

FIG. 1.**(1) Wild Type IL12B without signal (IL12B) Amino Acids (SEQ ID NO: 1):**

IWELKKDVYVVELDWYPDAPGEMVVLTCDTPEEDGITWTLDQSSEVLGSGKTLTIQVKEFGDAGQYTCHKGGEV
LSHSLLLLHKKEDGIWSTDILKDQKEPKNKTFLRCEAKNYSGRFTCWWLTTISTDLTFSVKSSRGSSDPQGVTC
GAATLSAERVVRGDNKEYEYSVEQCEDSACPAAEESLPIEVMDAVHKLKYENYTSFFIRDI IKPDPPKNLQLK
PLKNSRQVEVSWEYPDTWSTPHSYFSLTFCVQVQGKSKREKKDRVFTDKTSATVICRKNASISVRAQDRYSSS
WSEWASVPCS

(2) Wild Type IL12B without signal (IL12B) Nucleic Acids (SEQ ID NO: 2):

ATATGGGAAGCTGAAGAAAGATGTTTATGTCGTAGAATTGGATTGGTATCCGGATGCCCCCTGGAGAAATGGTGGT
CCTCACCTGTGACACCCCTGAAGAAGATGGTATCACCTGGACCTTGGACCAGAGCAGTGAGGTCTTAGGCTCTG
GCAAAACCCTGACCATCCAAGTCAAAGAGTTTGGAGATGCTGGCCAGTACACCTGTCACAAAGGAGGCGAGGTT
CTAAGCCATTCGCTCCTGCTGCTTACAAAAAGGAAGATGGAATTTGGTCCACTGATATTTTAAAGGACCAGAA
AGAACCCAAAAATAAGACCTTTCTAAGATGCGAGGCCAAGAATTATTCTGGACGTTTCACCTGCTGGTGGCTGA
CGACAATCAGTACTGATTTGACATTCACTGTCAAAGCAGCAGAGGCTCTTCTGACCCCCAAGGGGTGACGTGC
GGAGCTGCTACACTCTCTGCAGAGAGAGTCAGAGGGGACAACAAGGAGTATGAGTACTCAGTGGAGTGCCAGGA
GGACAGTGCCTGCCAGCTGCTGAGGAGAGTCTGCCCATGAGGTGATGGTGGATGCCGTTTACAAAGCTCAAGT
ATGAAAACCTACACCAGCAGCTTCTTCATCAGGGACATCATCAAACCTGACCCACCCAAGAACTTGCAGCTGAAG
CCATTAAAGAATTCTCGGCAGGTGGAGGTCAGCTGGGAGTACCCTGACACCTGGAGTACTCCACATTCCTACTT
CTCCCTGACATTCTGCGTTTCAGGTCCAGGGCAAGAGCAAGAGAGAAAAAGAAAGATAGAGTCTTACGGACAAGA
CCTCAGCCACGGTCATCTGCCGCAAAAATGCCAGCATTAGCGTGCGGGCCCAGGACCGCTACTATAGCTCATCT
TGGAGCGAATGGGCATCTGTGCCCTGCAGT

(3) Wild Type IL12A without signal peptide Amino acids (SEQ ID NO: 3):

RNLFPVATPDPMFPLHHSQNLLRAVSNMLQKARQTFLEFYPCTSEEIDHEDITKDKTSTVEACLPLELTKNES
LNSRETSFITNGSCLASRKTSFMMALCLSSIIYEDLKMYQVEFKTMNAKLLMDPKRQIFLDQNMLAVIDELMQAL
NFNSETVPQKSSLEEPDFYKTKIKLCILLHAFRIRAVTIDRVMSYLNAS

(4) Wild Type IL12A without signal peptide Nucleic acids (SEQ ID NO: 4):

AGAAACCTCCCCGTGGCCACTCCAGACCCAGGAATGTTCCCATGCCTTCACCACTCCCCAAACCTGCTGAGGGCC
GTCAGCAACATGCTCCAGAAGGCCAGACAACTCTAGAATTTTACCCTTGCACTTCTGAAGAGATTGATCATGAA
GATATCACAAAAGATAAAACCAGCACAGTGGAGGCCTGTTTACCATTGGAATTAACCAAGAATGAGAGTTGCCTA
AATTCCAGAGAGACCTCTTTCATAACTAATGGGAGTTGCCTGGCCTCCAGAAAGACCTCTTTTATGATGGCCCTG
TGCCTTAGTAGTATTTATGAAGACTTGAAGATGTACCAGGTGGAGTTCAAGACCATGAATGCAAAGCTTCTGATG
GATCCTAAGAGGCAGATCTTCTAGATCAAAACATGCTGGCAGTTATTGATGAGCTGATGCAGGCCCTGAATTC
AACAGTGAGACTGTGCCACAAAAATCCTCCCTTGAAGAACCGGATTTTATAAAACTAAAATCAAGCTCTGCATA
CTTCTTCATGCTTTCAGAATTCGGGCAGTGACTATTGATAGAGTGATGAGCTATCTGAATGCTTCC

(5) Wild Type IL12B signal peptide Amino acids: (SEQ ID NO: 45)

MCHQQLVISWFSLVFLASPLVA

(6) Wild Type IL12B signal peptide Nucleic acids: (SEQ ID NO: 46)

ATGTGTCACCAGCAGTTGGTCATCTCTTGGTTTTCCCTGGTTTTTCTGGCATCTCCCCTCGTGGCC

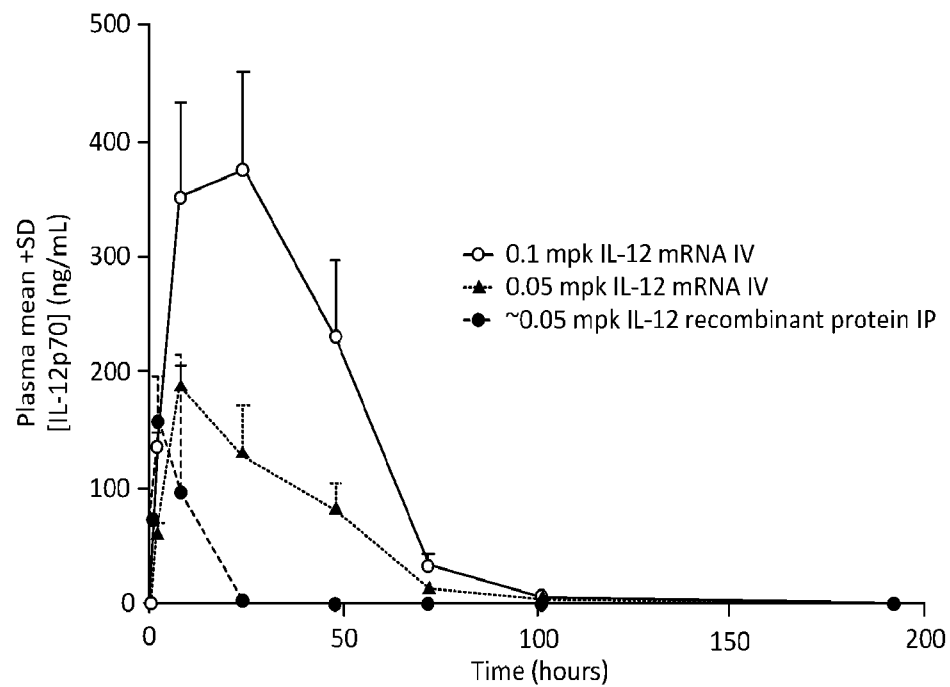
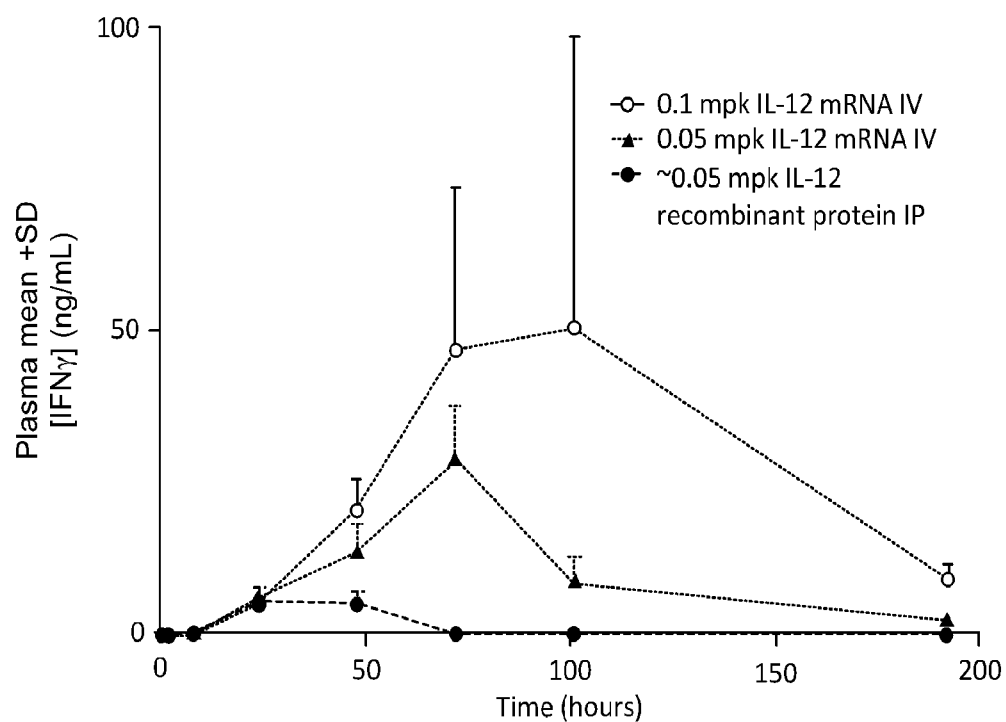
FIG. 2A. IL12 Levels**FIG. 2B. IFN γ Levels**

FIG. 3. A single IV dose of IL12 mRNA in LNP has robust efficacy

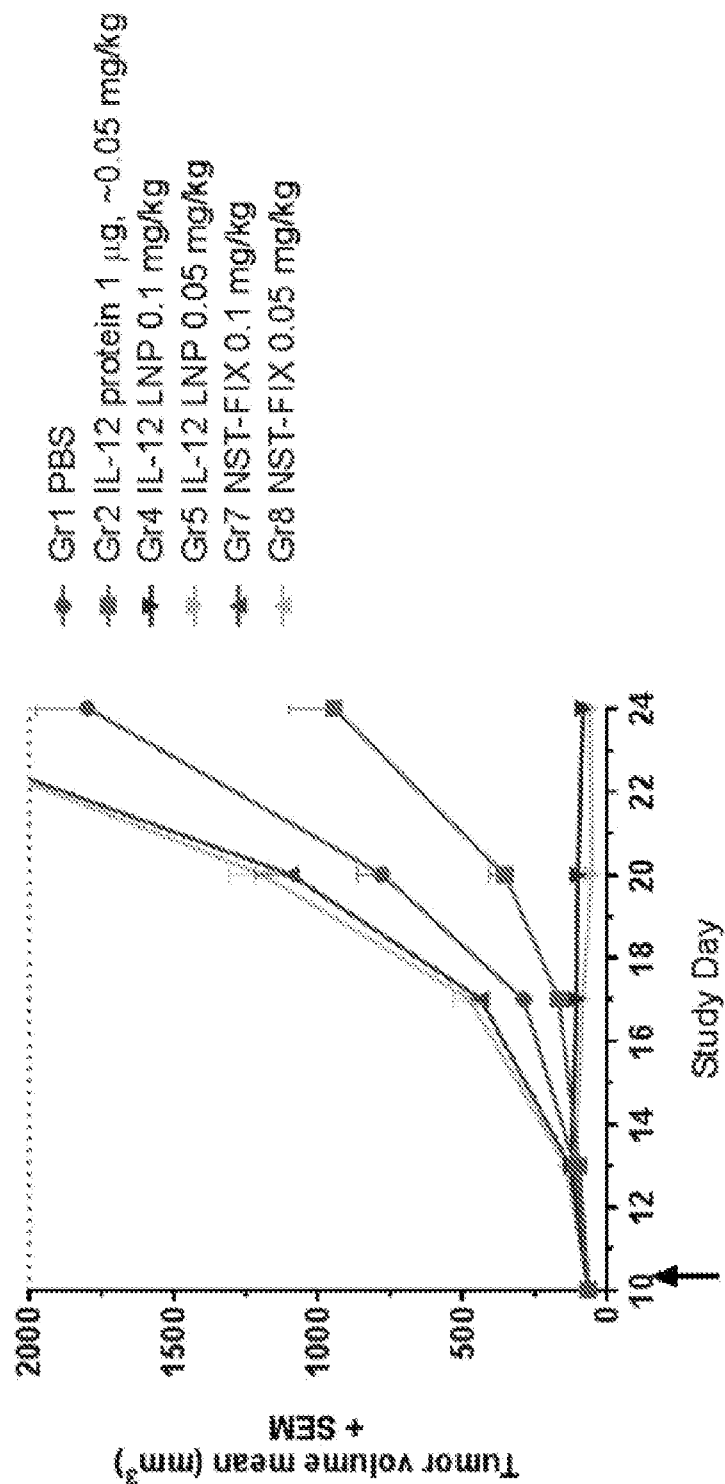


FIG. 4B

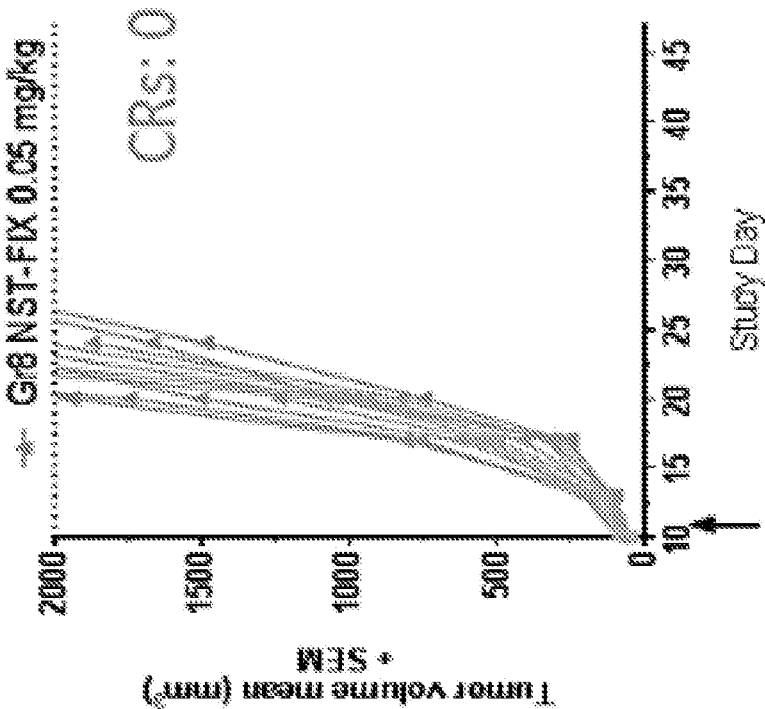


FIG. 4A

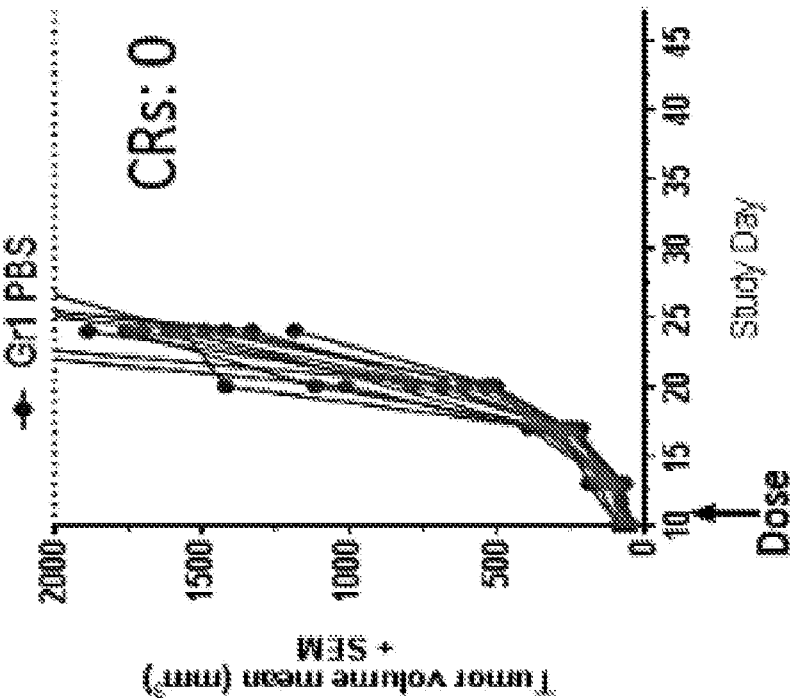


FIG. 4D

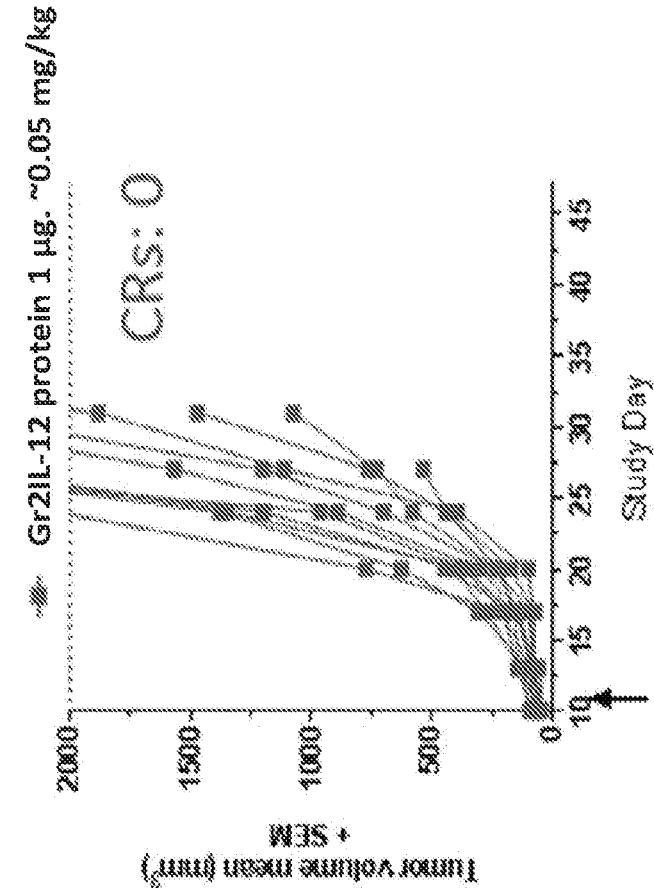


FIG. 4C

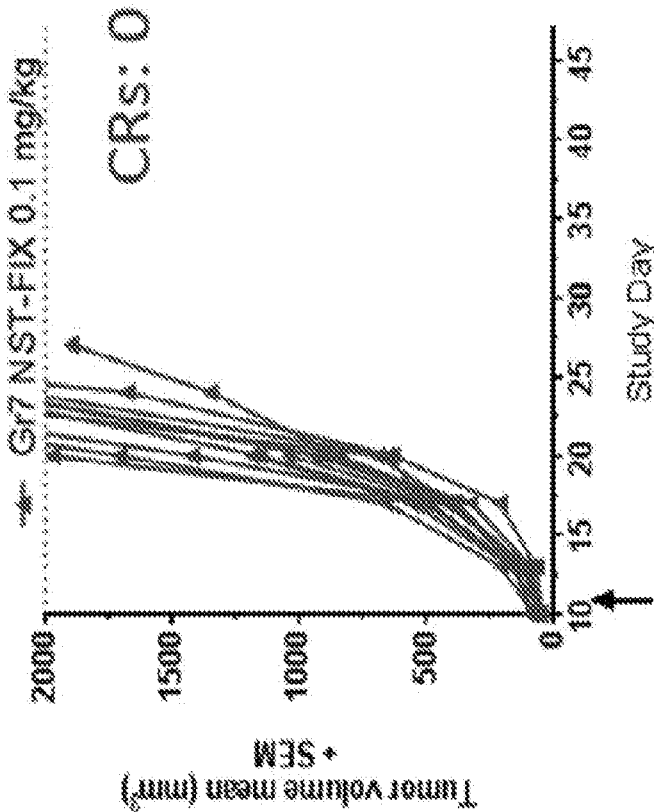


FIG. 4F

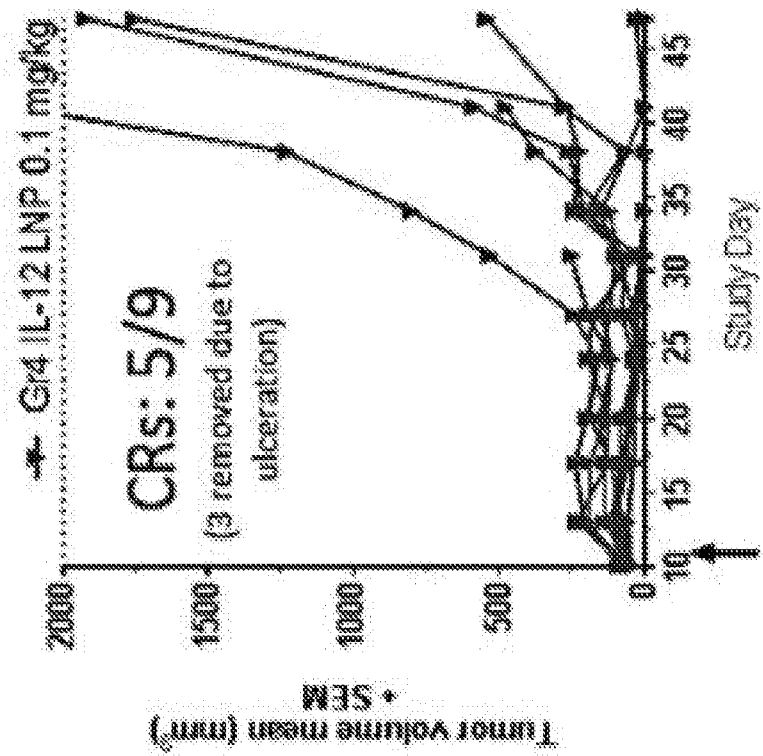


FIG. 4E

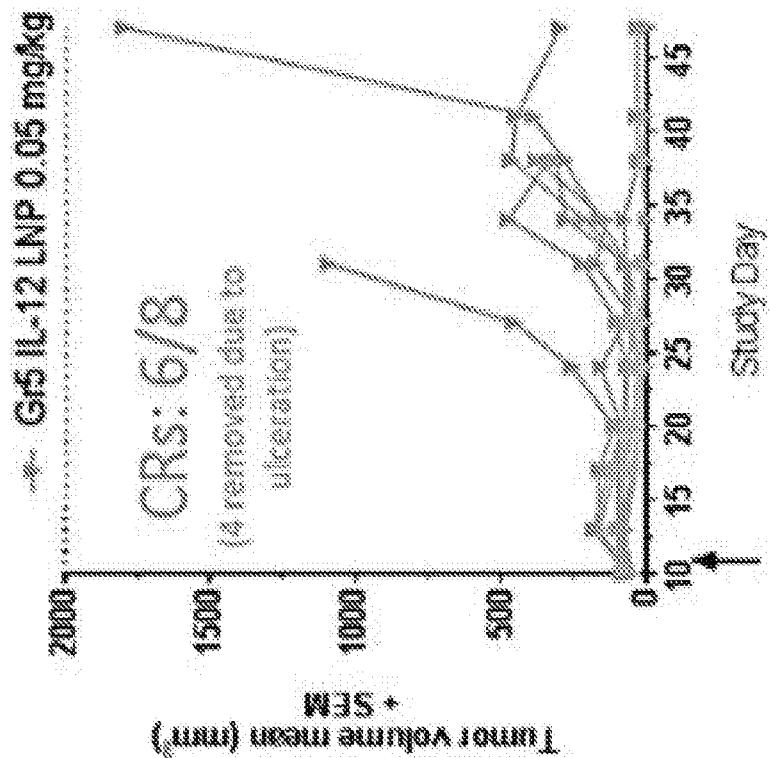


FIG. 5.

