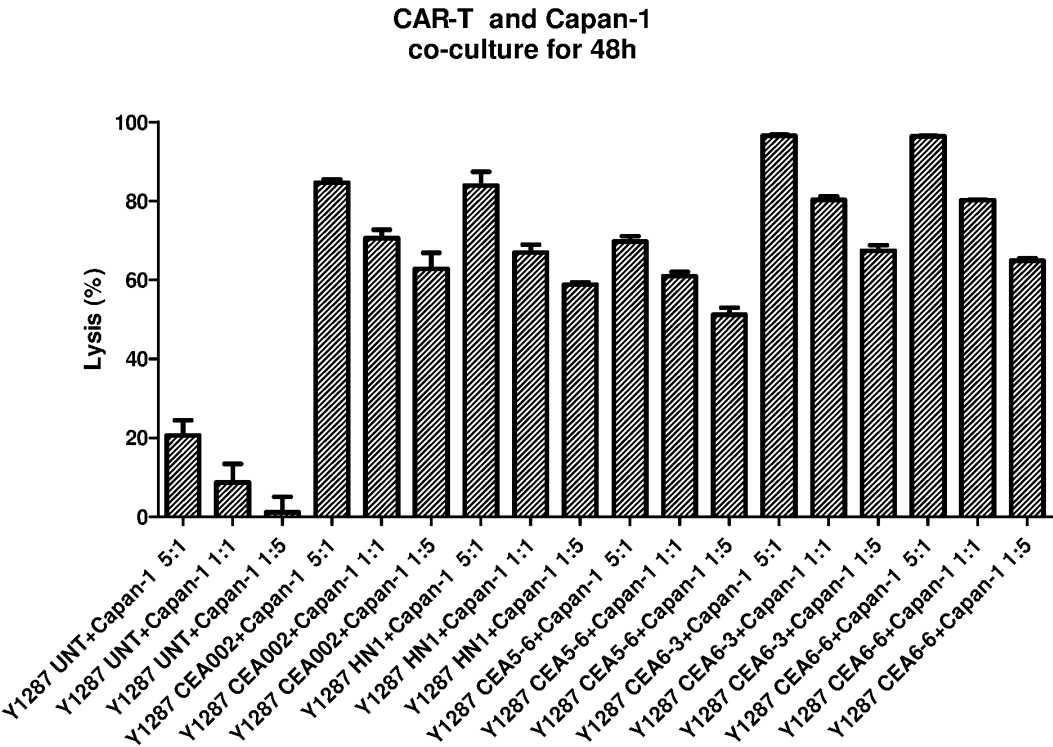
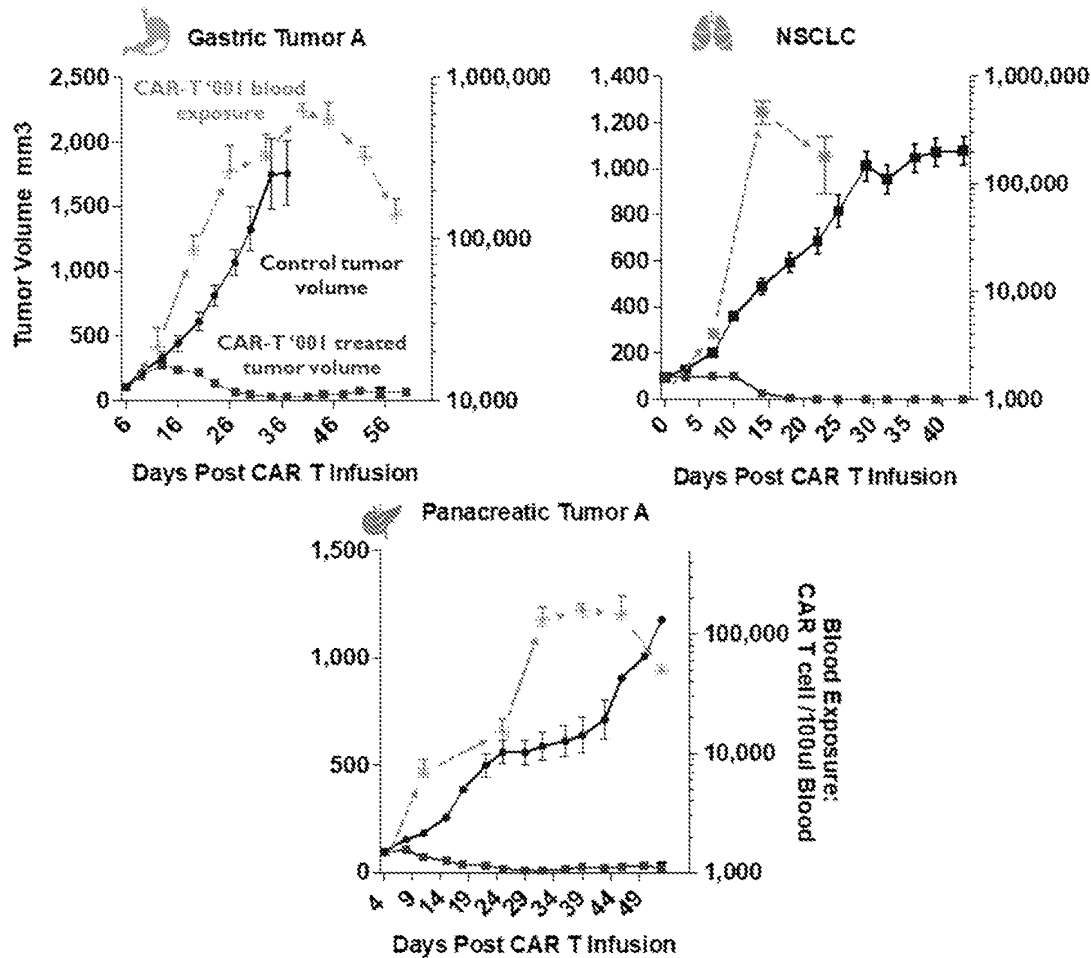