Kaplan-meir (Day 47)

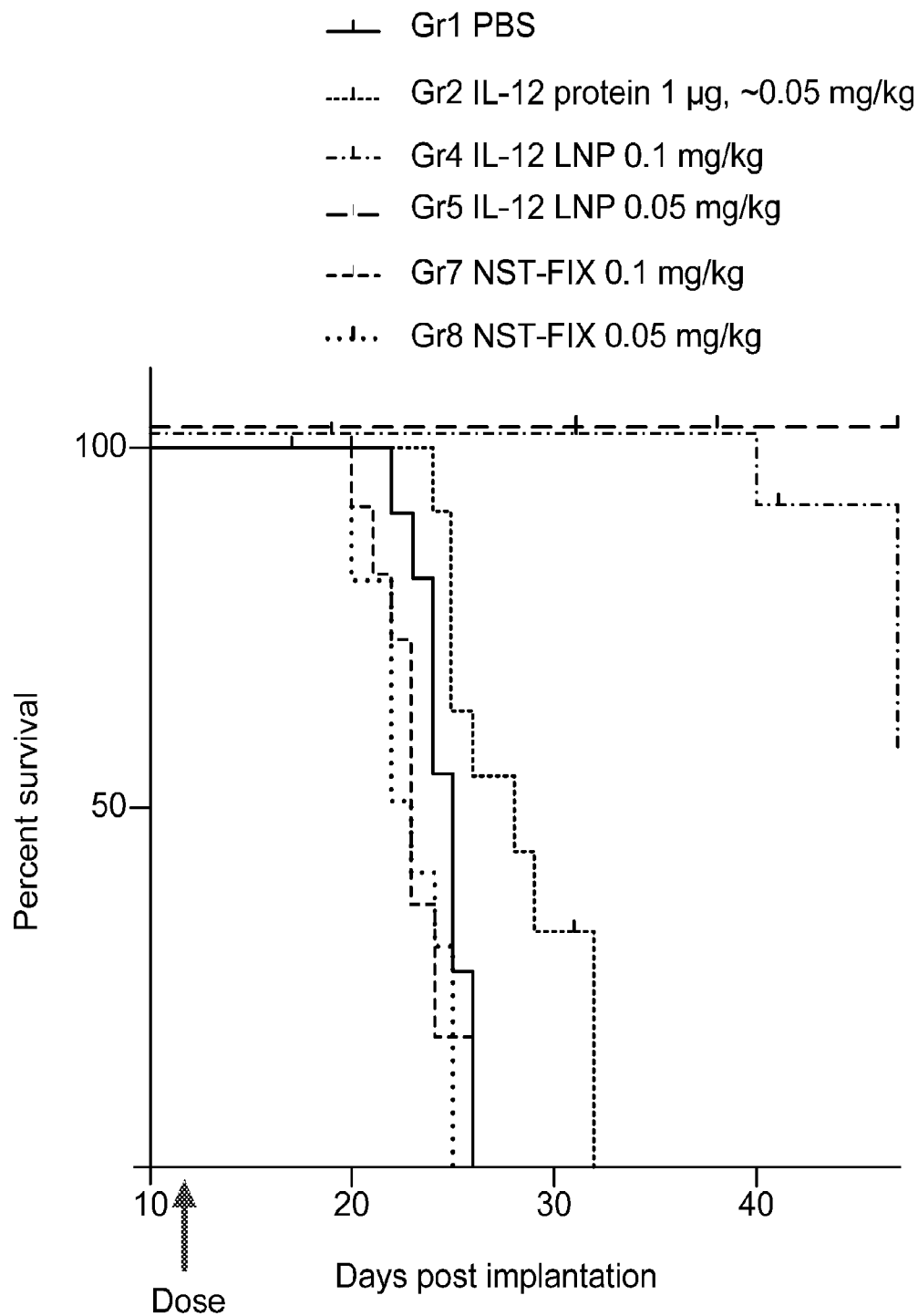


FIG. 6A

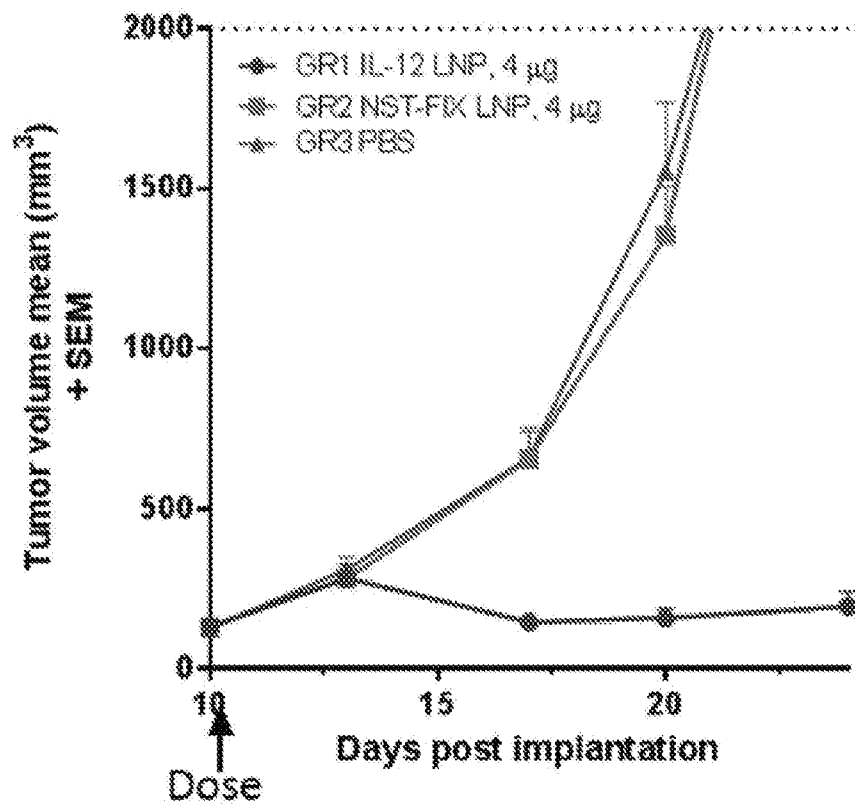


FIG. 6B

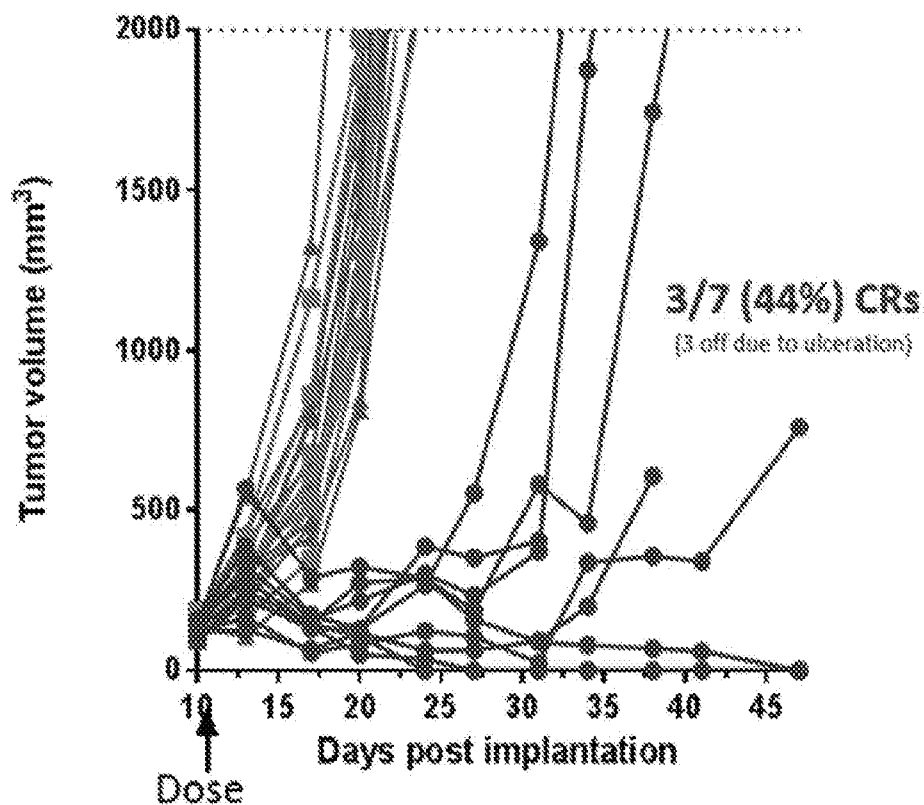


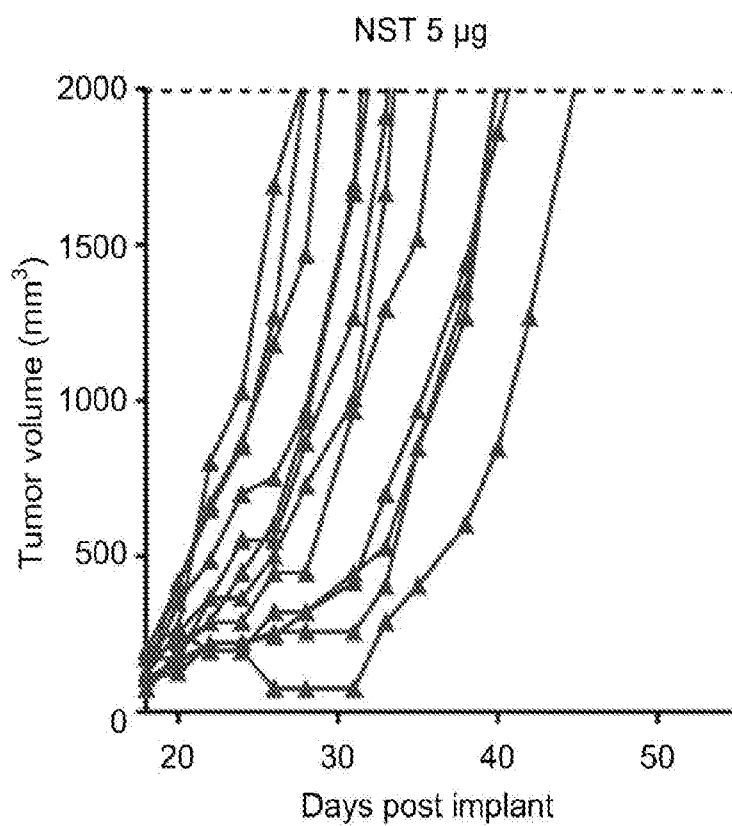
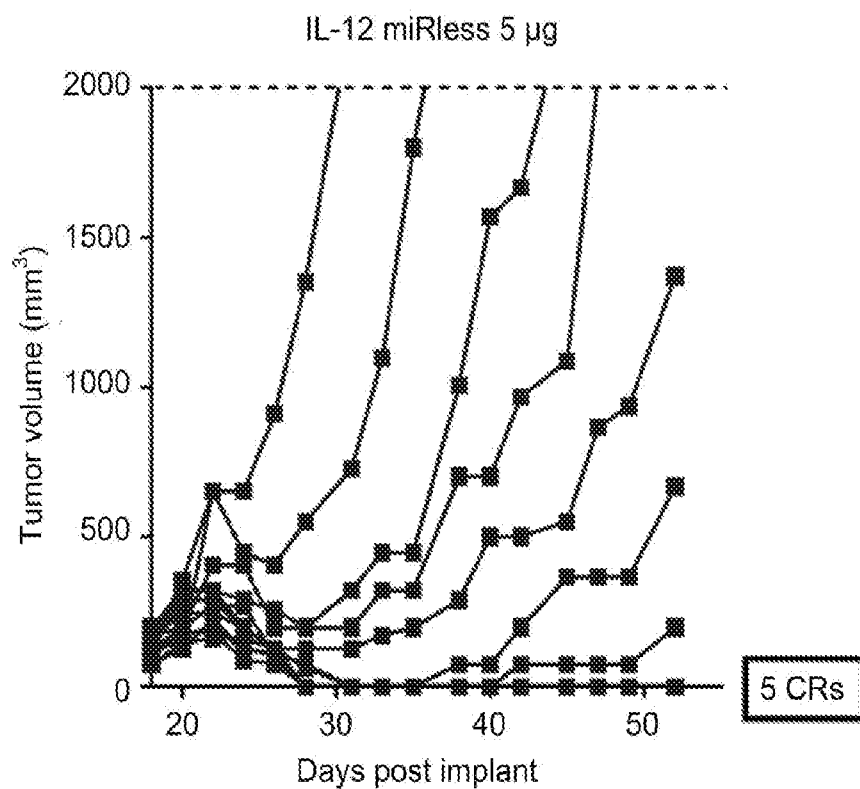
FIG. 7A**FIG. 7B**

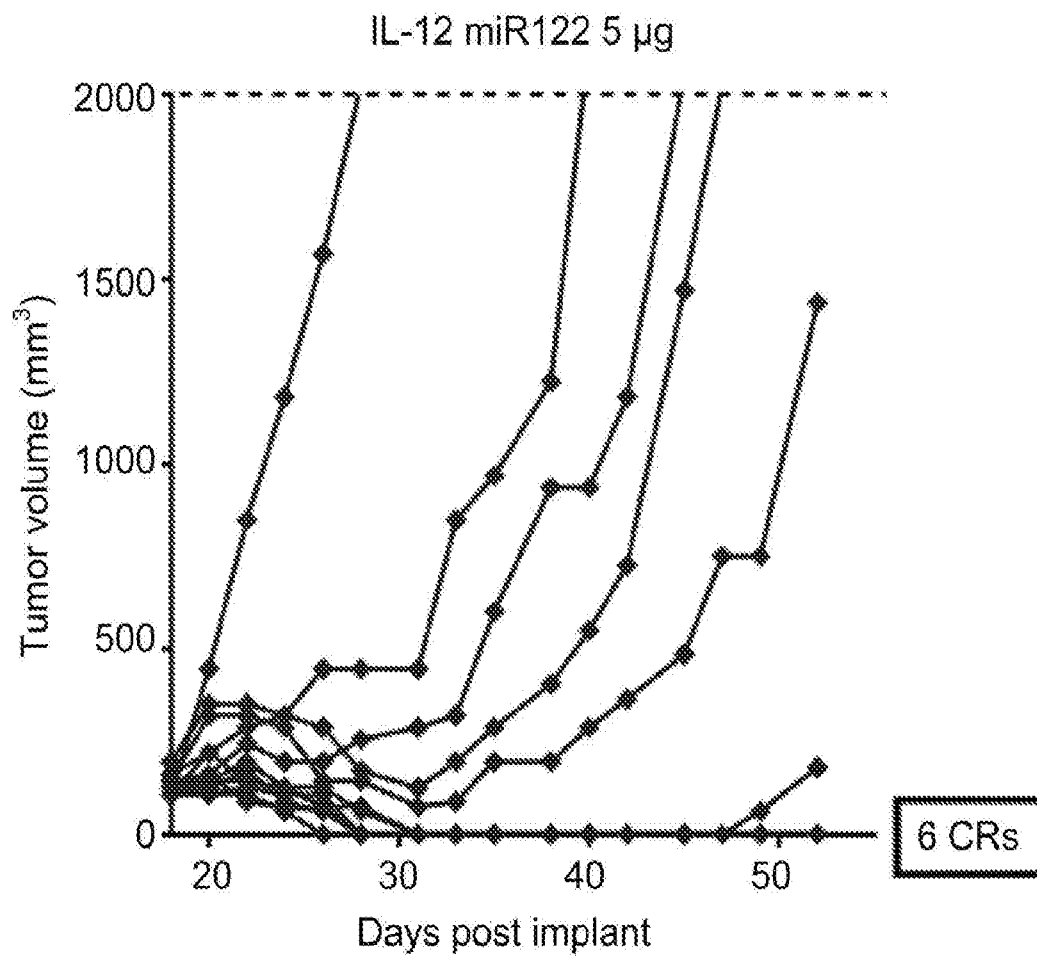
FIG. 7C

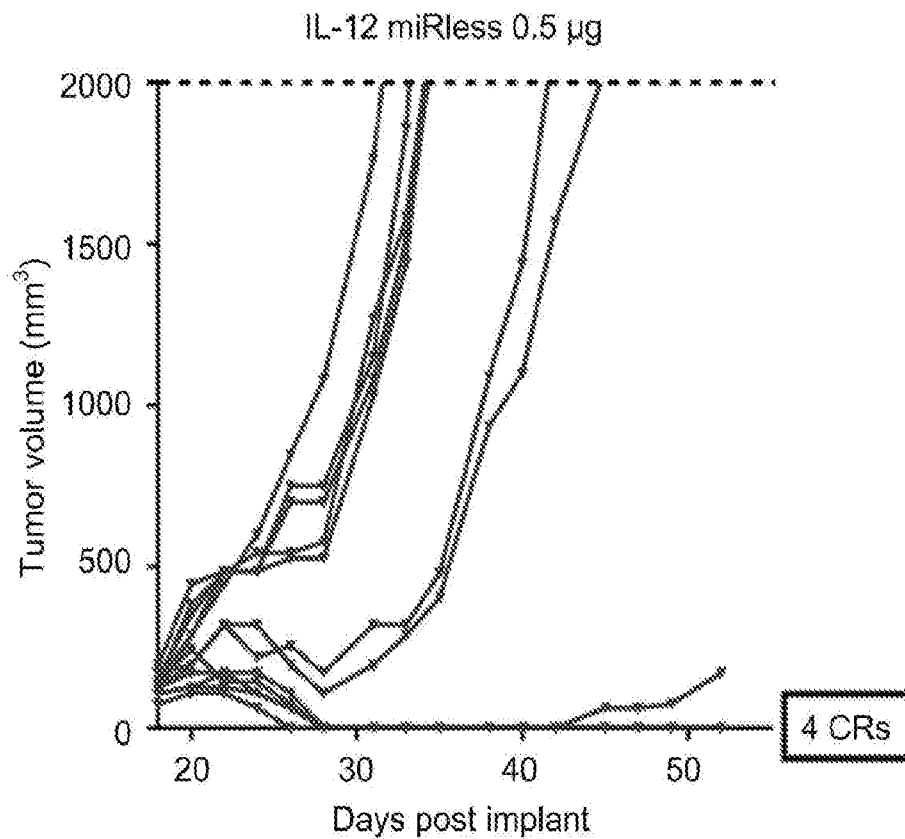
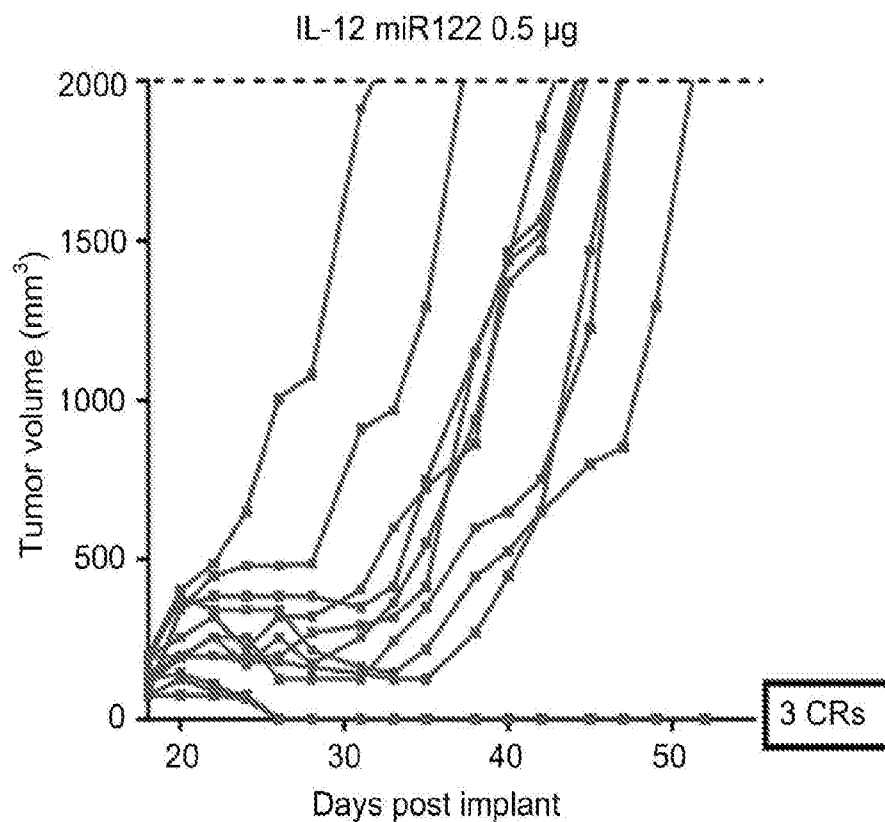
FIG. 8A**FIG. 8B**

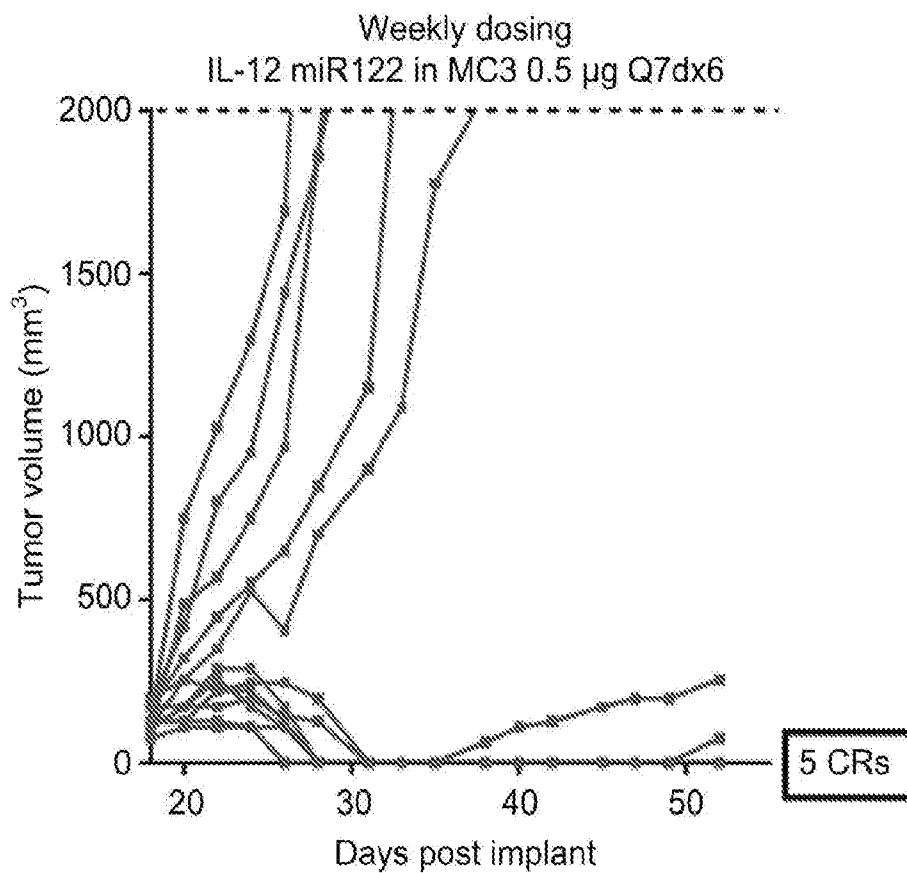
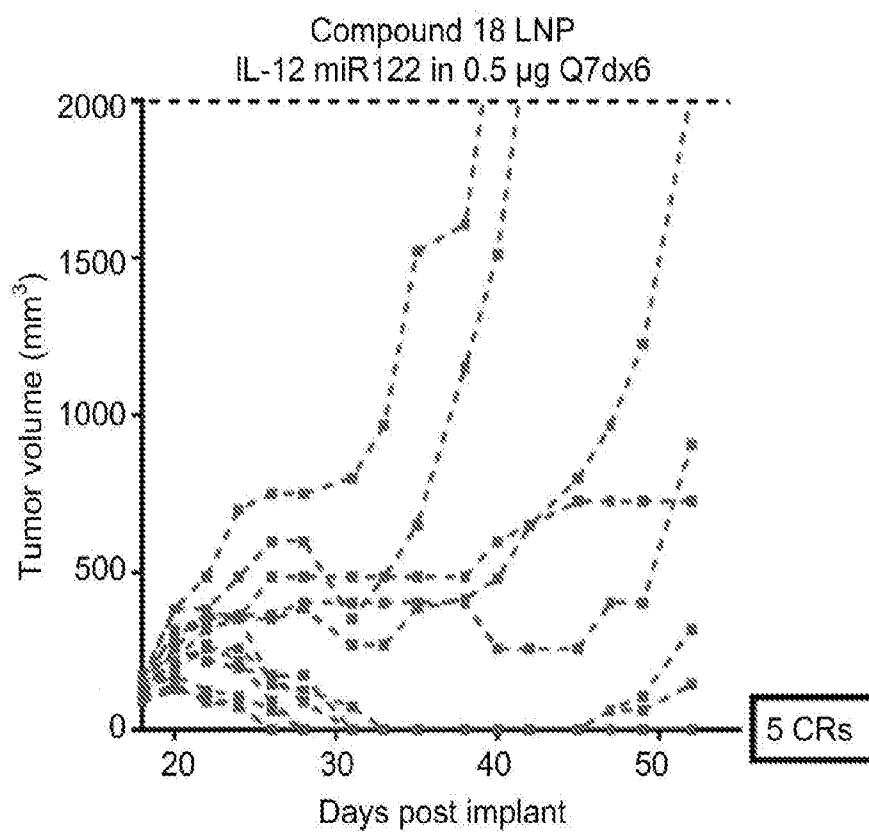
FIG. 8C**FIG. 8D**

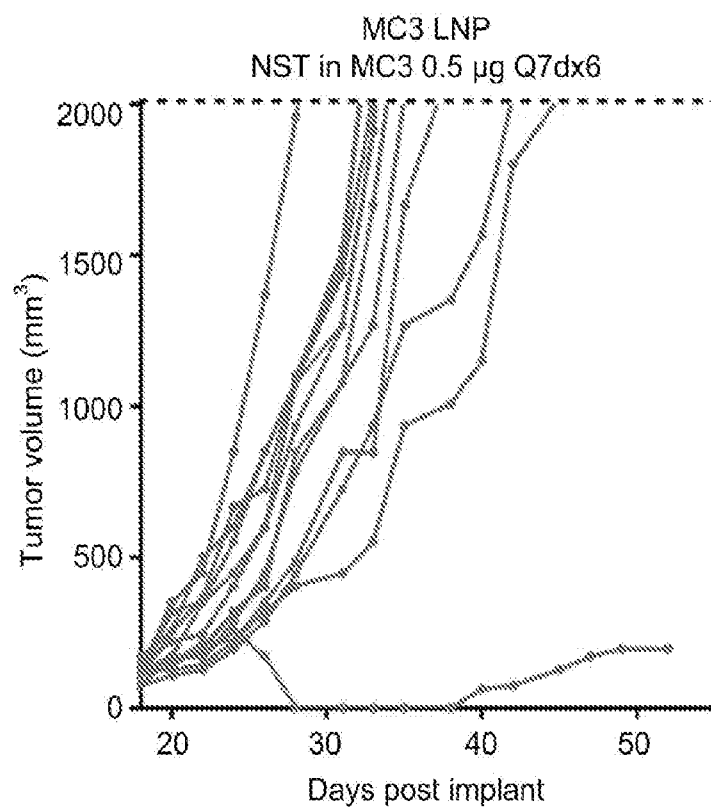
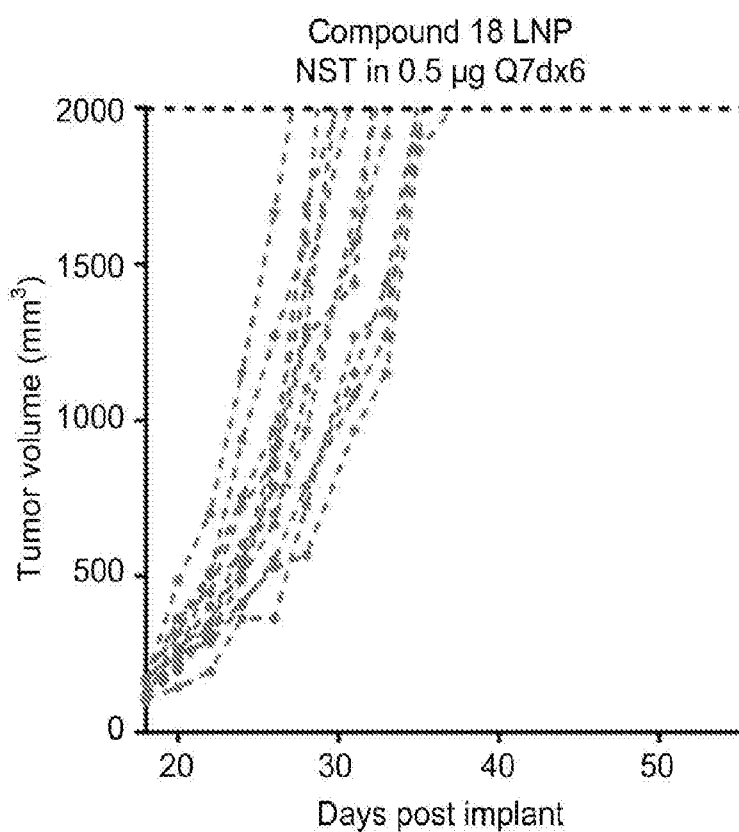
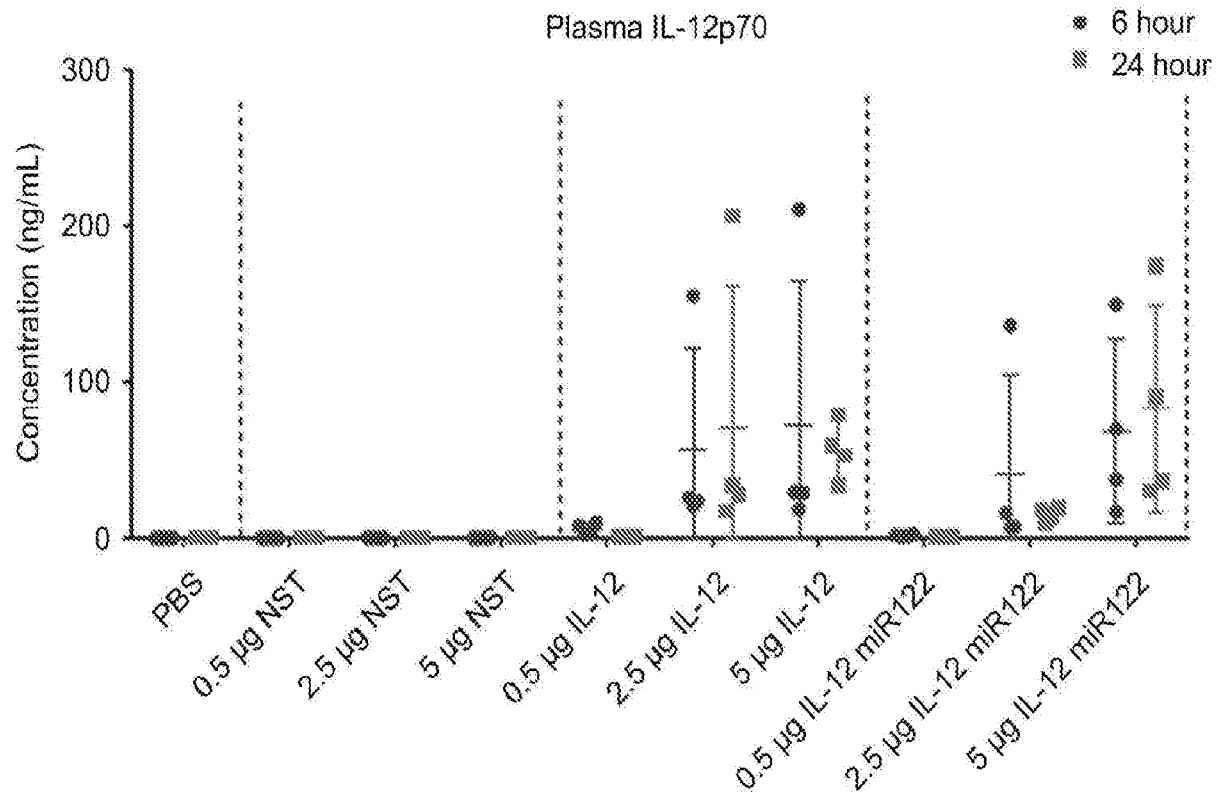
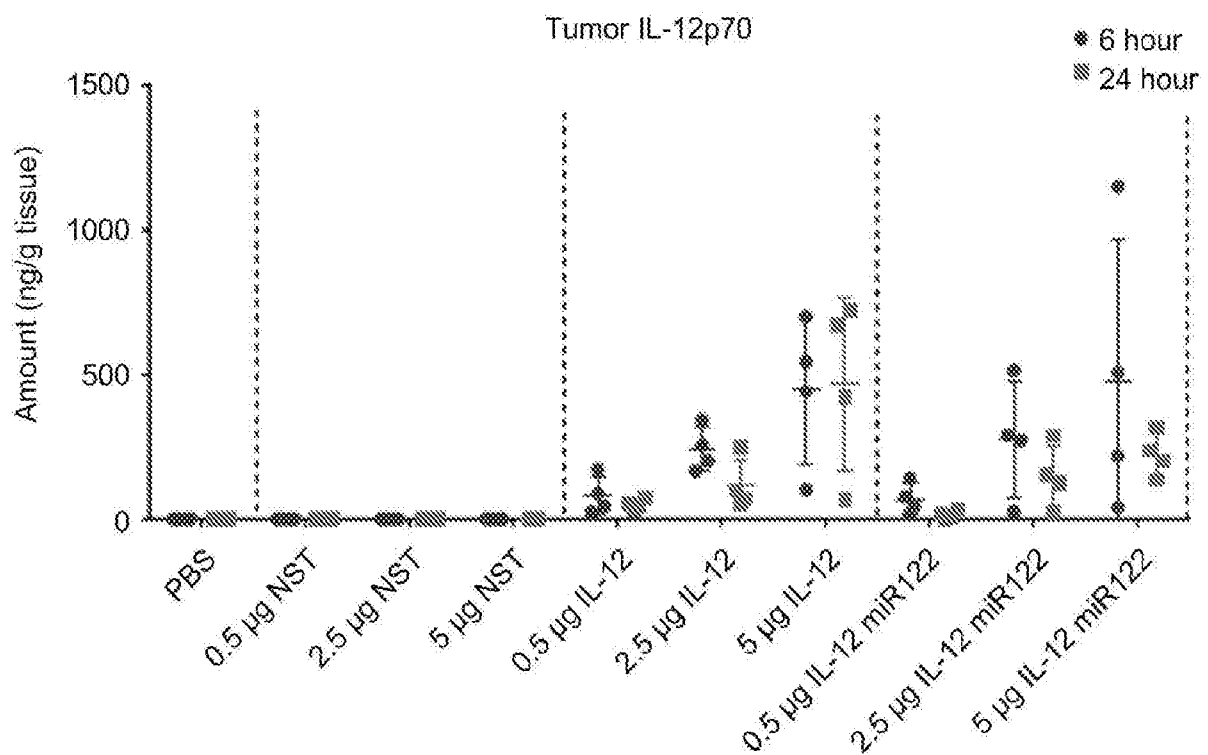
FIG. 8E**FIG. 8F**

FIG. 9A**FIG. 9B**

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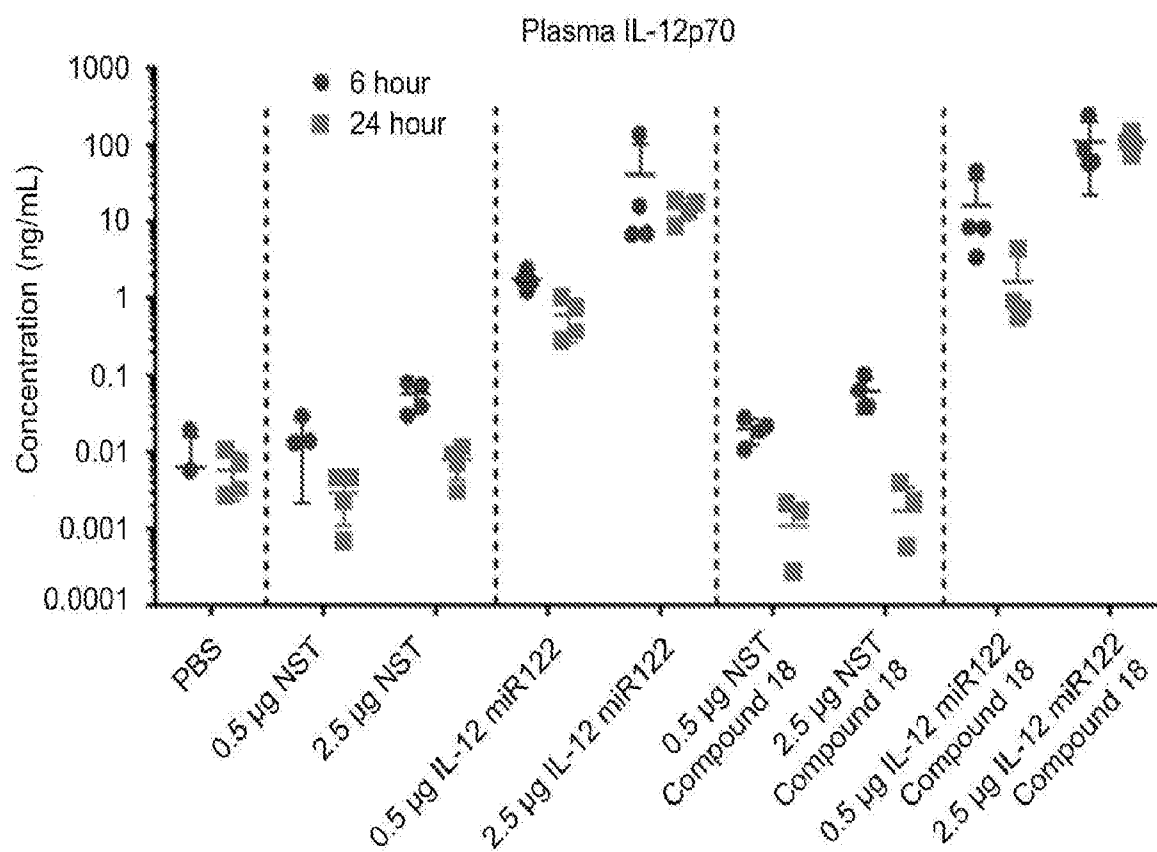
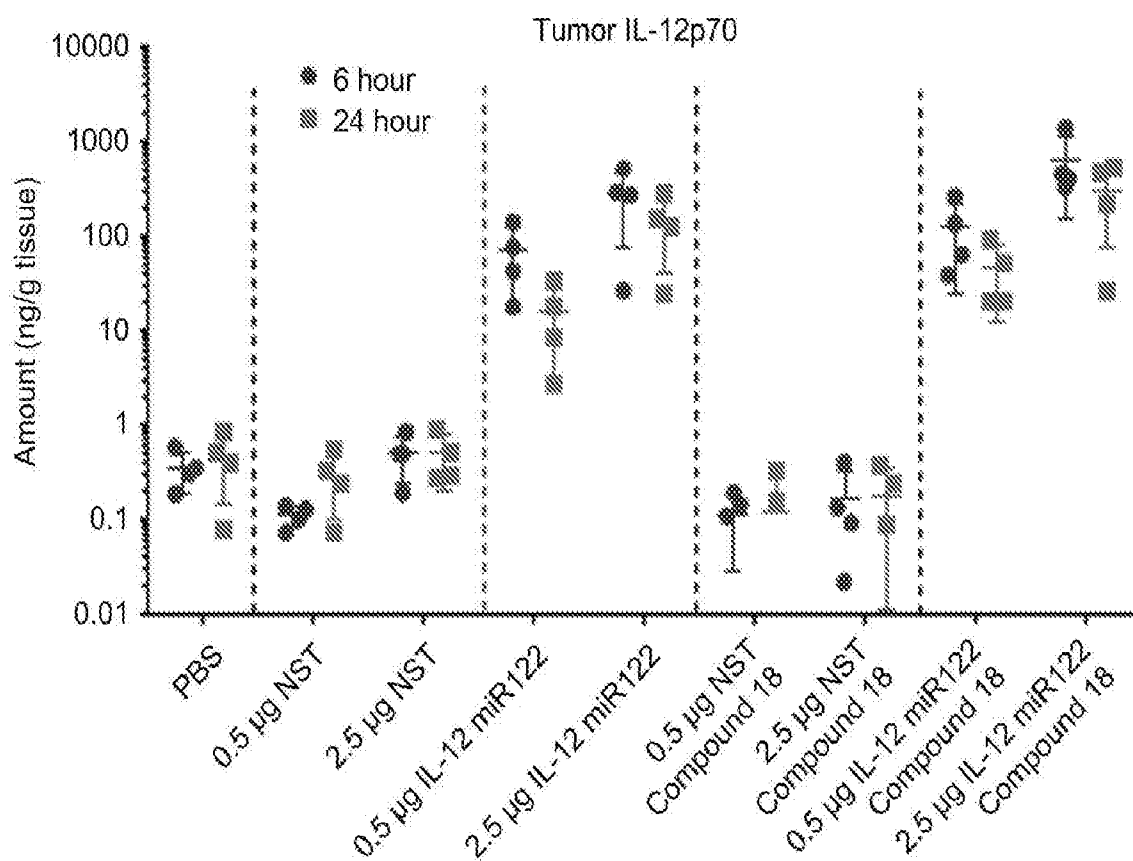
FIG. 9C**FIG. 9D**

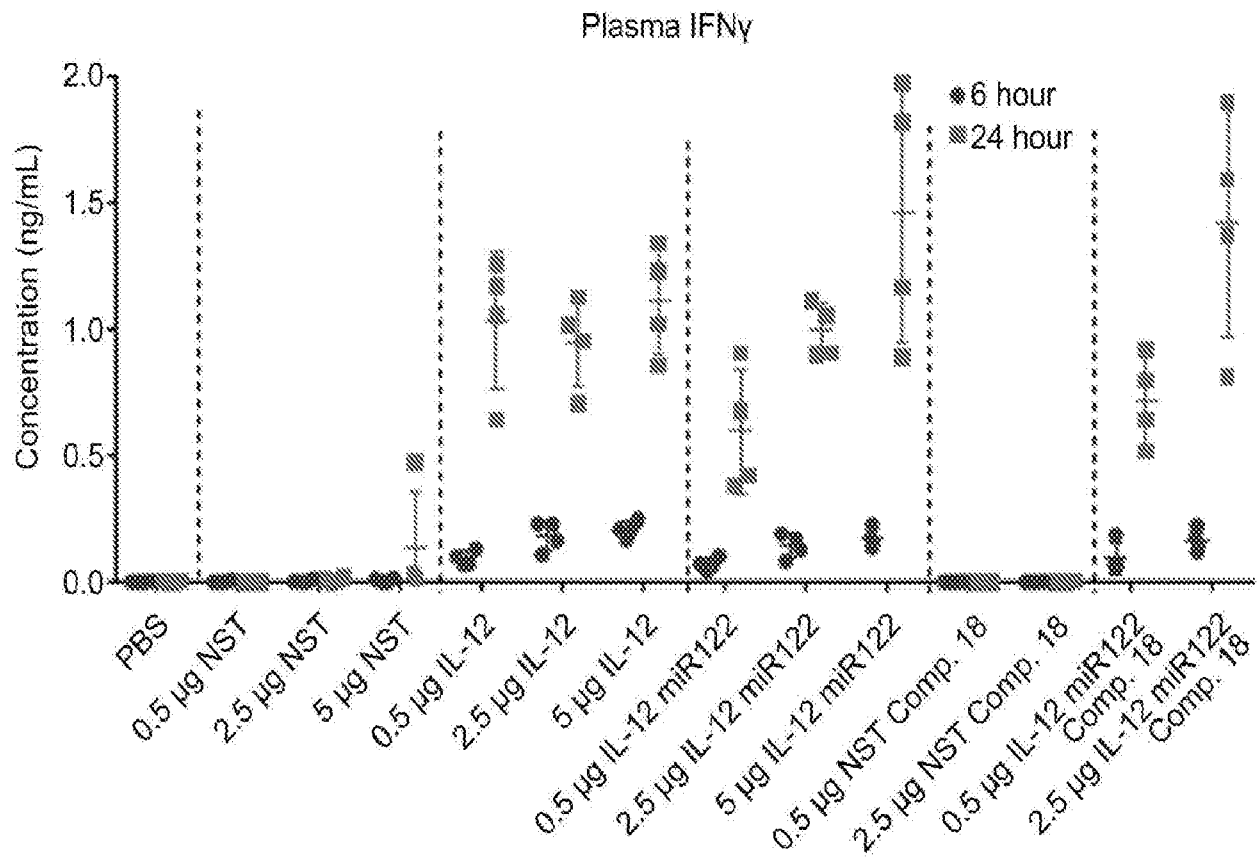
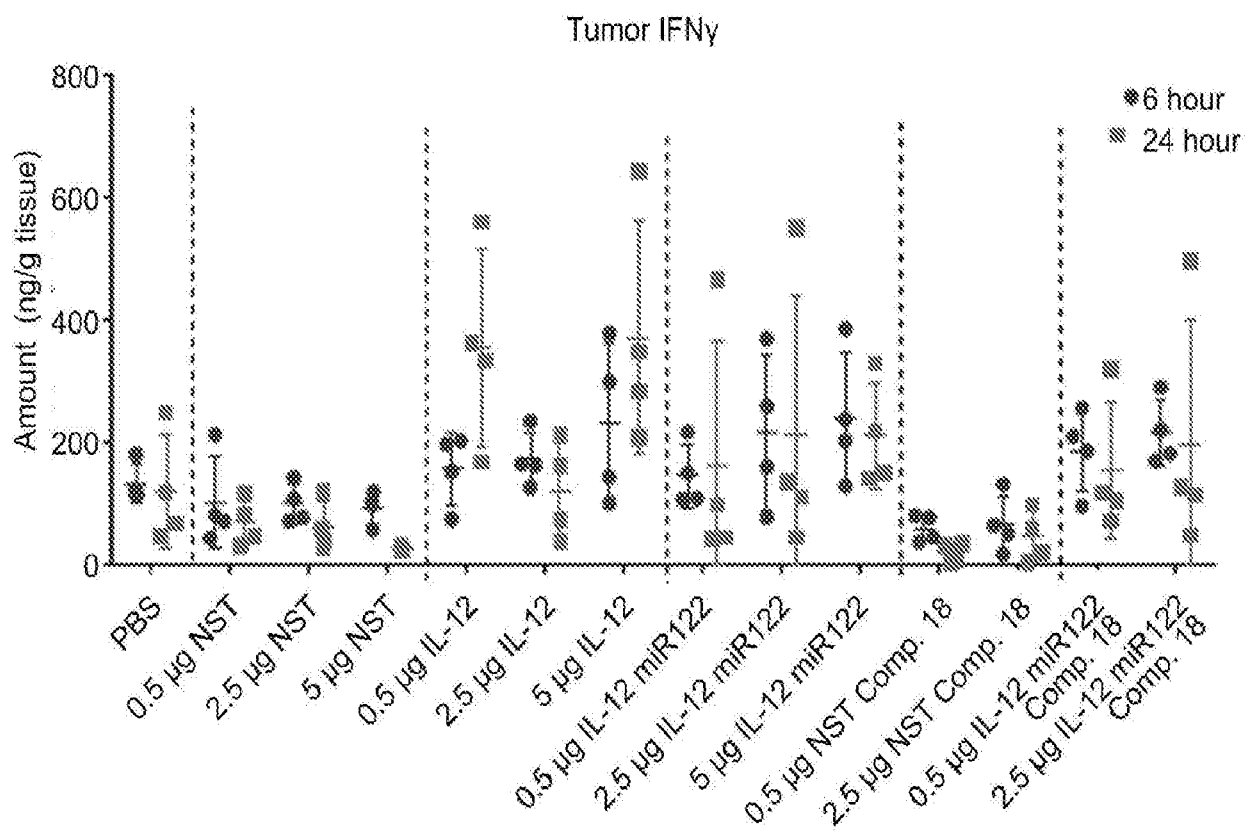
FIG. 9E**FIG. 9F**

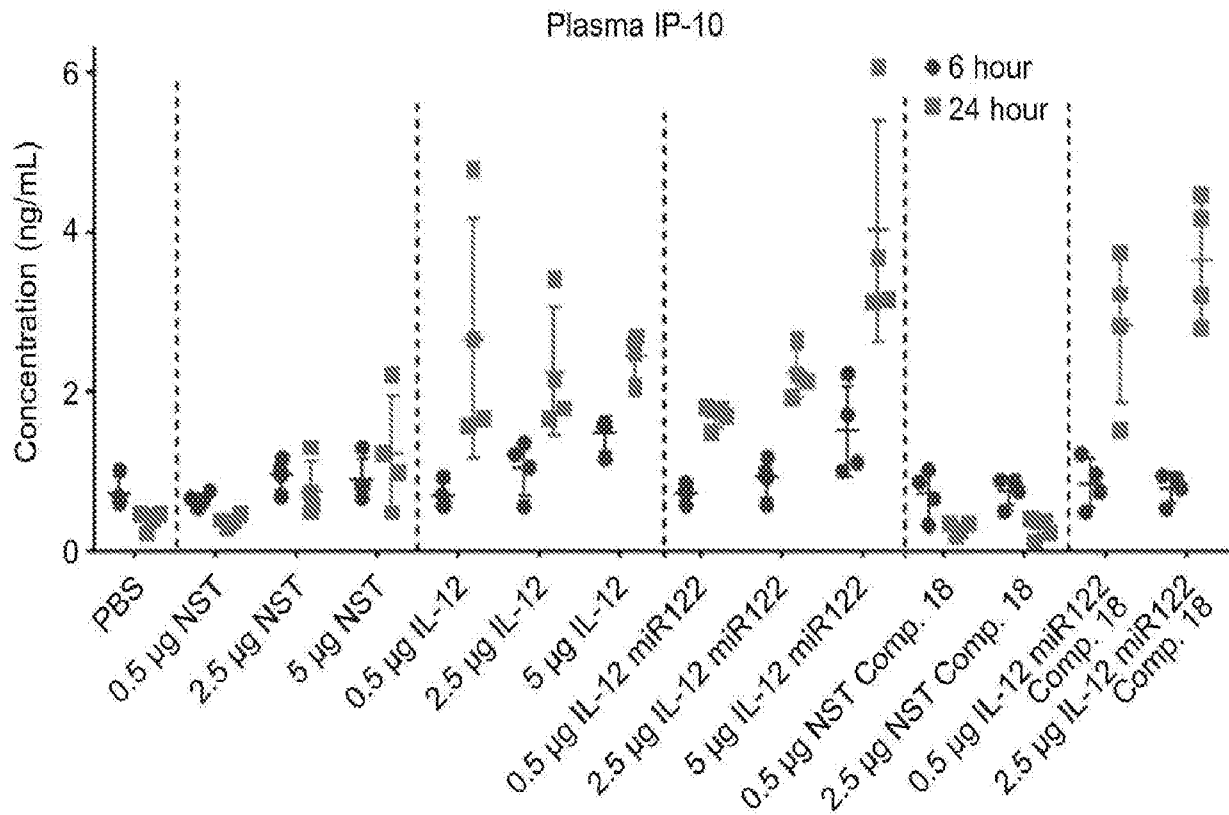
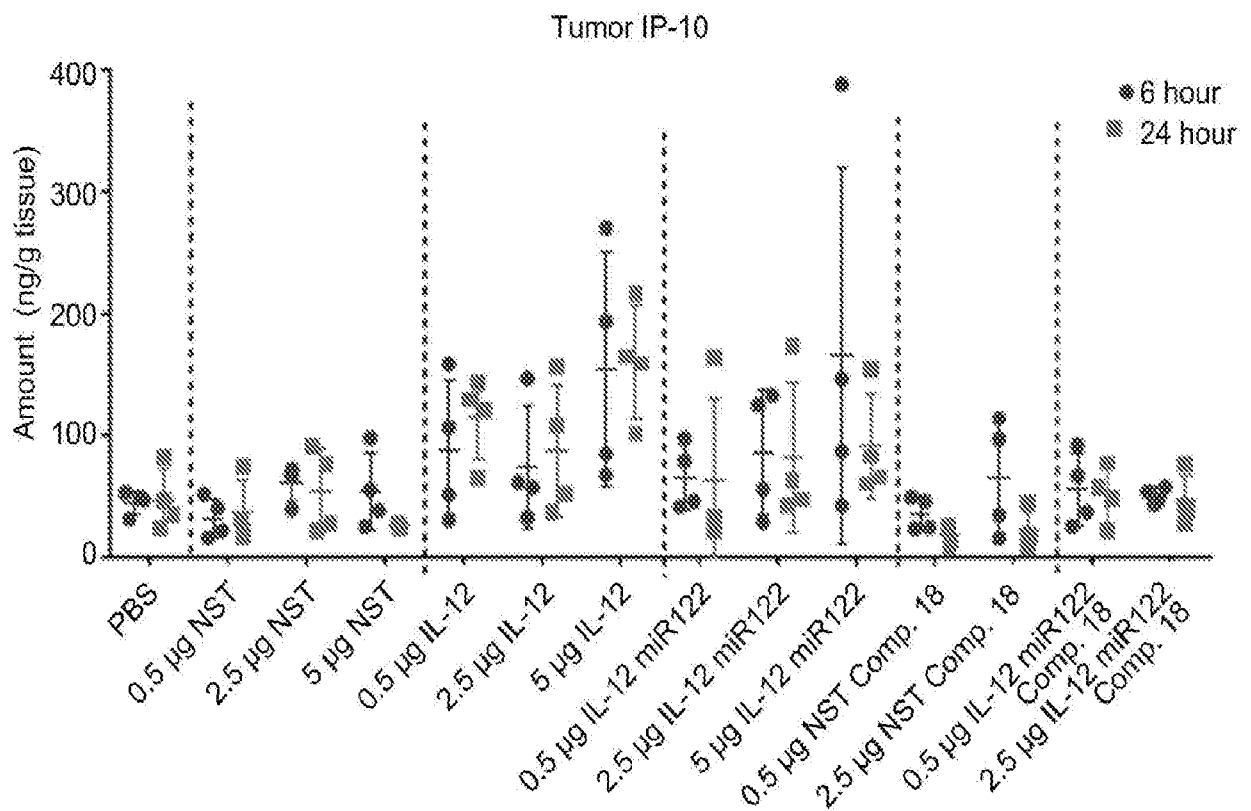
FIG. 9G**FIG. 9H**