FIG. 2F

Xenograft CDX Models



Note: each tumor type study included NSG mice (n=5)

FIG. 3A

Pancreatic Orthotopic model

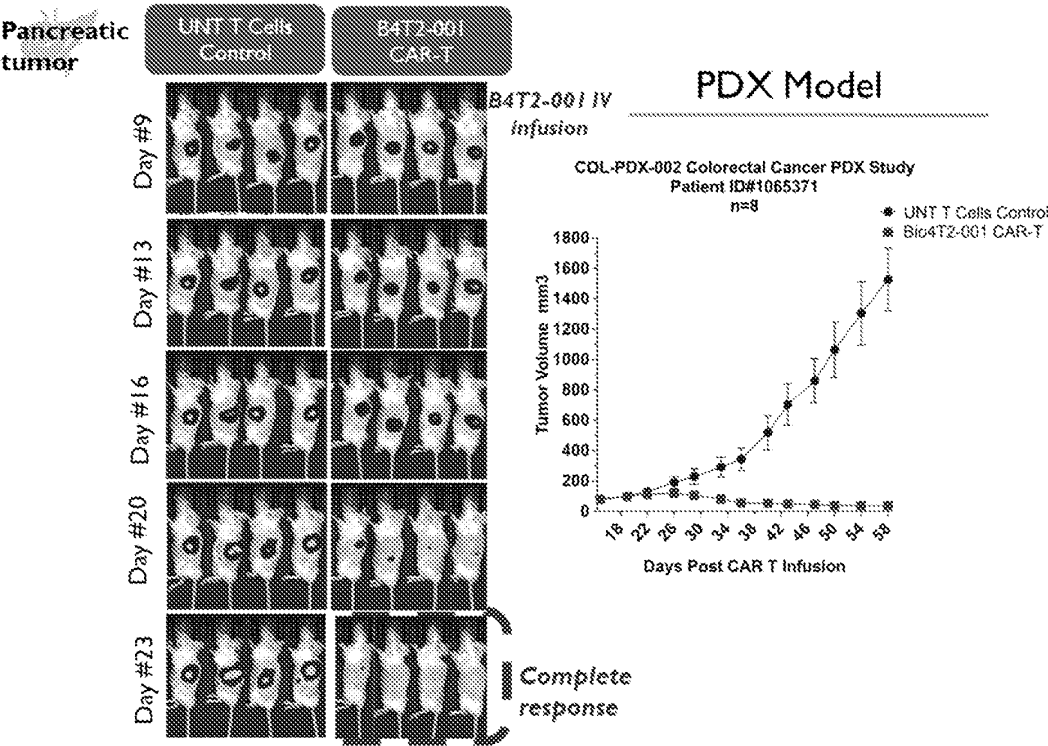


FIG. 3B

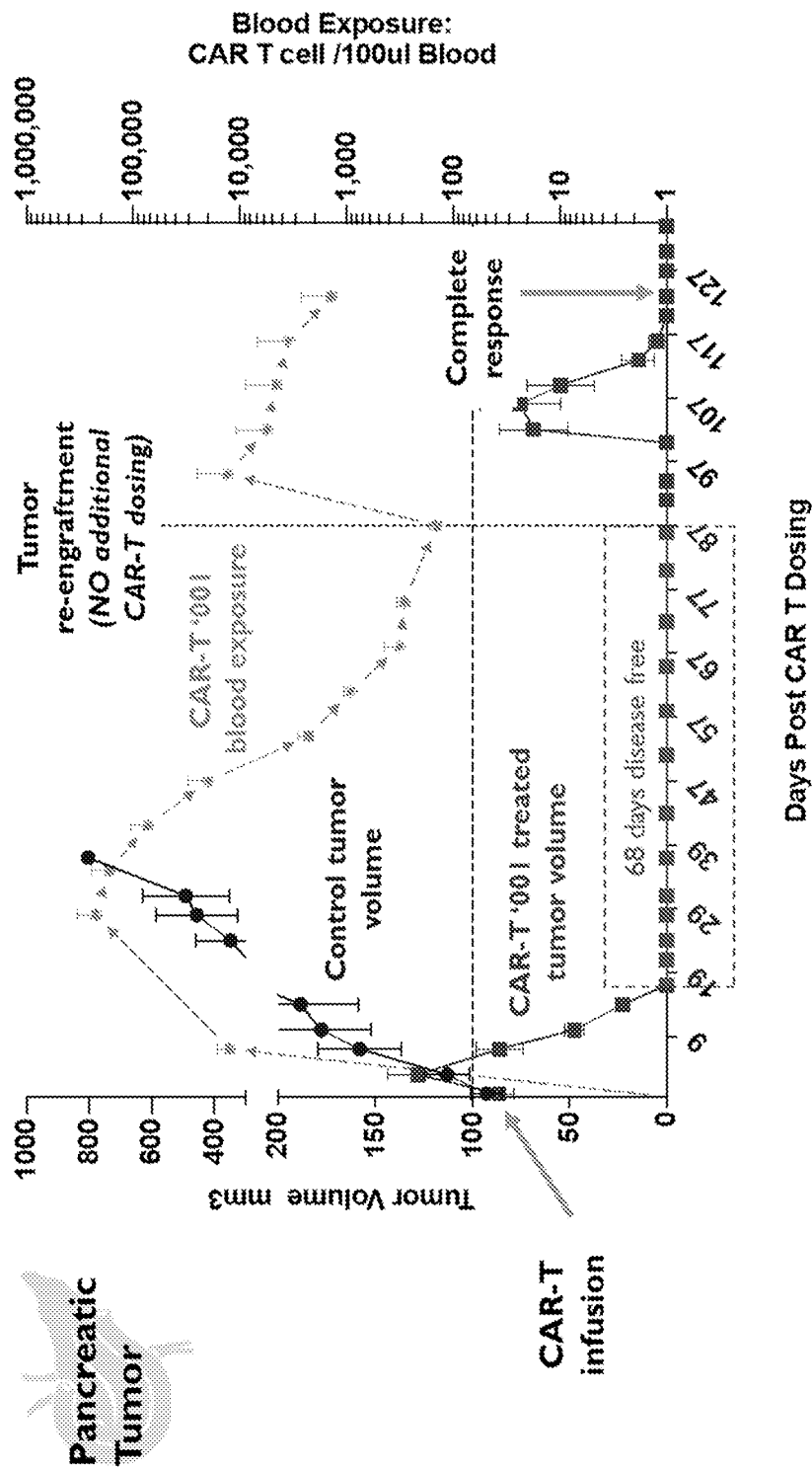


FIG. 4