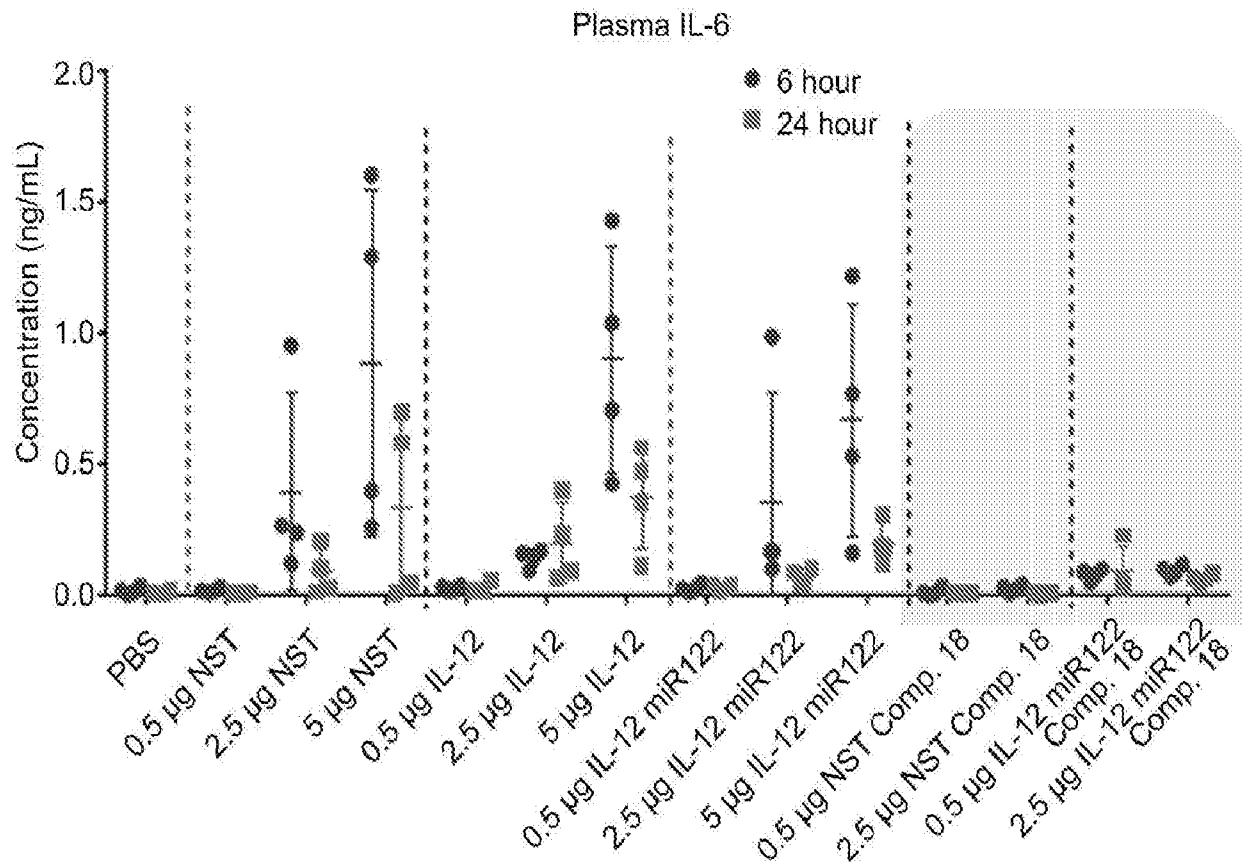
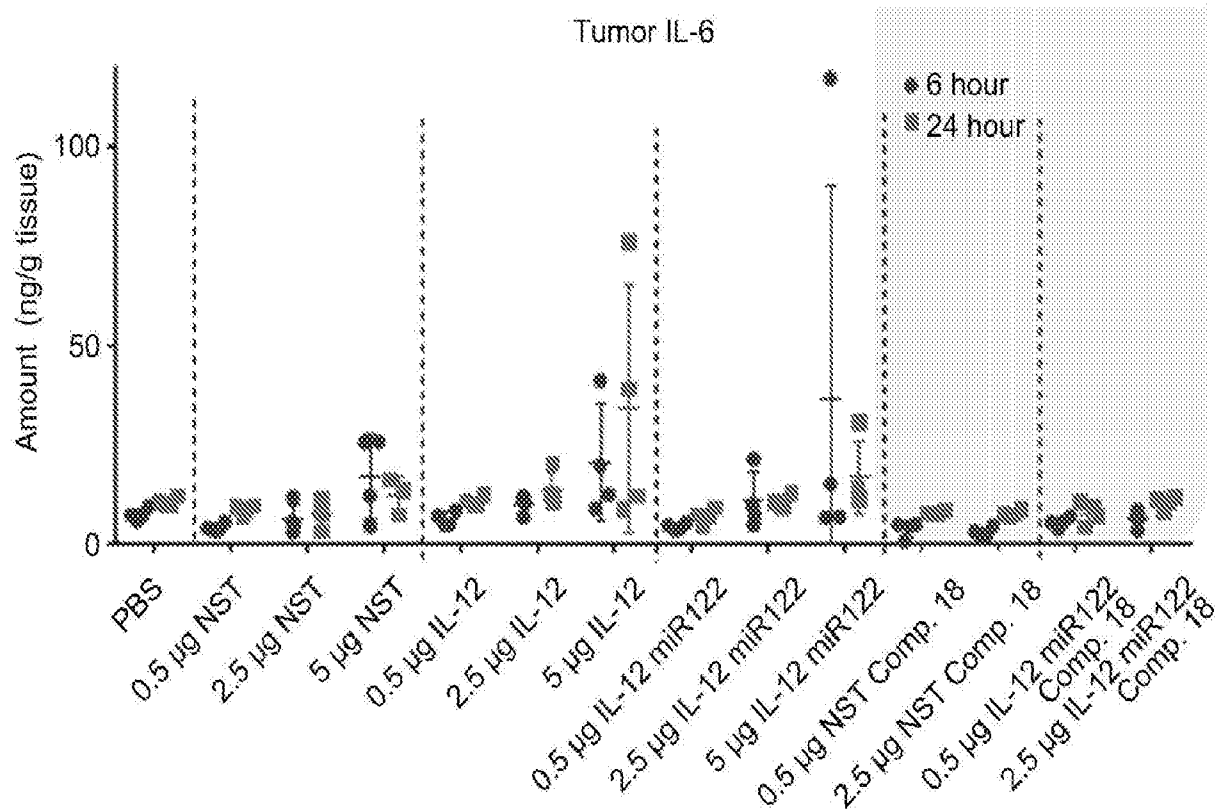
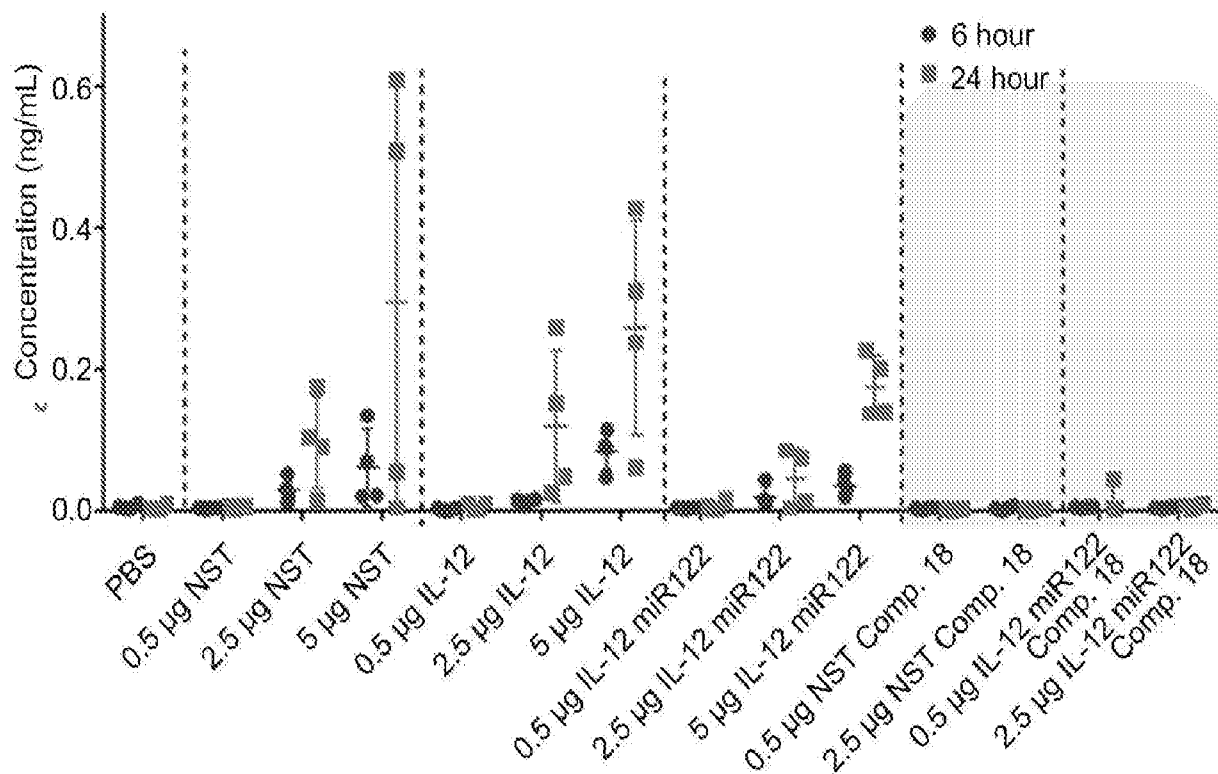
FIG. 9I**FIG. 9J**

FIG. 9K

Plasma G-CSF

**FIG. 9L**

Tumor G-CSF

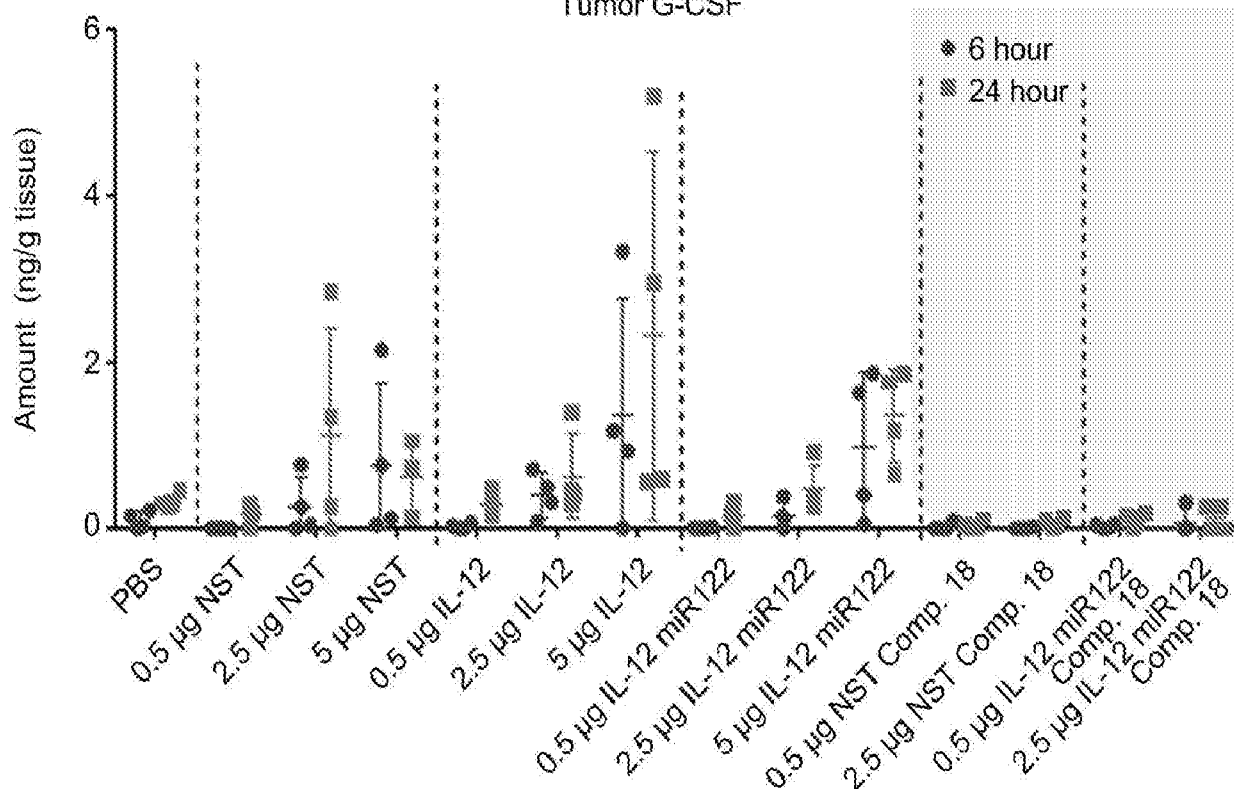


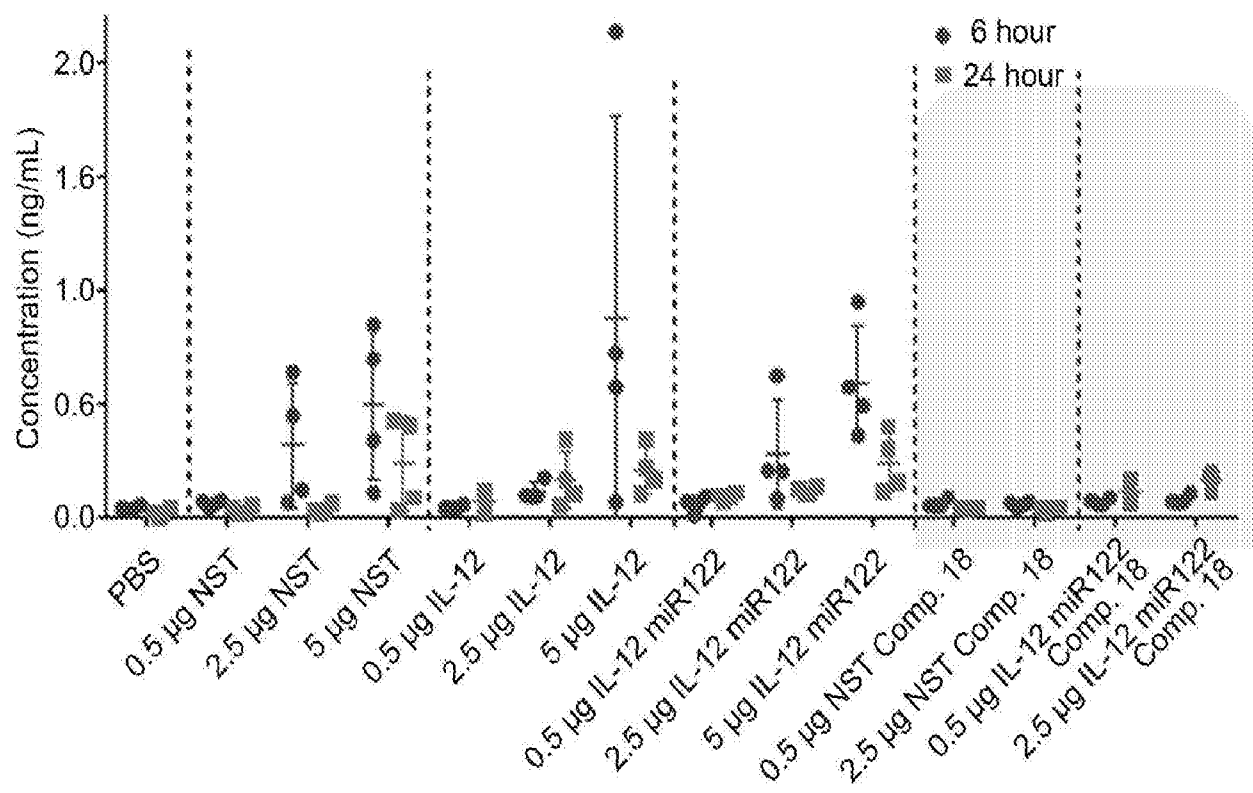
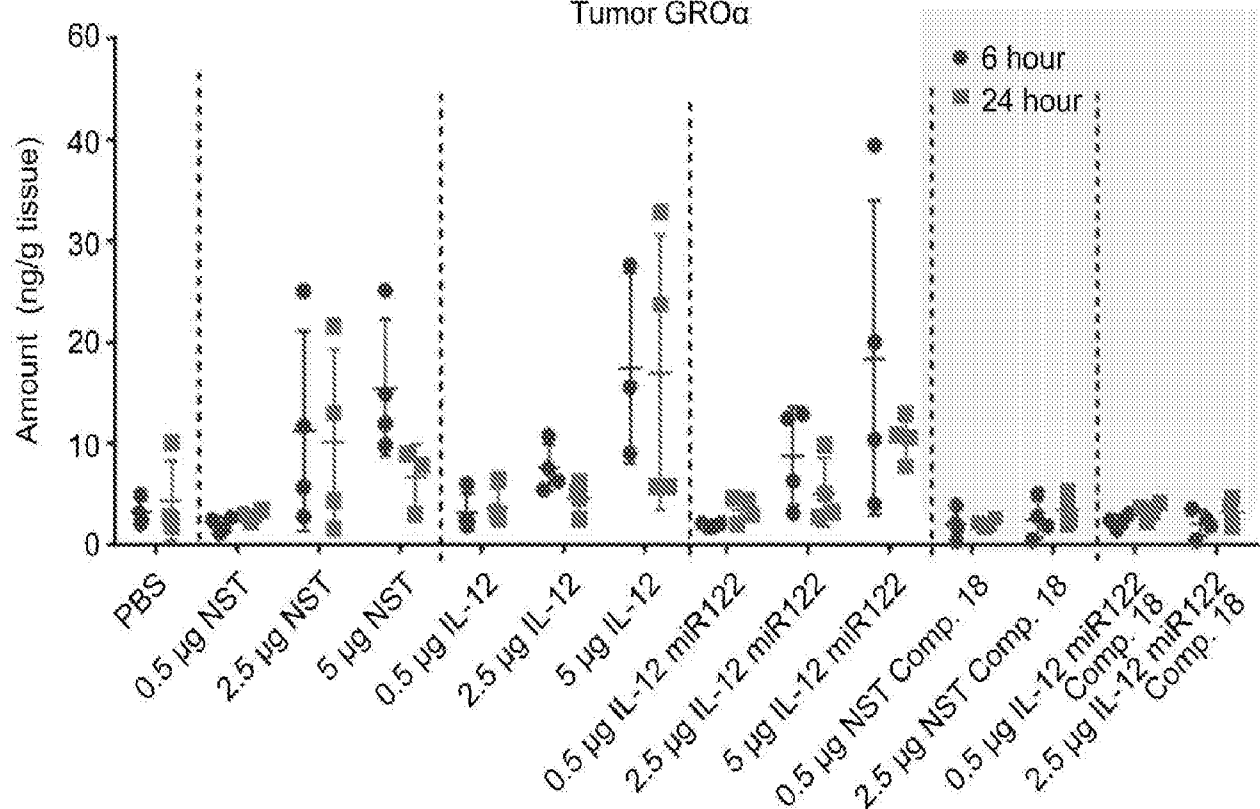
FIG. 9MPlasma GRO α **FIG. 9N**Tumor GRO α 

FIG. 10A

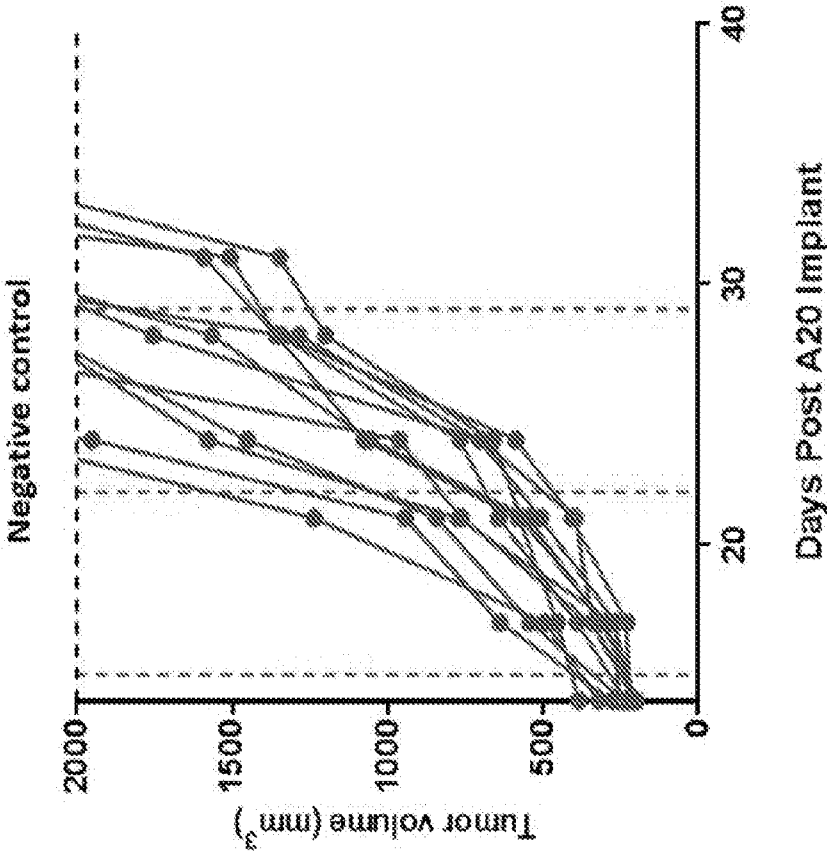


FIG. 10B

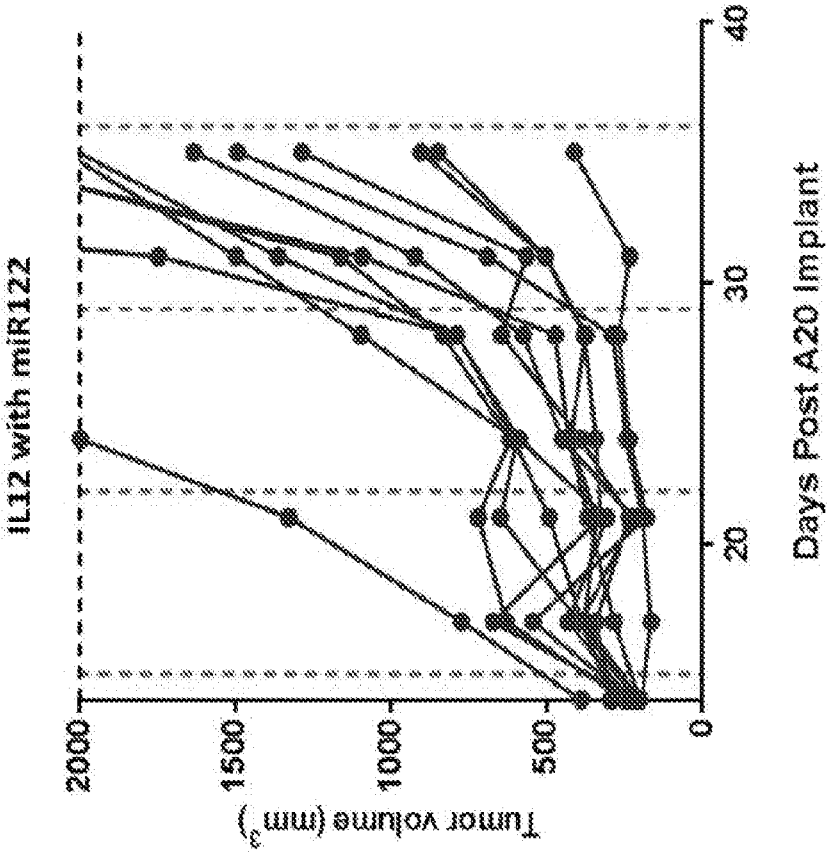


FIG.10D

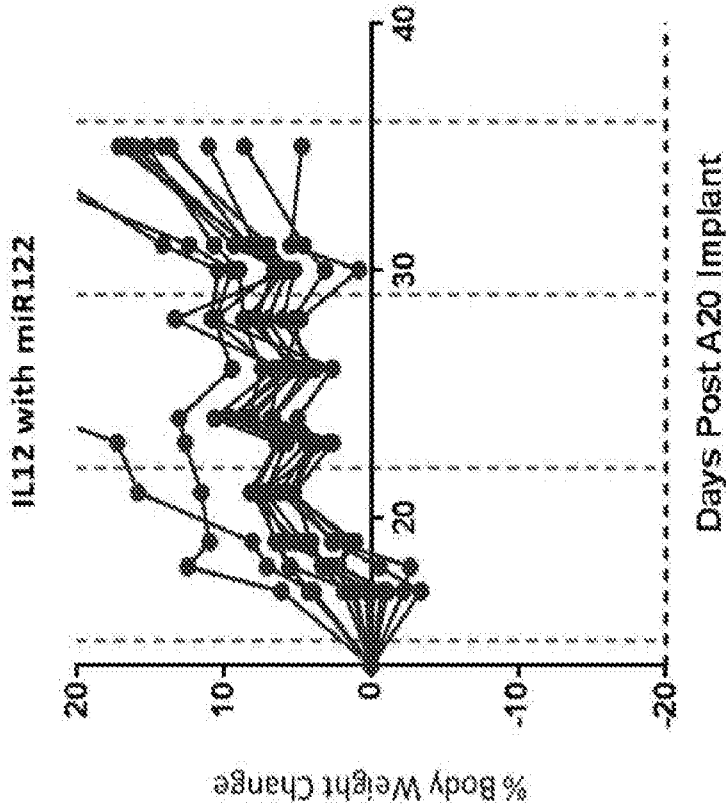


FIG.10C

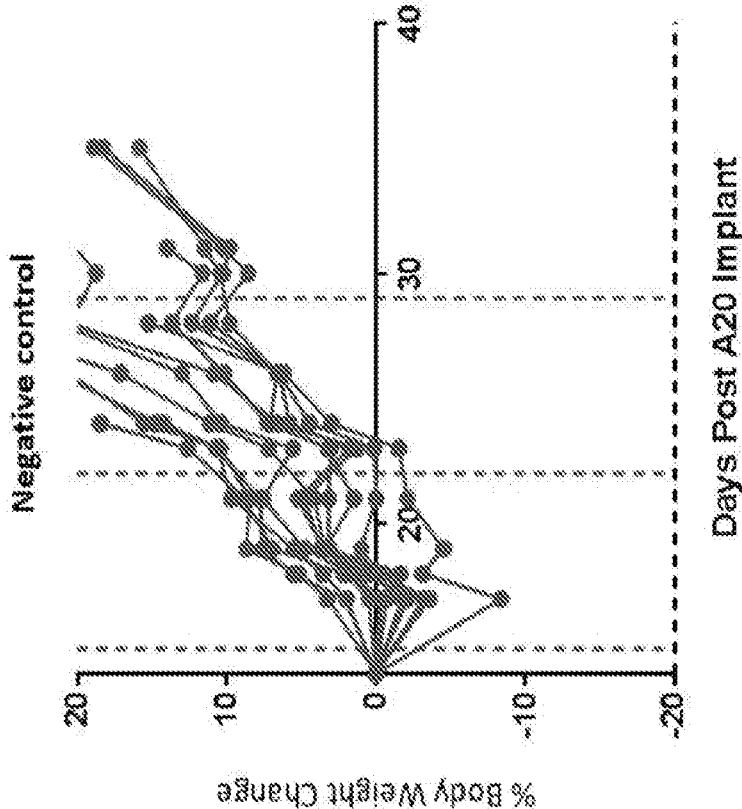


FIG. 11A

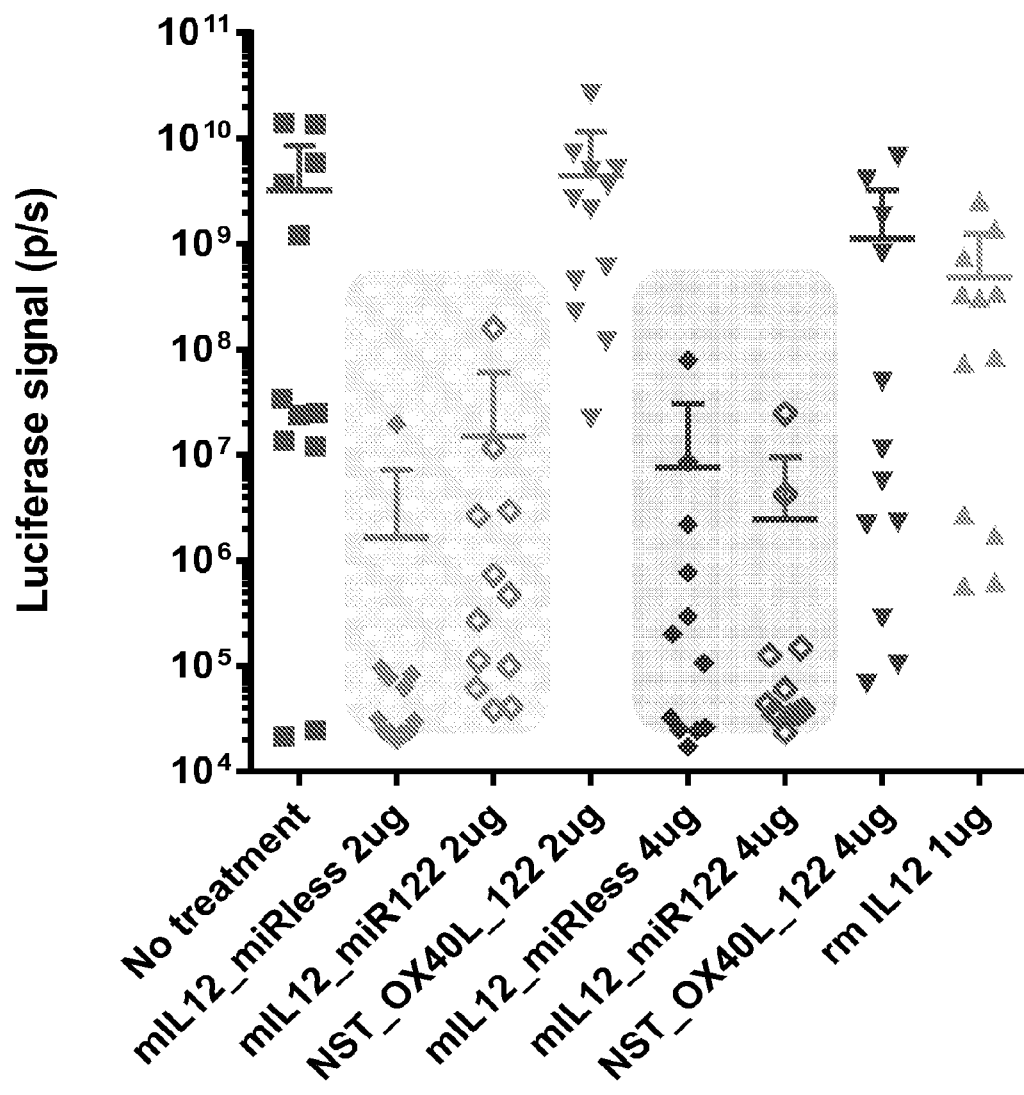
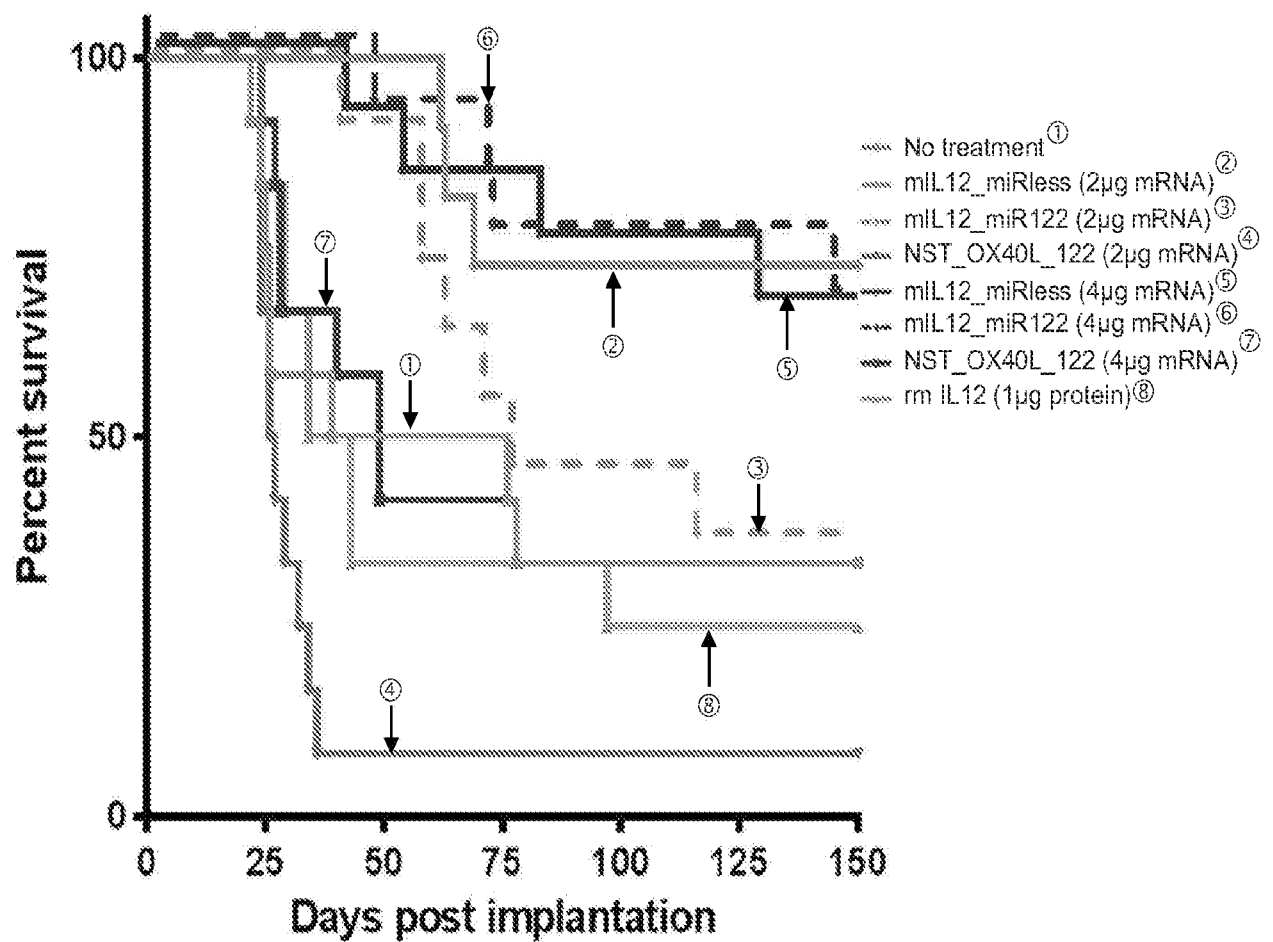


FIG. 11B

Survival to Day 150:



Protein	Length	Theoretical Minimum U (%)	Theoretical Minimum U (abs)		
IL12B WT	306	10.78%	99		
Nucleic Acid	Length			U Content v WT (%)	U Content v Theoretical Minimum (%)
	918	195	21.24%	100.00%	196.97%
	MAX	275	29.96%	141.03%	277.78%
	MIN	109	11.87%	55.90%	110.10%
	MEAN	173.25	18.87%	88.85%	175.00%
	MEDIAN	176	19.17%	90.26%	177.78%
	STD DEV	44.19	4.81%	22.66%	44.63%

FIG. 12A

Protein	Length	Theoretical Maximum G (%)	Theoretical Maximum G (abs)		
IL12B WT	306	40.74%	374		
Nucleic Acid	Length			G Content v WT(%)	G Content v Theoretical Maximum (%)
	918	243	26.47%	100.00%	64.97%
	MAX	374	40.74%	153.91%	100.00%
	MIN	219	23.86%	90.12%	58.56%
	MEAN	260.55	28.38%	107.22%	69.67%
	MEDIAN	249.5	27.18%	102.67%	66.71%
	STD DEV	35.31	3.85%	14.53%	9.44%

FIG. 12B

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Protein	Length	Theoretical Maximum C (%)	Theoretical Maximum C (abs)		
IL12B WT	306	41.61%	382		
Nucleic Acid	Length	C Content (abs)	C Content (%)	C Content v WT (%)	C Content v Theoretical Maximum (%)
	918	223	24.29%	100.00%	58.38%
	MAX	279	30.39%	125.11%	73.04%
	MIN	213	23.20%	95.52%	55.76%
	MEAN	238.05	25.93%	106.75%	62.32%
	MEDIAN	238.5	25.98%	106.96%	62.43%
	STD DEV	18.41	2.01%	8.26%	4.82%

FIG. 12C

Protein	Length	Theoretical Maximum GC (%)	Theoretical Maximum GC (abs)		
IL12B WT	306	65.25%	599		
Nucleic Acid	Length	GC Content (abs)	GC Content (%)	GC Content v WT (%)	GC Content v Theoretical Minimum (%)
	918	466	50.76%	100.00%	77.80%
	MAX	587	63.94%	125.97%	98.00%
	MIN	449	48.91%	96.35%	74.96%
	MEAN	498.6	54.31%	107.00%	83.24%
	MEDIAN	489.5	53.32%	105.04%	81.72%
	STD DEV	43.23	4.71%	9.28%	7.22%

FIG. 12D

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Protein	Length	Theoretical Minimum U (%)	Theoretical Minimum U (abs)		
IL12B WT	306	10.78%	99		
Nucleic Acid	Length	U Content (abs)	U Content (%)	U Content v WT (%)	U Content v Theoretical Minimum (%)
	918	195	21.24%	100.00%	196.97%
	MAX	149	16.23%	76.41%	150.51%
	MIN	131	14.27%	67.18%	132.32%
	MEAN	141.1	15.37%	72.36%	142.53%
	MEDIAN	141	15.36%	72.31%	142.42%
	STD DEV	5.37	0.58%	2.75%	5.42%

FIG. 13A

Protein	Length	Theoretical Maximum G (%)	Theoretical Maximum G (abs)		
IL12B WT	306	40.74%	374		
Nucleic Acid	Length	G Content (abs)	G Content (%)	G Content v WT (%)	G Content v Theoretical Minimum (%)
	918	243	26.47%	100.00%	64.97%
	MAX	294	32.03%	120.99%	78.61%
	MIN	279	30.39%	114.81%	74.60%
	MEAN	287.35	31.30%	118.25%	76.83%
	MEDIAN	288	31.37%	118.52%	77.01%
	STD DEV	4.63	0.50%	1.90%	1.24%

FIG. 13B

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Protein	Length	Theoretical Maximum C (%)	Theoretical Maximum C (abs)		
IL12B WT	306	41.61%	382		
Nucleic Acid	Length	C Content (abs)	C Content (%)	C Content v WT (%)	C Content v Theoretical Maximum (%)
	918	223	24.29%	100.00%	58.38%
	MAX	279	30.39%	125.11%	73.04%
	MIN	262	28.54%	117.49%	68.59%
	MEAN	270.15	29.43%	121.14%	70.72%
	MEDIAN	270	29.41%	121.08%	70.68%
	STD DEV	4.76	0.52%	2.13%	1.25%

FIG. 13C

Protein	Length	Theoretical Maximum GC (%)	Theoretical Maximum GC (abs)		
IL12B WT	306	65.25%	599		
Nucleic Acid	Length	GC Content (abs)	GC Content (%)	GC Content v WT (%)	GC Content v Theoretical Maximum (%)
	918	466	50.76%	100.00%	77.80%
	MAX	565	61.55%	121.24%	94.32%
	MIN	547	59.59%	117.38%	91.32%
	MEAN	557.5	60.73%	119.64%	93.07%
	MEDIAN	557	60.68%	119.53%	92.99%
	STD DEV	4.67	0.51%	1.00%	0.78%

FIG. 13D

Protein	Length	Theoretical Minimum U (%)	Theoretical Minimum U (abs)		
IL12A WT	197	12.86%	76		
Nucleic Acid	Length			U Content v WT (%)	U Content v Theoretical Minimum (%)
	591	151	25.55%	100.00%	198.68%
	MAX	189	31.98%	125.17%	248.68%
	MIN	83	14.04%	54.97%	109.21%
	MEAN	126.3	21.37%	83.64%	166.18%
	MEDIAN	125	21.15%	82.78%	164.47%
	STD DEV	28.03	4.74%	18.56%	36.88%

FIG. 14A

Protein	Length	Theoretical Maximum G (%)	Theoretical Maximum G (abs)		
IL12A WT	197	34.69%	205		
Nucleic Acid	Length			G Content v WT (%)	G Content v Theoretical Maximum (%)
	591	123	20.81%	100.00%	60.00%
	MAX	205	34.69%	166.67%	100.00%
	MIN	108	18.27%	87.80%	52.68%
	MEAN	134.7	22.79%	109.51%	65.71%
	MEDIAN	131	22.17%	106.50%	63.90%
	STD DEV	22.88	3.87%	18.60%	11.16%

FIG. 14B

Protein	Length	Theoretical Maximum C (%)	Theoretical Maximum C (abs)		
IL12A WT	197	43.82%	259		
Nucleic Acid	Length			C Content v WT (%)	C Content v Theoretical Maximum (%)
	591	138	23.35%	100.00%	53.28%
	MAX	191	32.32%	138.41%	73.75%
	MIN	140	23.69%	101.45%	54.05%
	MEAN	164.8	27.88%	119.42%	63.63%
	MEDIAN	164	27.75%	118.84%	63.32%
	STD DEV	12.22	2.07%	8.85%	4.72%

FIG. 14C

Protein	Length	Theoretical Maximum GC (%)	Theoretical Maximum GC (abs)		
IL12A WT	197	62.27%	368		
Nucleic Acid	Length			GC Content v WT (%)	GC Content v Theoretical Maximum (%)
	591	261	44.16%	100.00%	70.92%
	MAX	359	60.74%	137.55%	97.55%
	MIN	265	44.84%	101.53%	72.01%
	MEAN	299.5	50.68%	114.75%	81.39%
	MEDIAN	293	49.58%	112.26%	79.62%
	STD DEV	29.23	4.95%	11.20%	7.94%

FIG. 14D

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Protein	Length	Theoretical Minimum U (%)	Theoretical Minimum U (abs)		
IL12A WT	197	12.86%	76		
Nucleic Acid	Length			U Content v WT (%)	U Content v Theoretical Maximum (%)
	591	151	25.55%	100.00%	198.68%
	MAX	109	18.44%	72.19%	143.42%
	MIN	95	16.07%	62.91%	125.00%
	MEAN	101.9	17.24%	67.48%	134.08%
	MEDIAN	102	17.26%	67.55%	134.21%
	STD DEV	4.27	0.72%	2.83%	5.61%

FIG. 15A

Protein	Length	Theoretical Maximum G (%)	Theoretical Maximum G (abs)		
IL12A WT	197	34.69%	205		
Nucleic Acid	Length			G Content v WT (%)	G Content v Theoretical Maximum (%)
	591	123	20.81%	100.00%	60.00%
	MAX	160	27.07%	130.08%	78.05%
	MIN	147	24.87%	119.51%	71.71%
	MEAN	153.45	25.96%	124.76%	74.85%
	MEDIAN	153.5	25.97%	124.80%	74.88%
	STD DEV	3.38	0.57%	2.75%	1.65%

FIG. 15B

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Protein	Length	Theoretical Maximum C (%)	Theoretical Maximum C (abs)		
IL12A WT	197	43.82%	259		
Nucleic Acid	Length			C Content v WT (%)	C Content v Theoretical Maximum (%)
	591	138	23.35%	100.00%	53.28%
	MAX	195	32.99%	141.30%	75.29%
	MIN	179	30.29%	129.71%	69.11%
	MEAN	187.8	31.78%	136.09%	72.51%
	MEDIAN	187.5	31.73%	135.87%	72.39%
	STD DEV	4.05	0.68%	2.93%	1.56%

FIG. 15C

Protein	Length	Theoretical Maximum GC (%)	Theoretical Maximum GC (abs)		
IL12A WT	197	62.27%	368		
Nucleic Acid	Length			GC Content v WT (%)	GC Content v Theoretical Maximum (%)
	591	261	44.16%	100.00%	70.92%
	MAX	350	59.22%	134.10%	95.11%
	MIN	337	57.02%	129.12%	91.58%
	MEAN	341.25	57.74%	130.75%	92.73%
	MEDIAN	340	57.53%	130.27%	92.39%
	STD DEV	4.08	0.69%	1.56%	1.11%

FIG. 15D

IL12B (w/o SP) G6 SEQUENCES					
Sequence	GC	GC 1st	GC 2nd	GC 3rd	
IL12B_WT	50.76	48.04	43.14	61.11	
MAX	61.55	51.63	43.14	91.83	
MIN	59.59	49.35	43.14	85.29	
MEAN	60.73	50.33	43.14	88.73	
MEDIAN	60.68	50.33	43.14	88.725	
STD DEV	0.51	0.61	0.00	1.60	

FIG. 16B

IL12B (w/o SP) G6 SEQUENCES					
Sequence	GC	GC 1st	GC 2nd	GC 3rd	
IL12A_WT	44.16	45.18	35.53	51.78	
MAX	59.22	51.27	35.53	93.91	
MIN	57.02	46.7	35.53	85.79	
MEAN	57.74	48.45	35.53	89.24	
MEDIAN	57.53	48.22	35.53	88.83	
STD DEV	0.69	1.02	0.00	1.80	

FIG. 16D

IL12B (w/o SP) G5 SEQUENCES					
Sequence	GC	GC 1st	GC 2nd	GC 3rd	
IL12B_WT	50.76	48.04	43.14	61.11	
MAX	63.94	50.65	43.14	100	
MIN	48.91	48.04	43.14	54.25	
MEAN	54.31	49.46	43.14	70.34	
MEDIAN	53.325	49.35	43.14	66.34	
STD DEV	4.71	0.80	0.00	14.42	

FIG. 16A

IL12A (w/o SP) G5 SEQUENCES					
Sequence	GC	GC 1st	GC 2nd	GC 3rd	
IL12A_WT	44.16	45.18	35.53	51.78	
MAX	60.74	48.73	35.53	100	
MIN	44.84	43.65	35.53	52.28	
MEAN	50.68	46.35	35.53	70.15	
MEDIAN	49.58	46.7	35.53	67.765	
STD DEV	4.94	1.44	0.00	14.35	

FIG. 16C

FIG. 17A

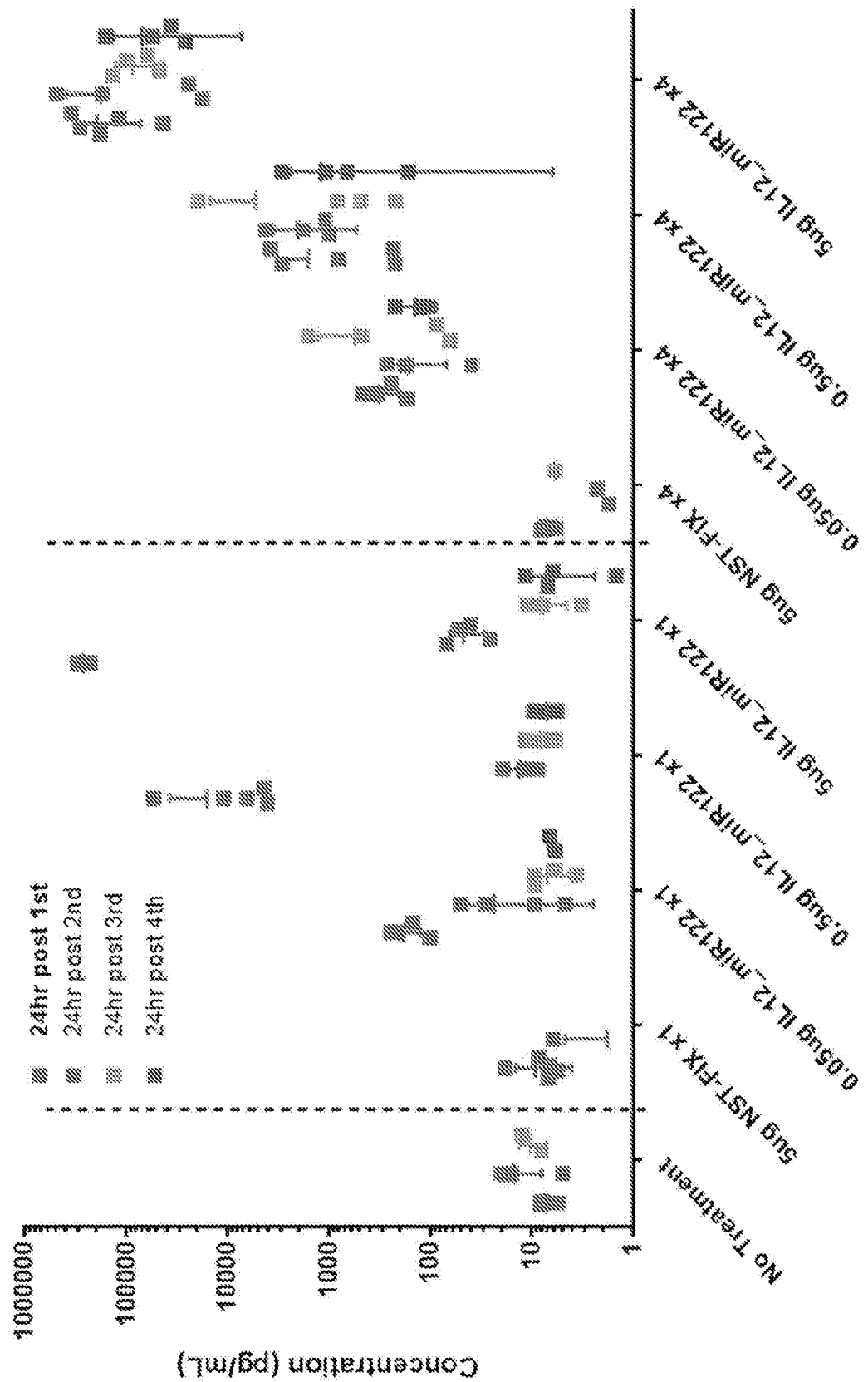


FIG. 17B

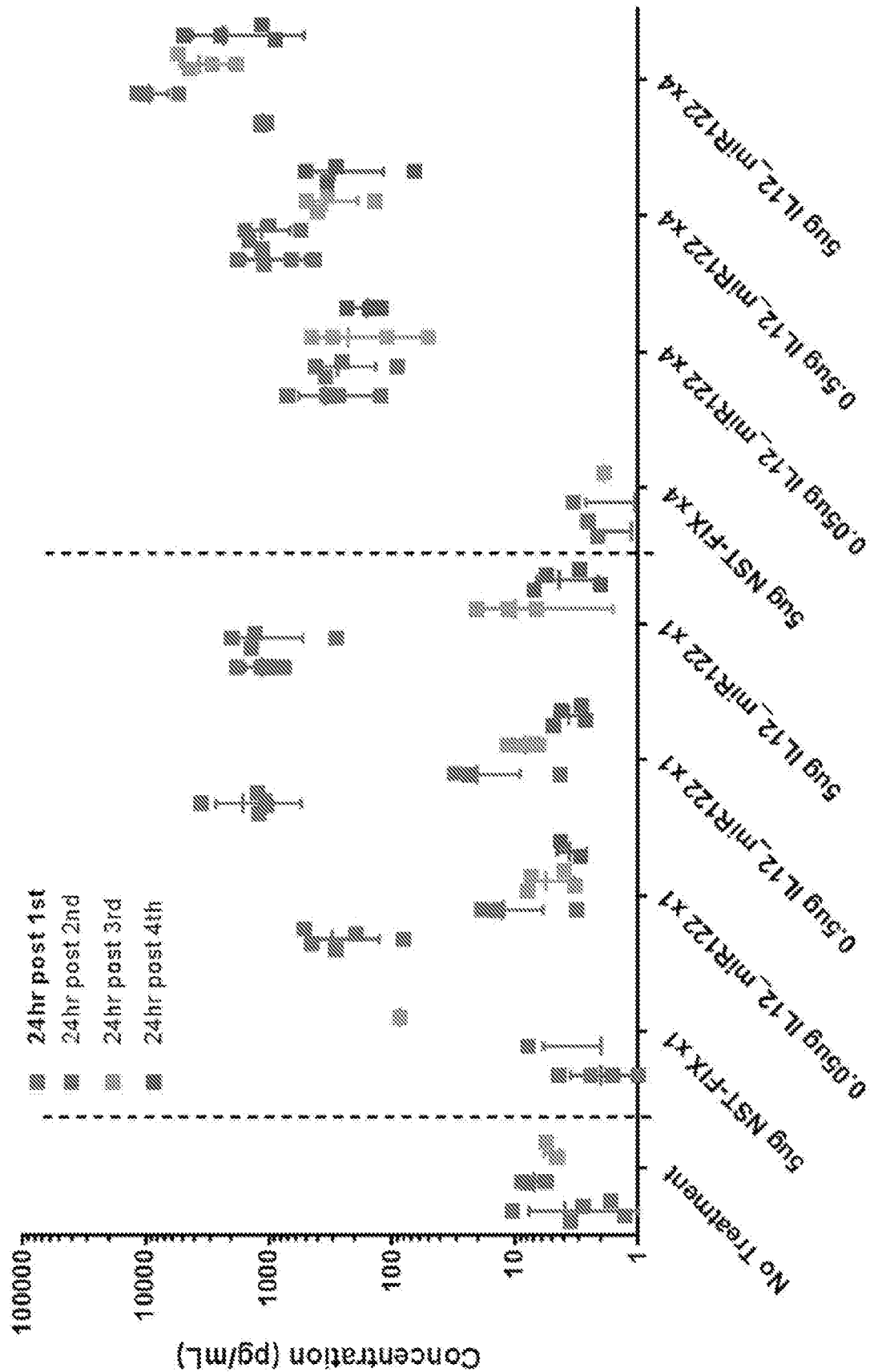


FIG. 18A

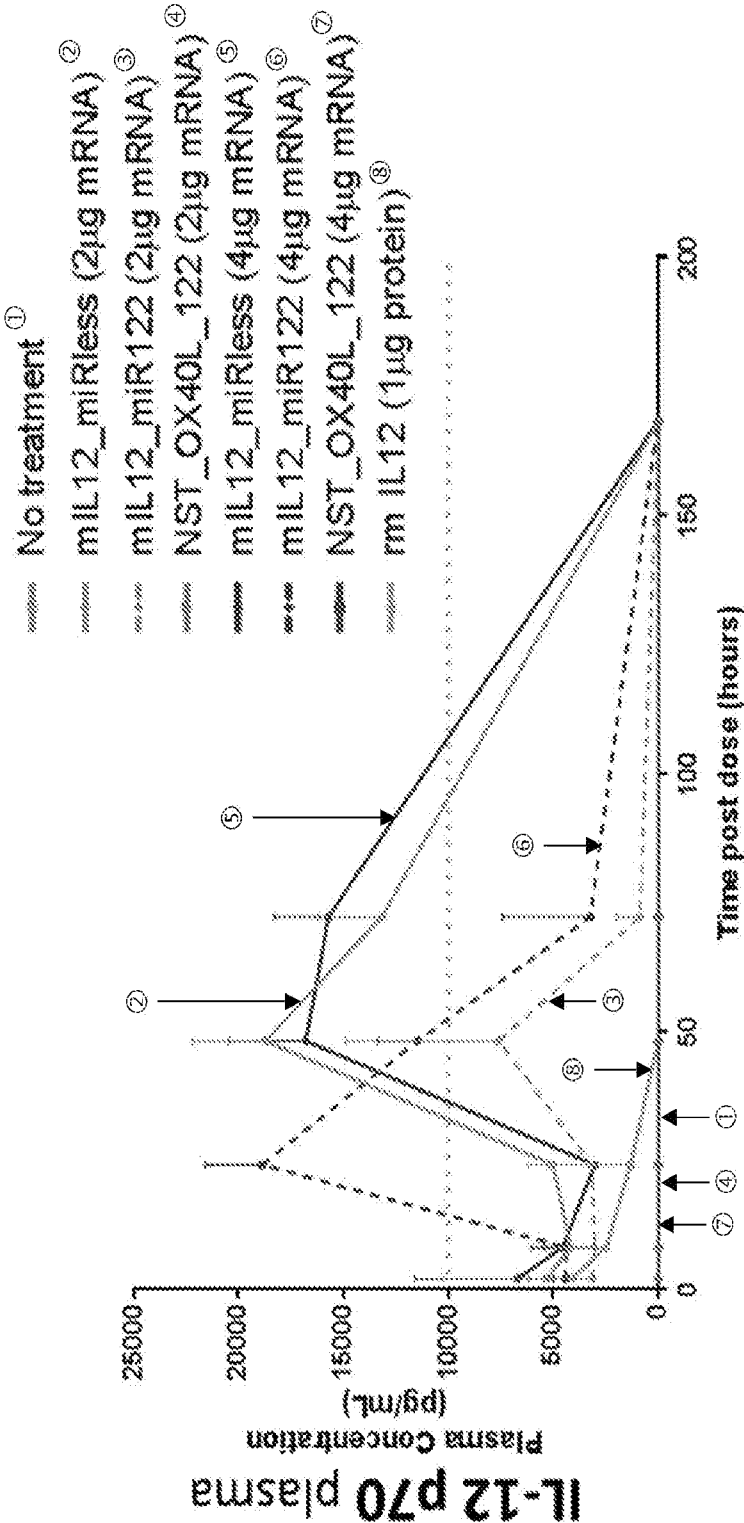


FIG. 18B

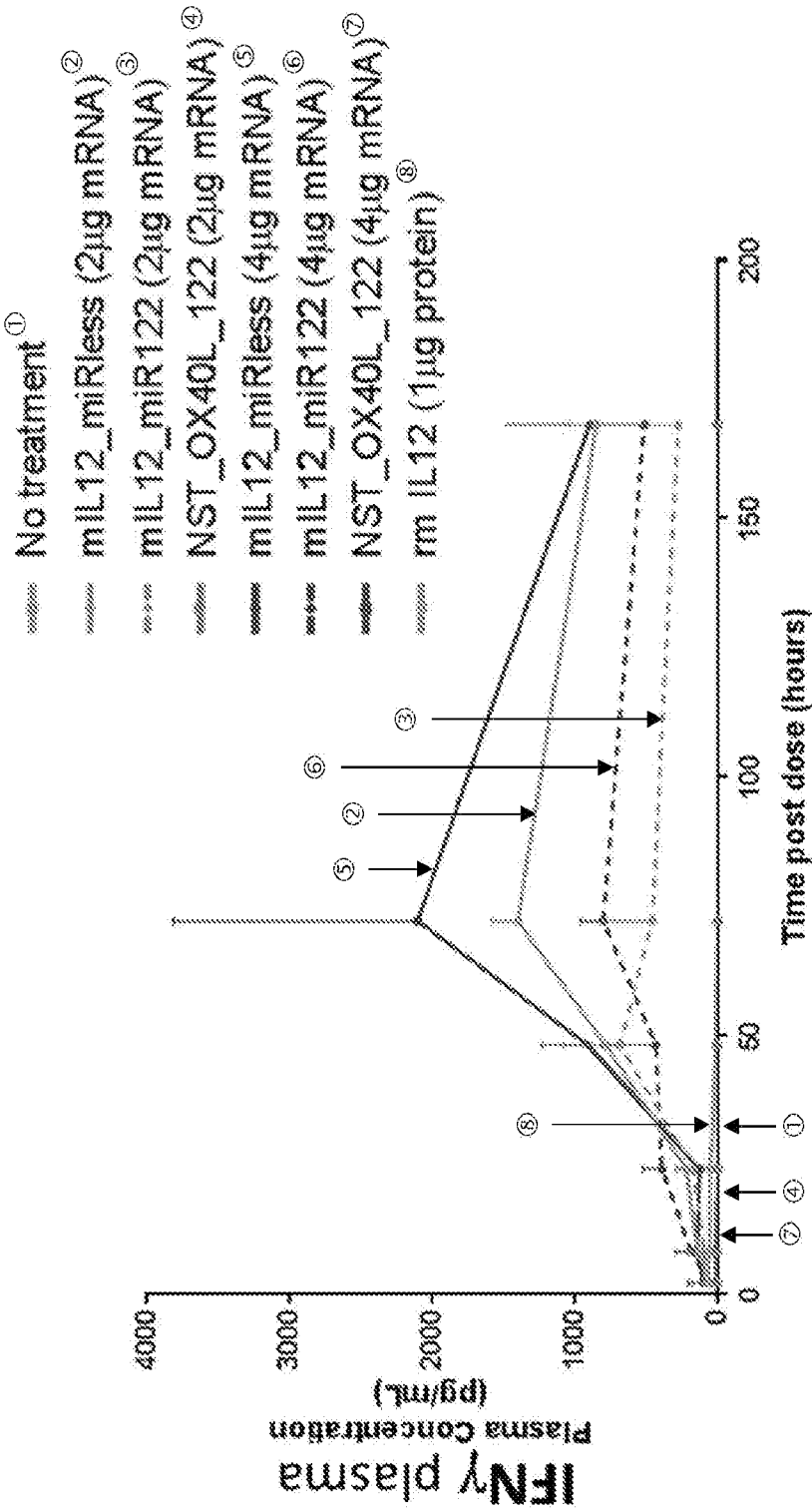


FIG. 19A

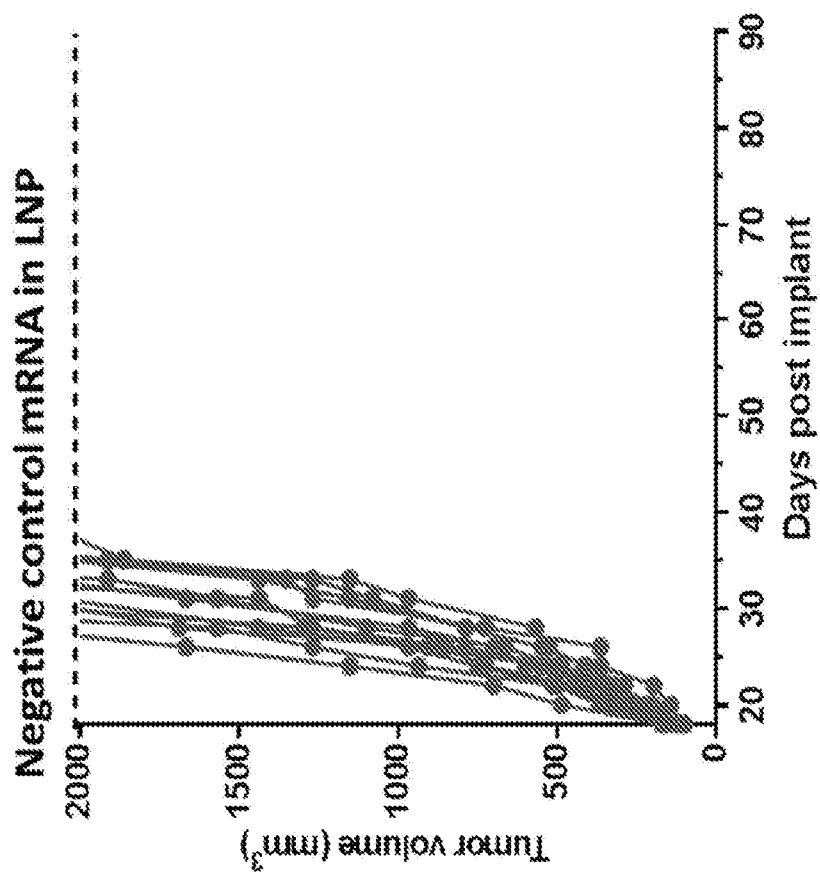


FIG. 19B

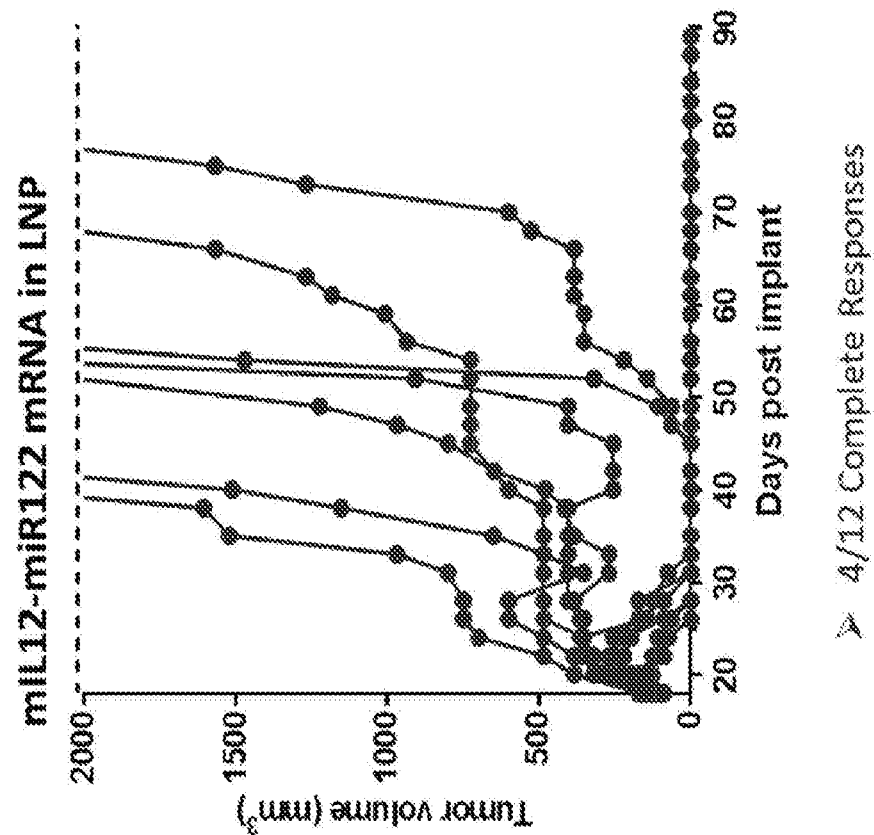
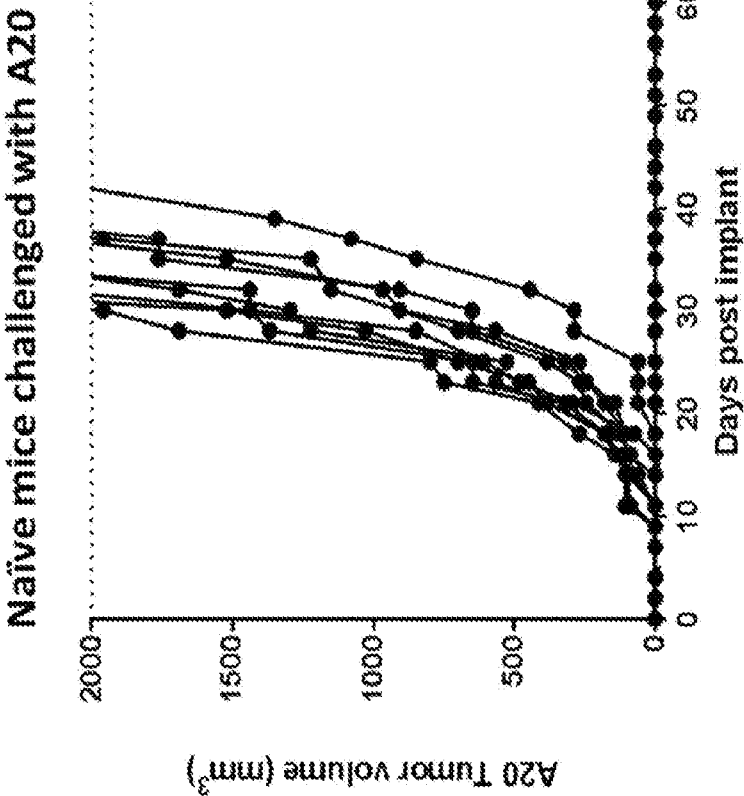
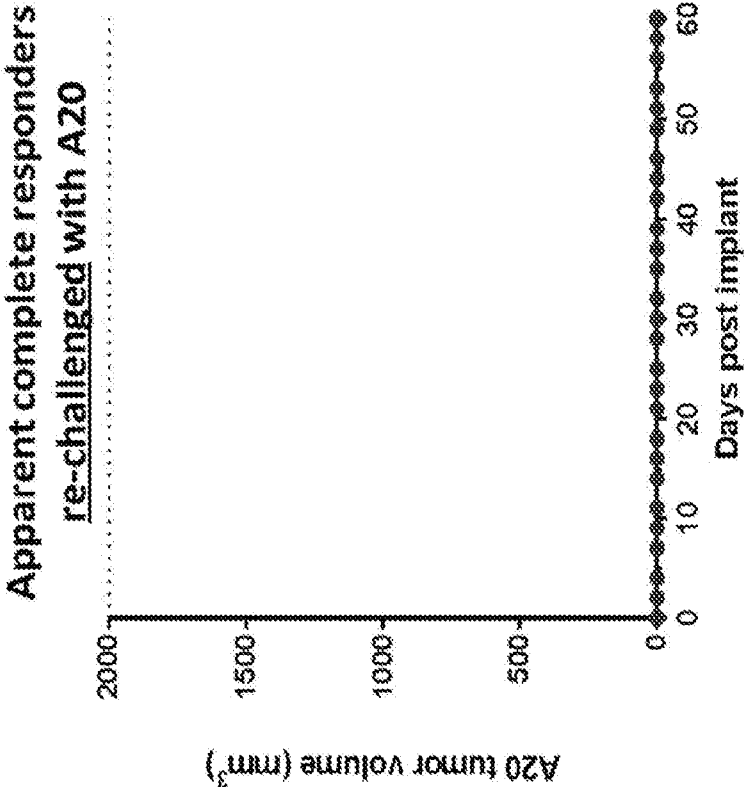


FIG. 19C



9/10 newly implanted tumors hit
end point by Day 45

FIG. 19D



➤ 0/4 newly implanted tumors grew

FIG. 20A
0.05 μ g IL12

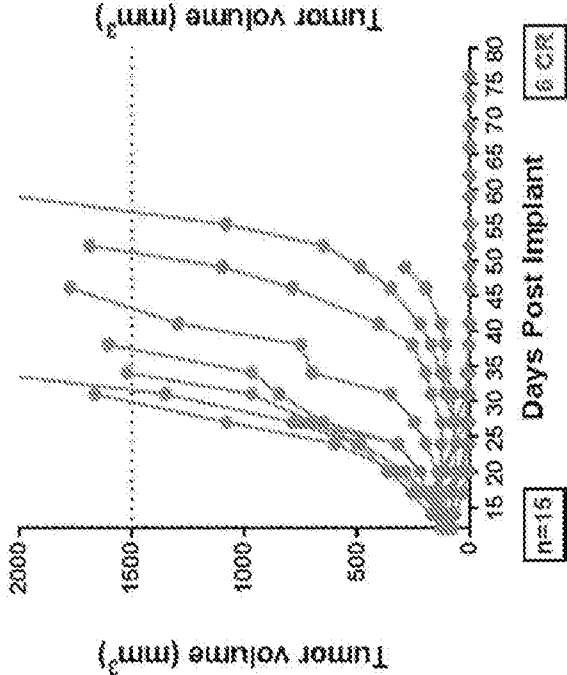


FIG. 20B
0.5 μ g IL12

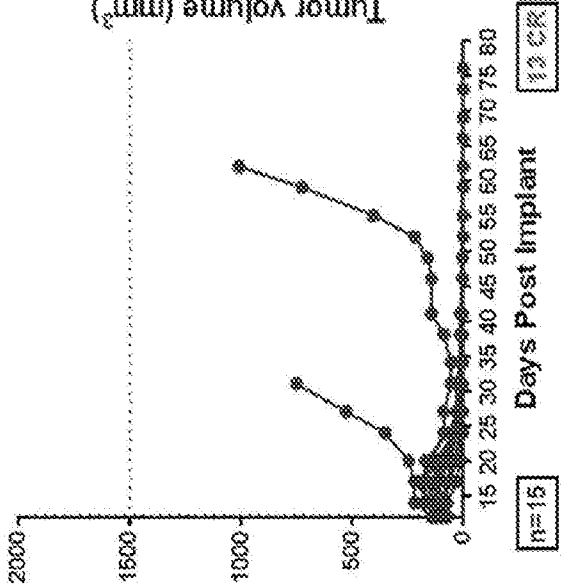
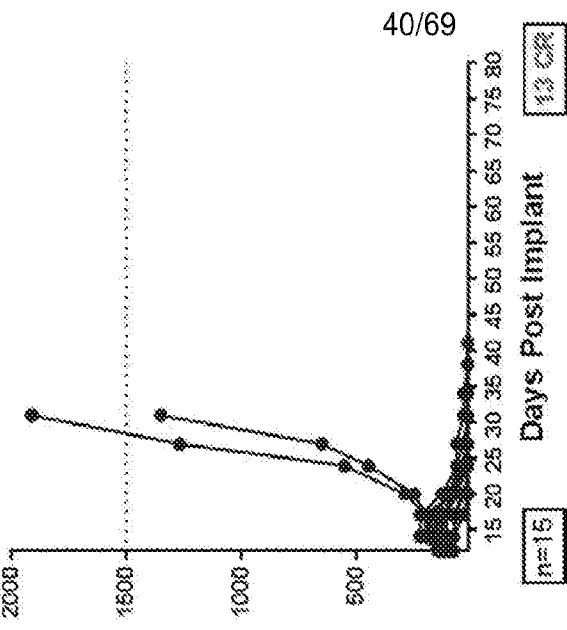


FIG. 20C
5 μ g IL12



40/69

Negative controls:

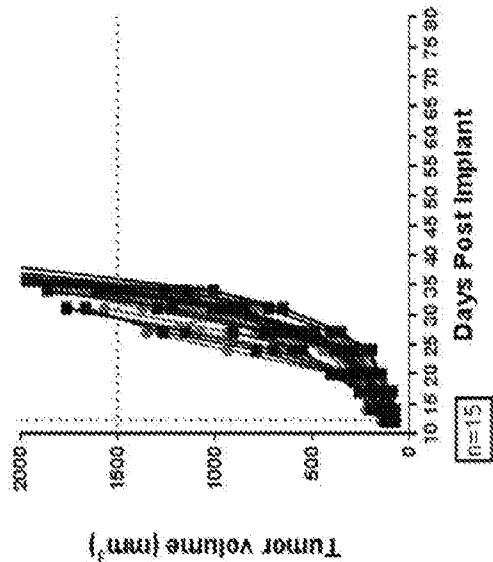


FIG. 20D

◆ NST-FIX 5ug iTu qdx1
■ Untreated

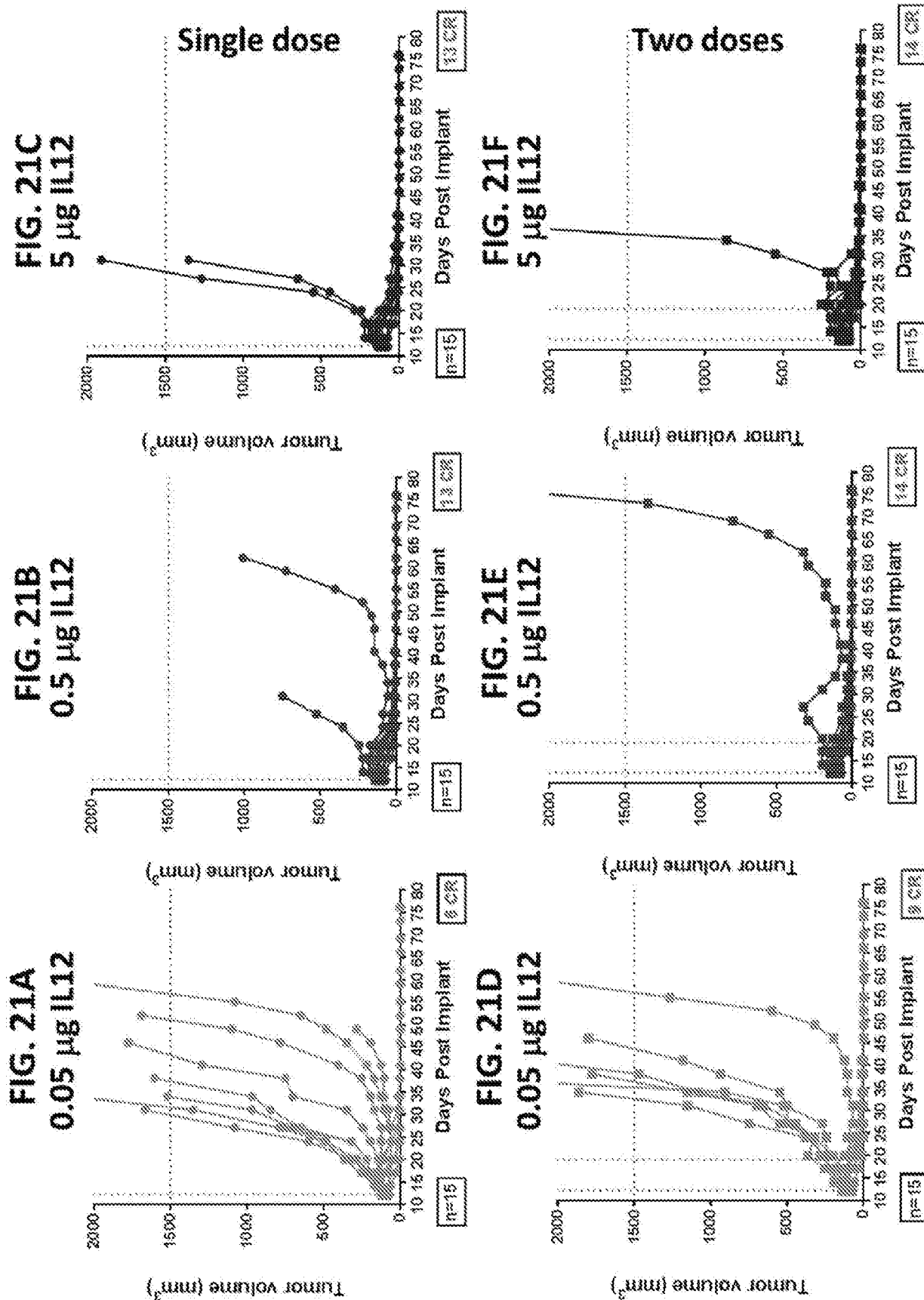


FIG. 22

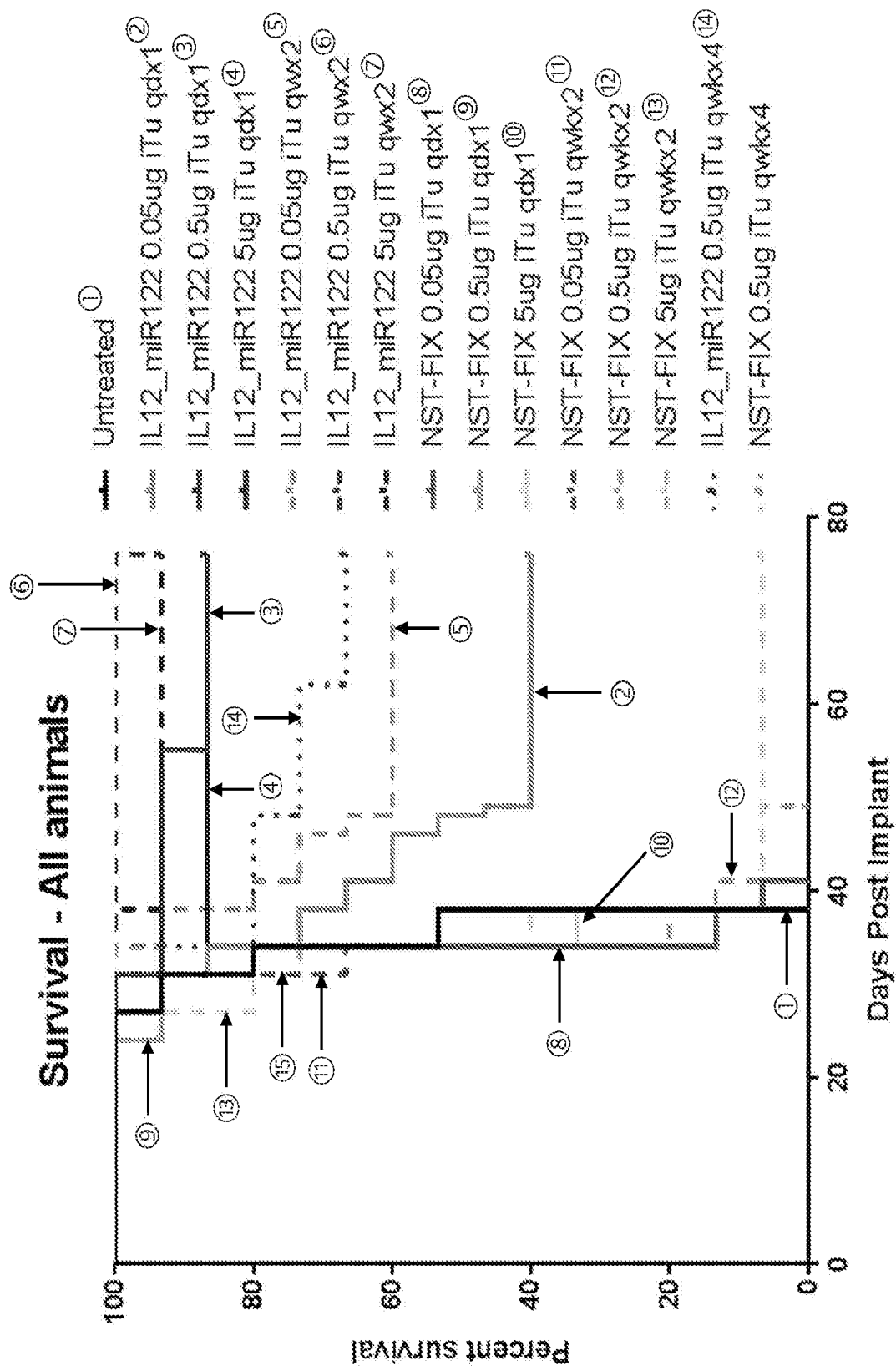
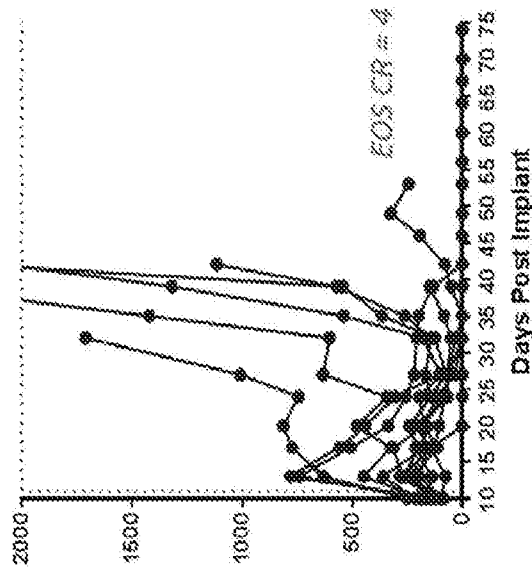
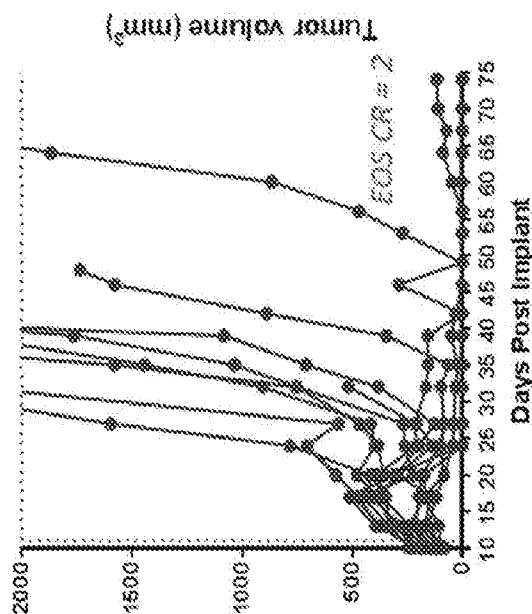


FIG. 23C
5 μ g IL12



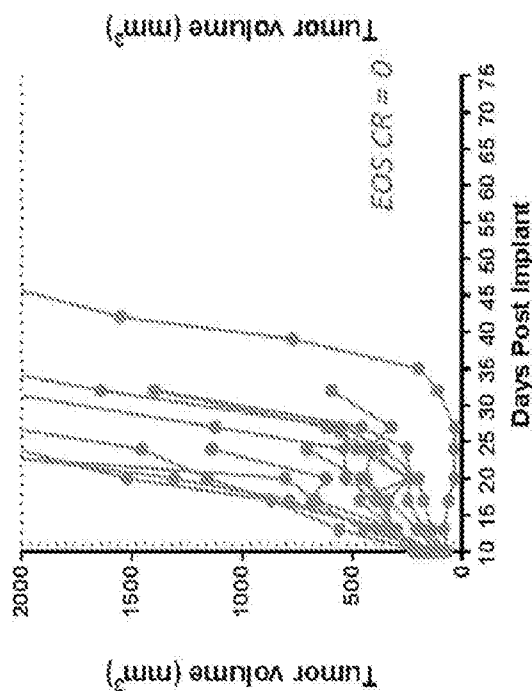
*EOS CR is End of
Study Complete
Response*

FIG. 23B
0.5 μ g IL12



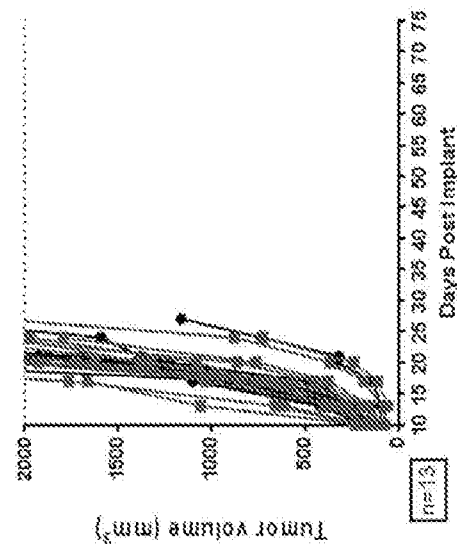
— NST-FIX_5ug_single dose
— Untreated

FIG. 23A
0.05 μ g IL12



Negative controls:

FIG. 23D



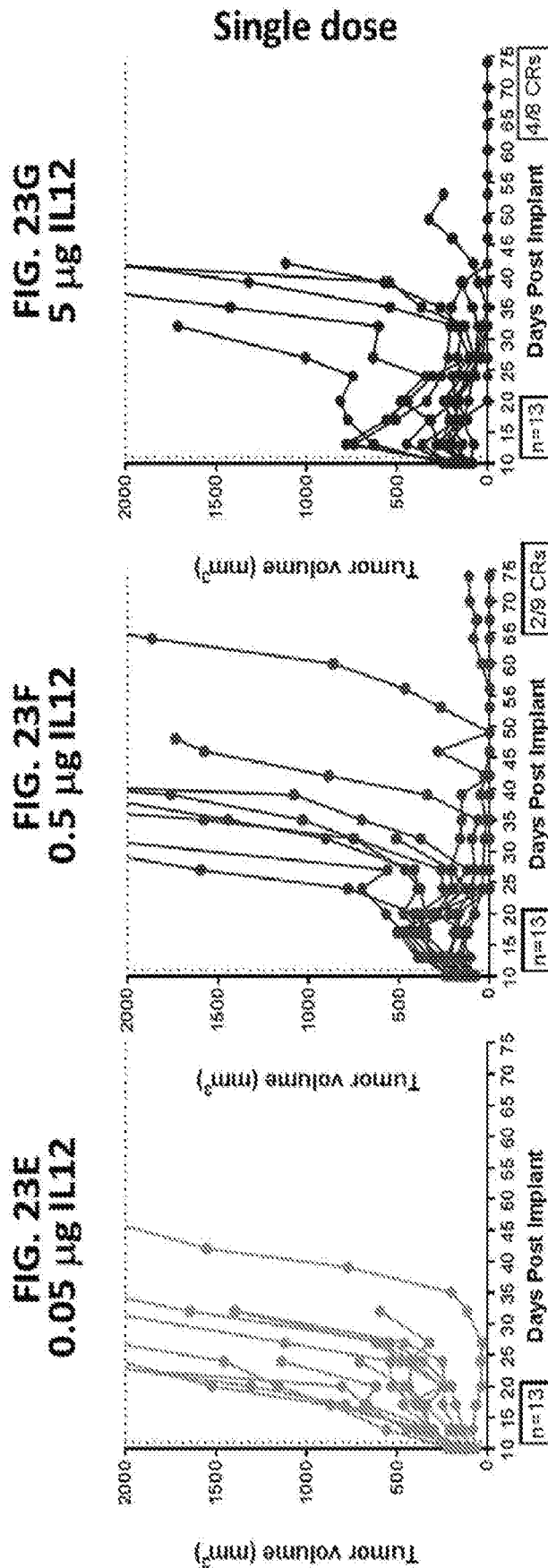
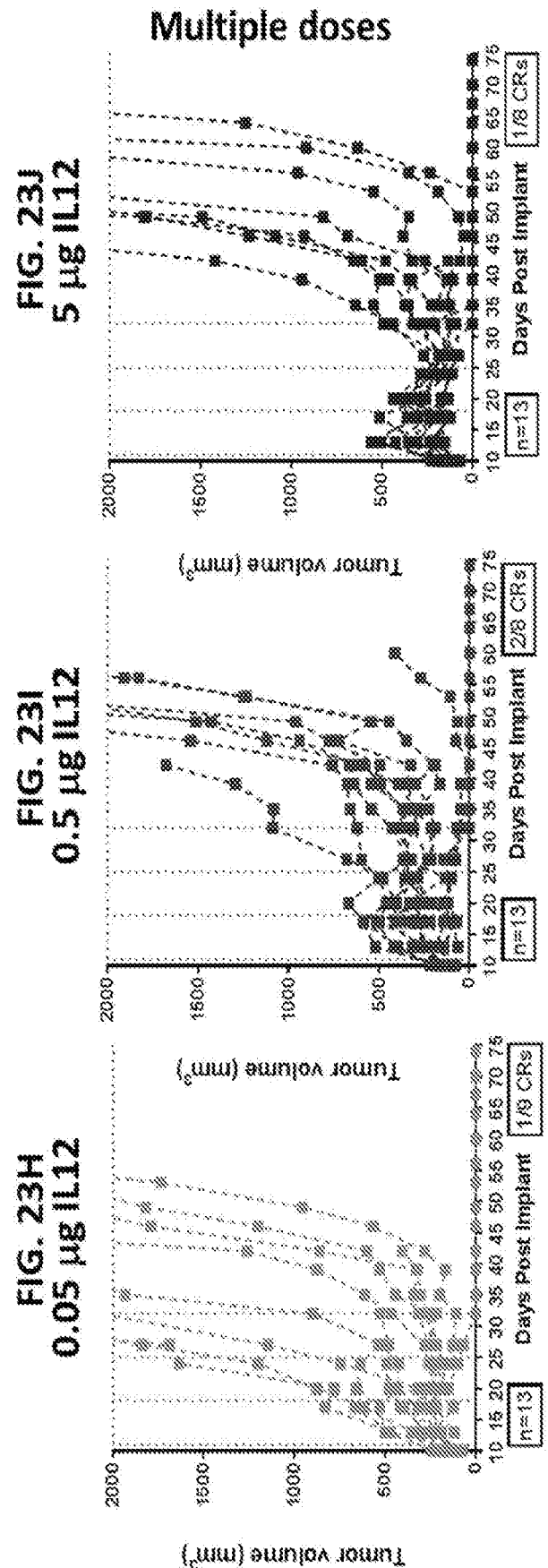


FIG. 24

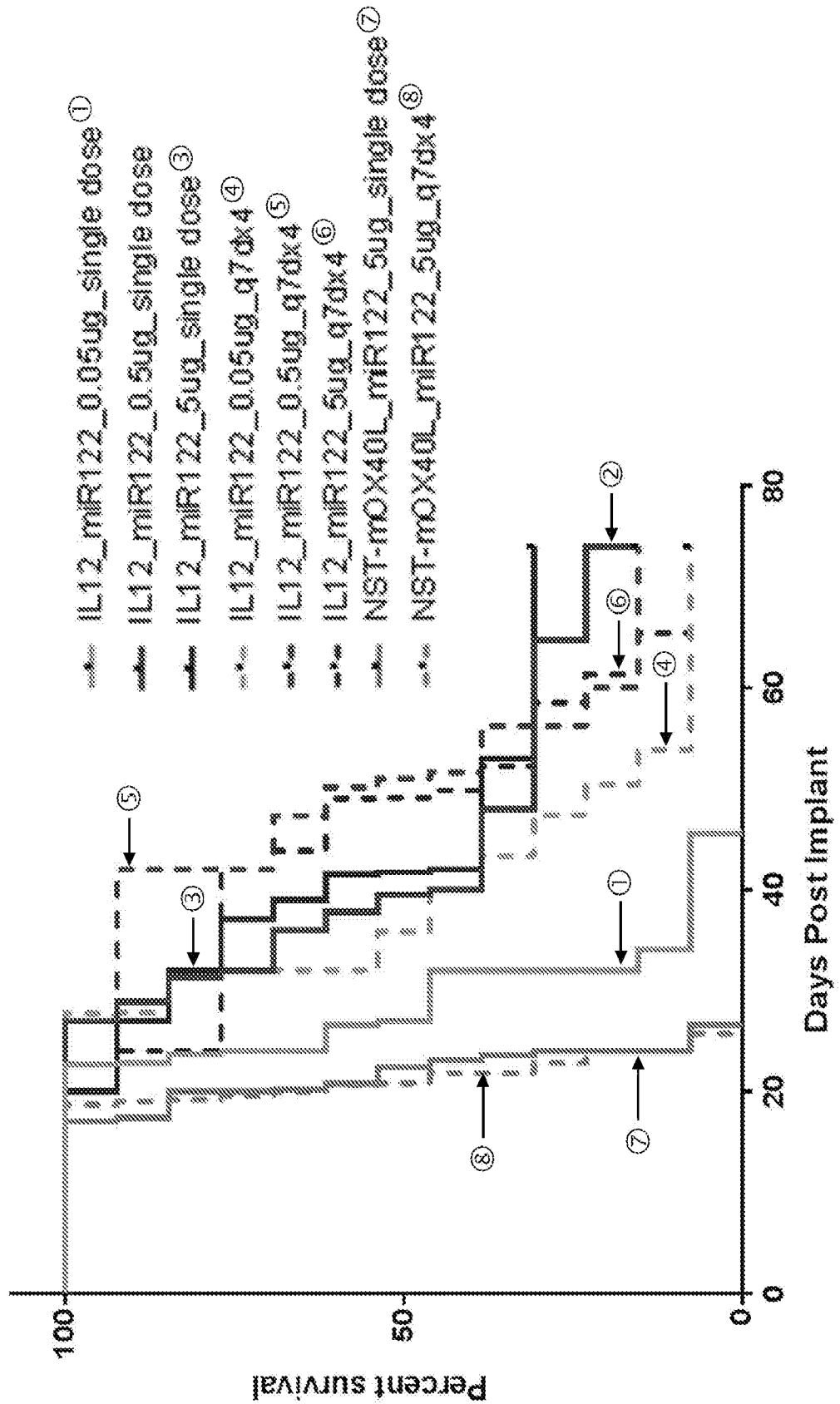


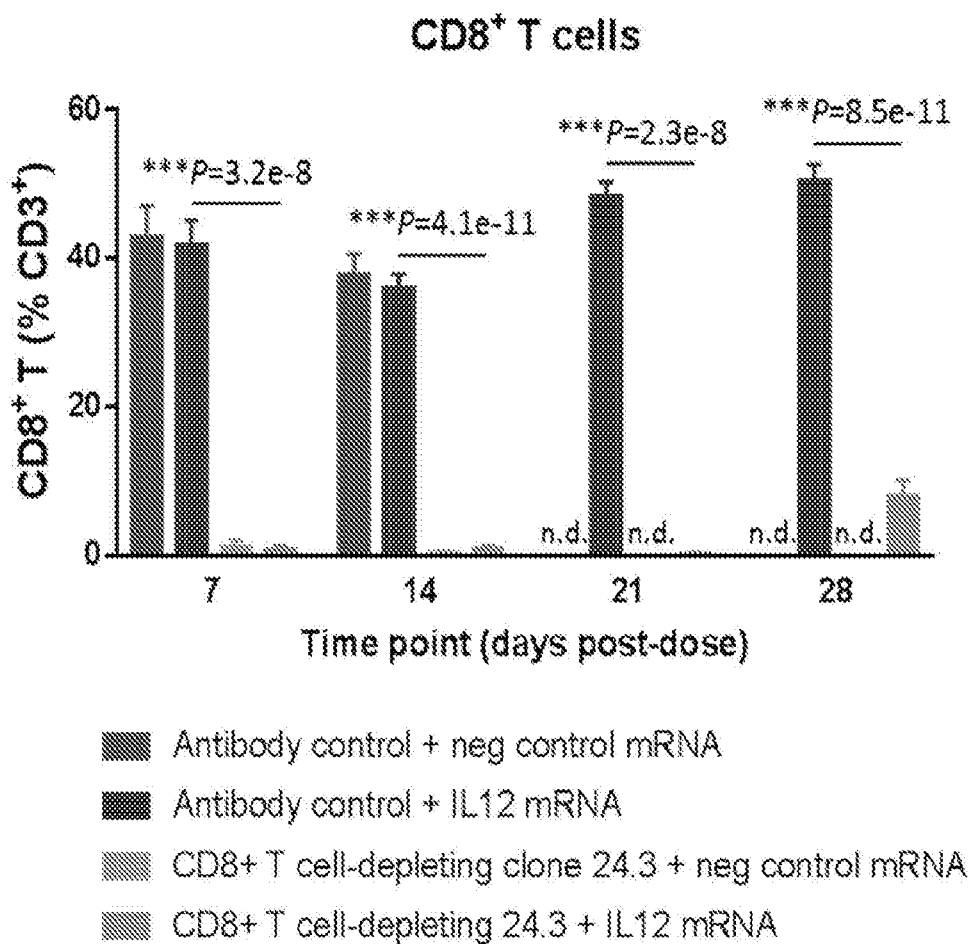
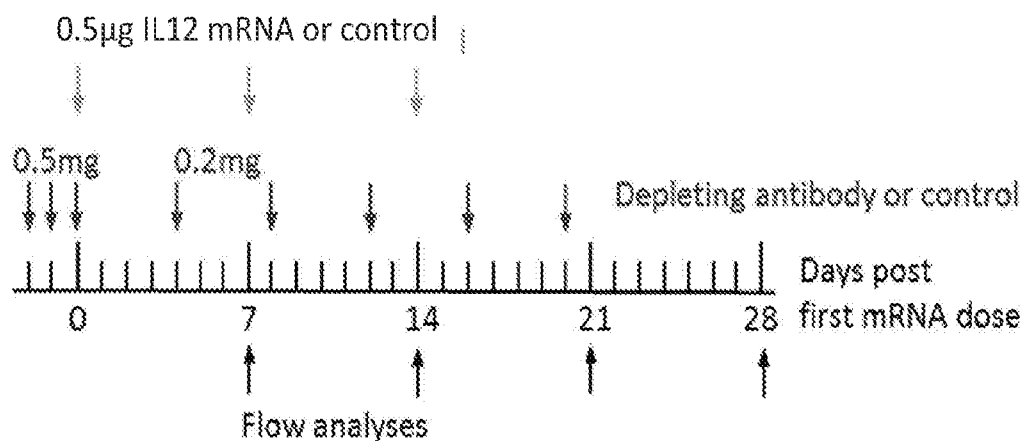
FIG. 25

FIG. 26A

FIG. 26B

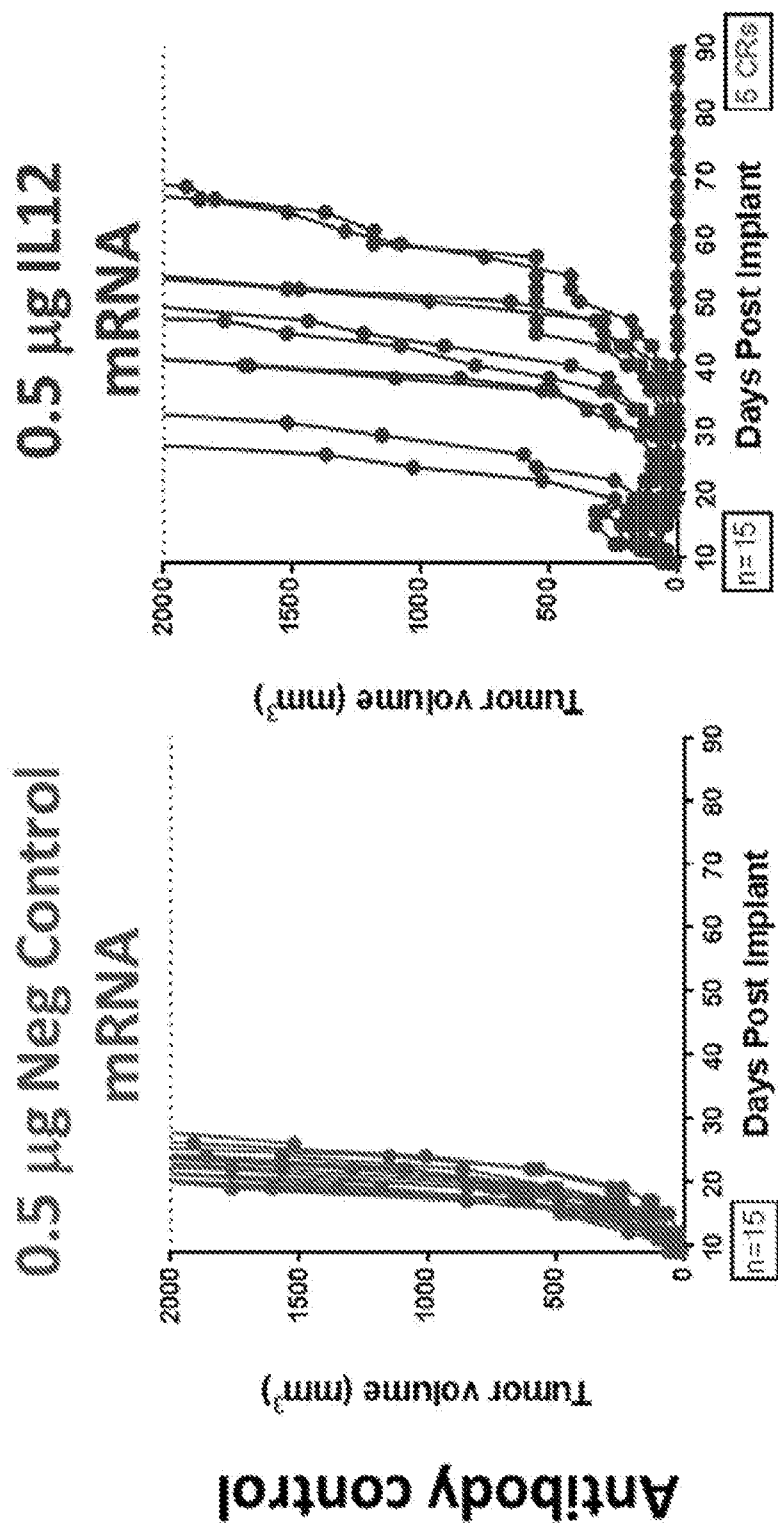


FIG. 26C

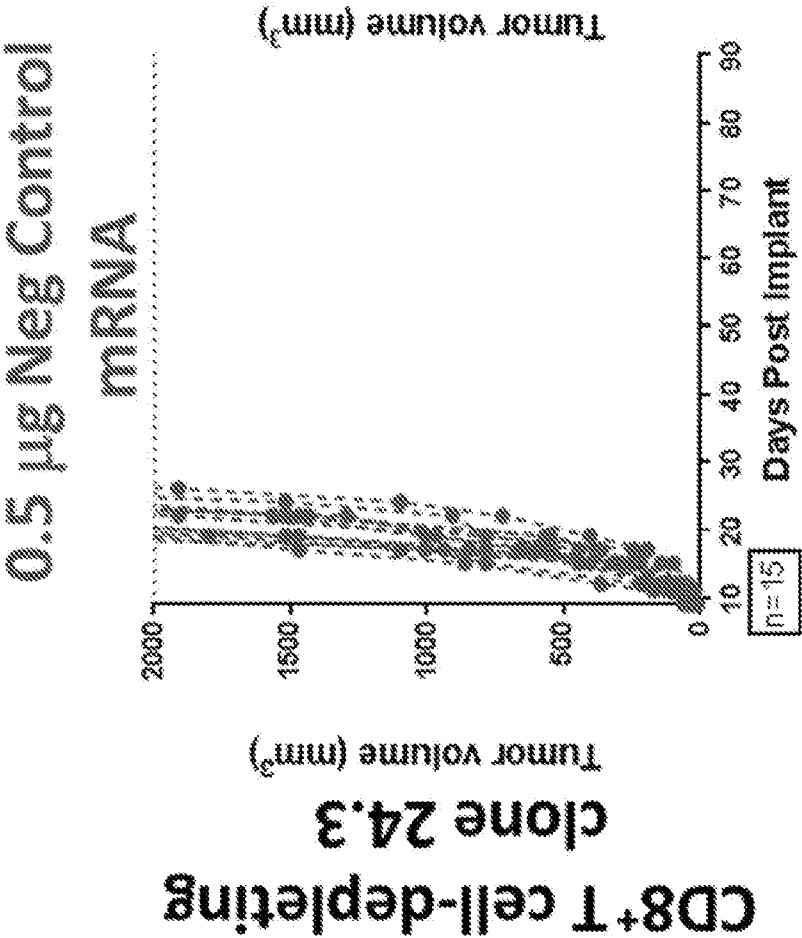


FIG. 26D

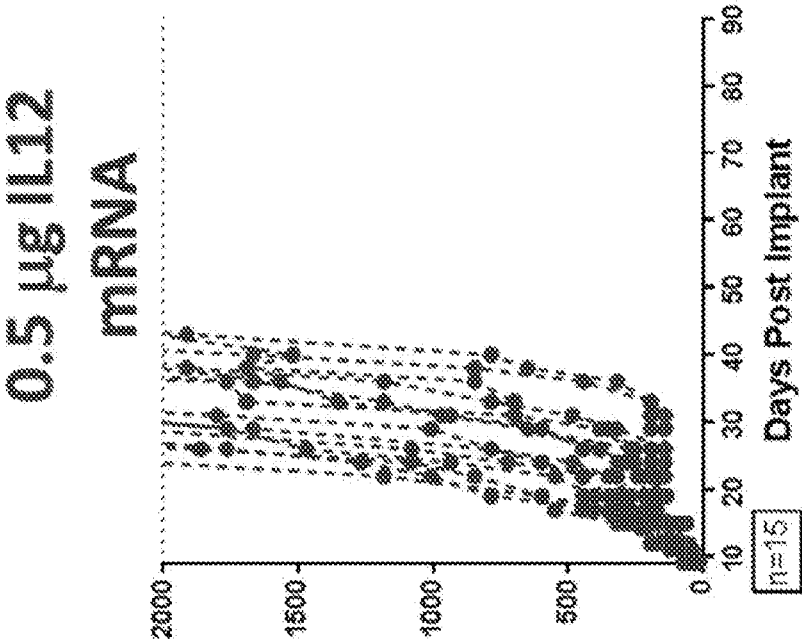


FIG. 26E

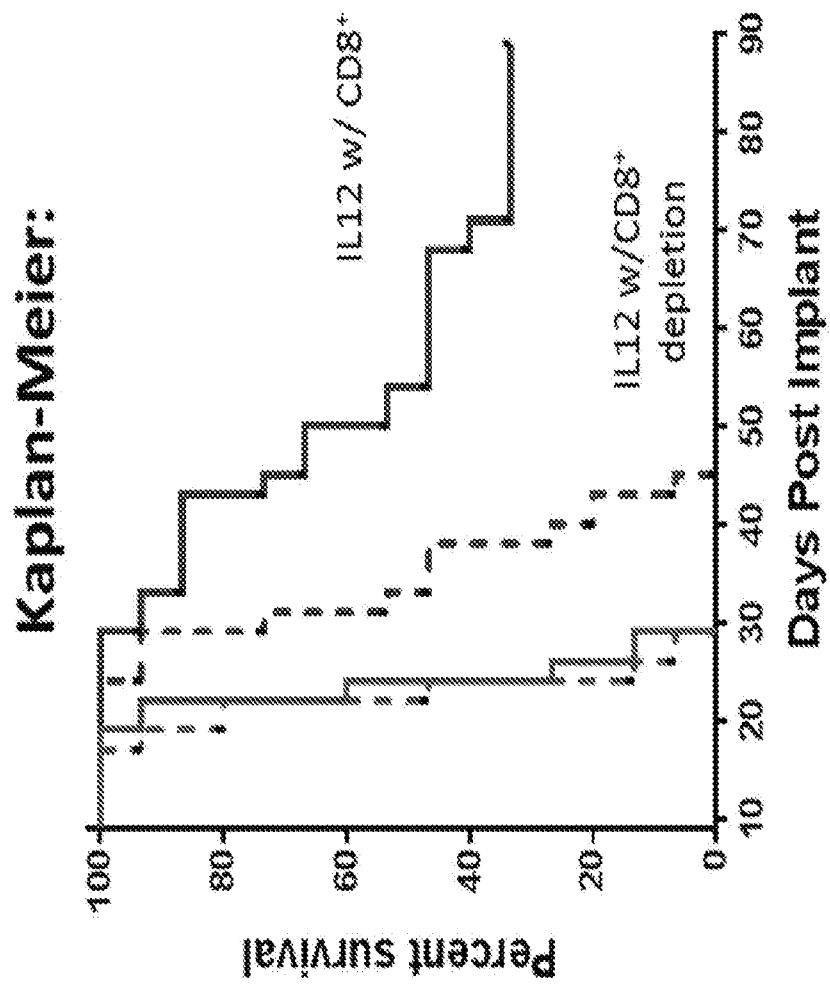


FIG. 27A

MC38-R

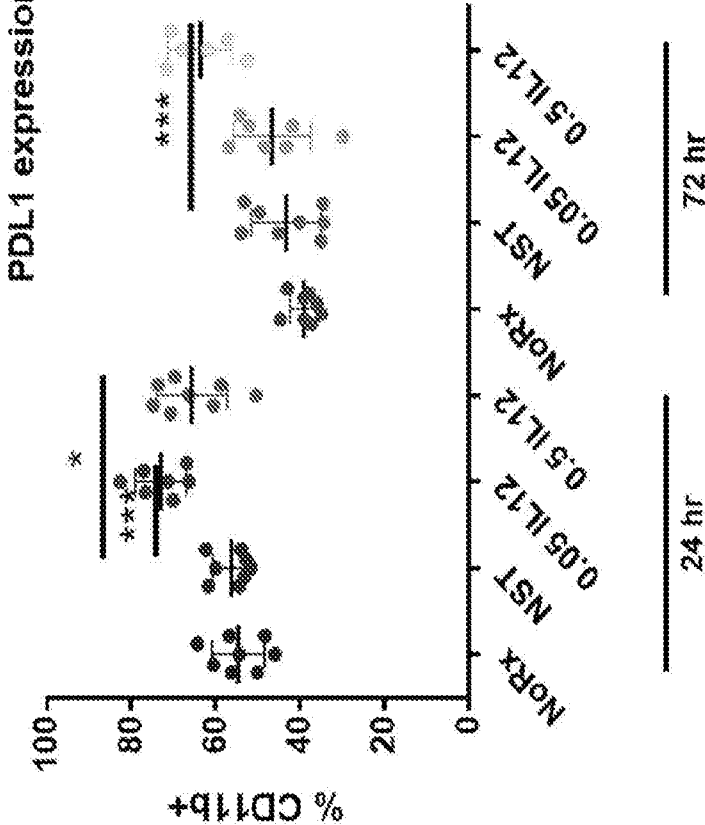


FIG. 27B

B16F10-AP3

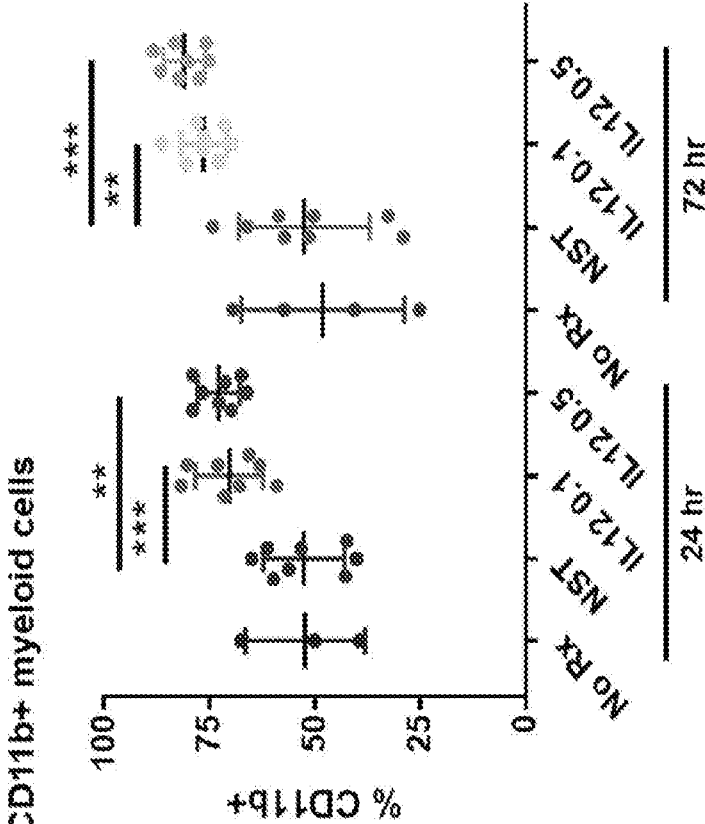


FIG. 28A

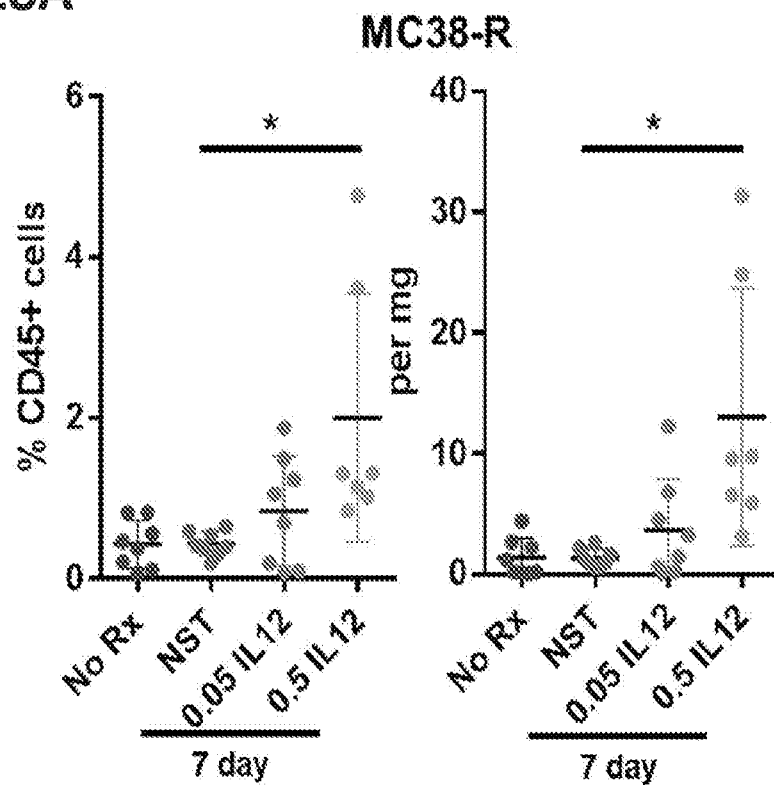


FIG. 28B

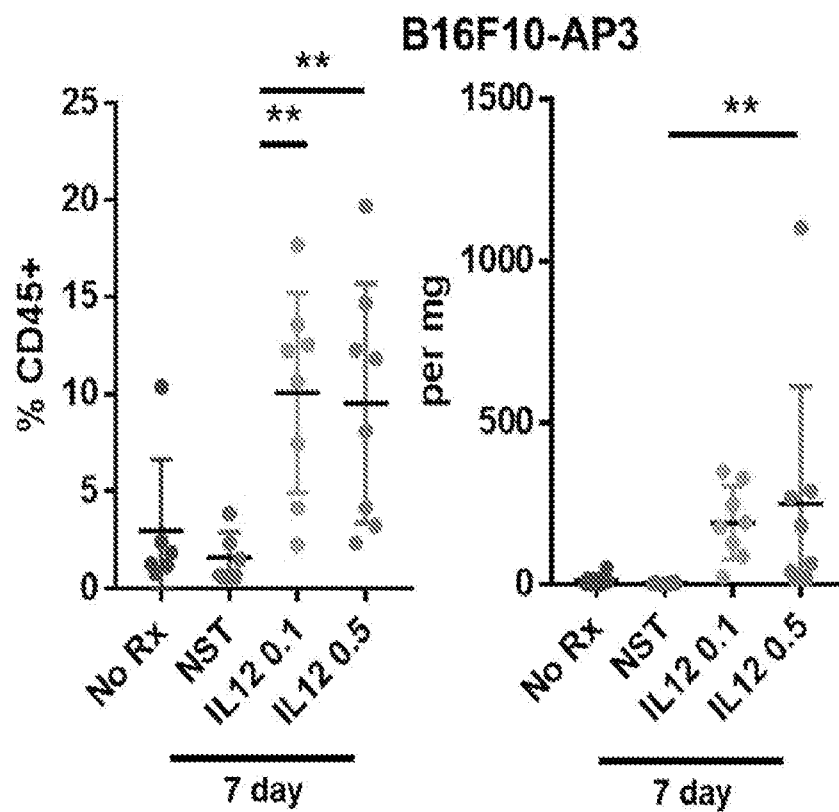


FIG. 29A

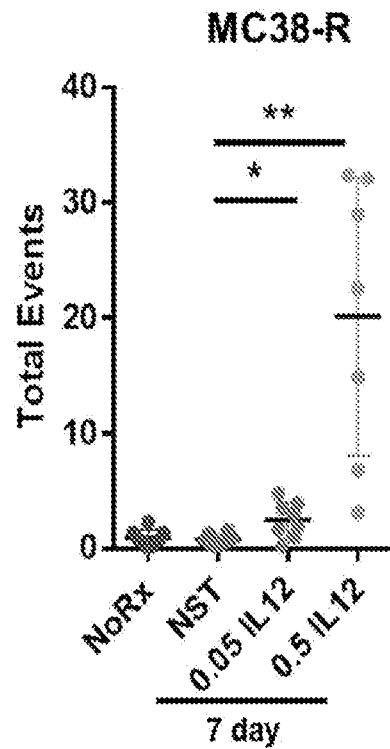


FIG. 29B

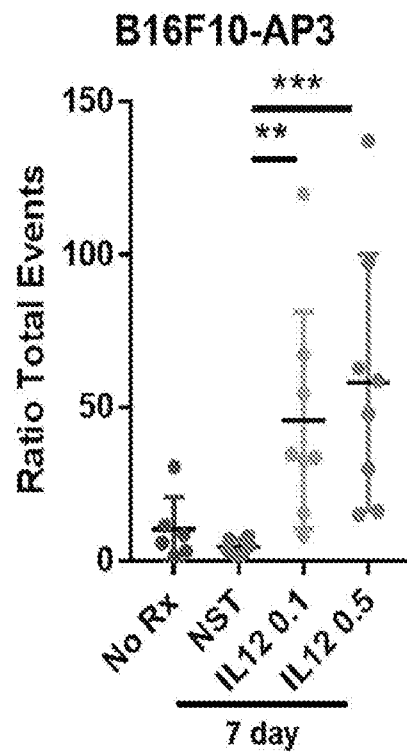


FIG. 30A

MC38-R

CD69 early activation marker expression on CD8+ T cells

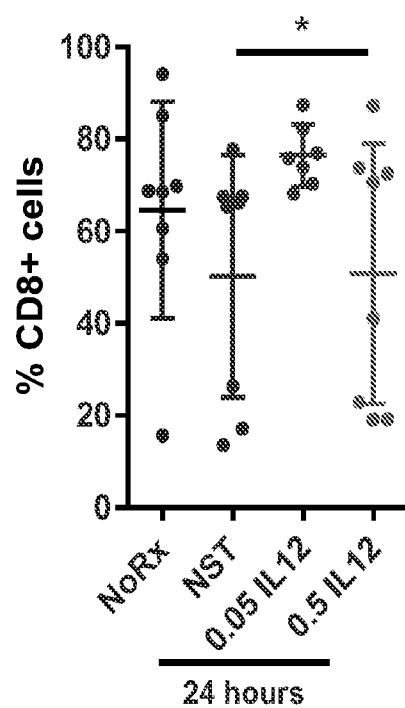


FIG. 30B

B16F10-AP3

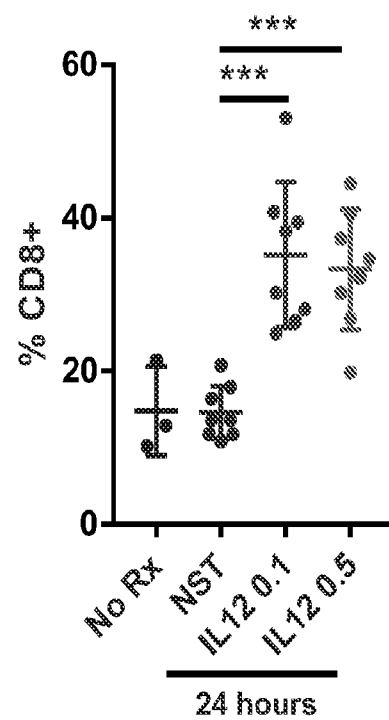


FIG. 31B

B16F10-AP3

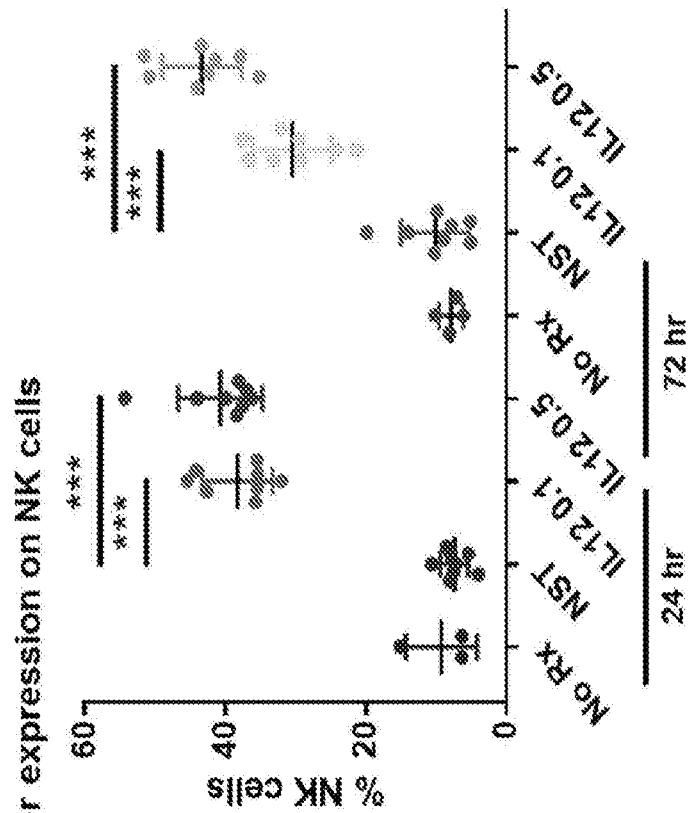


FIG. 31A

MC38-R

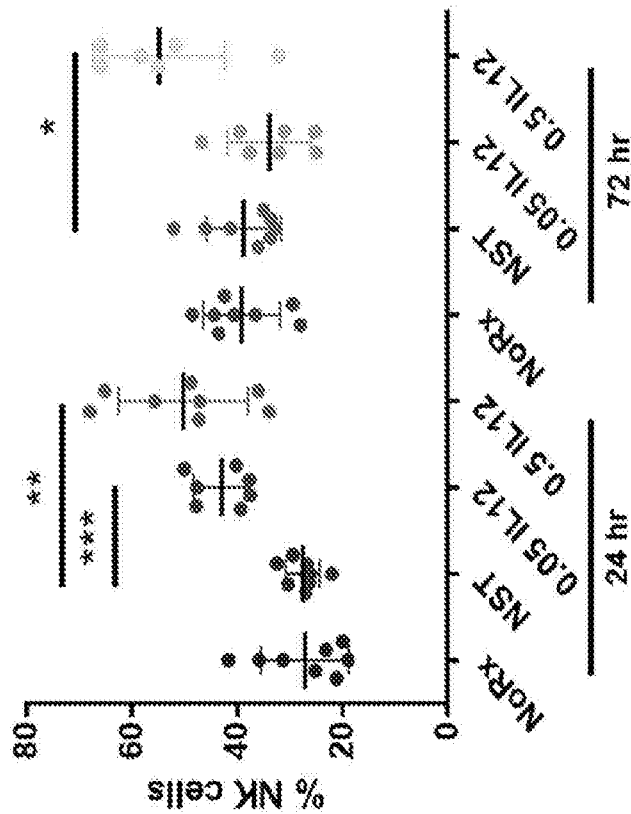


Fig. 32C

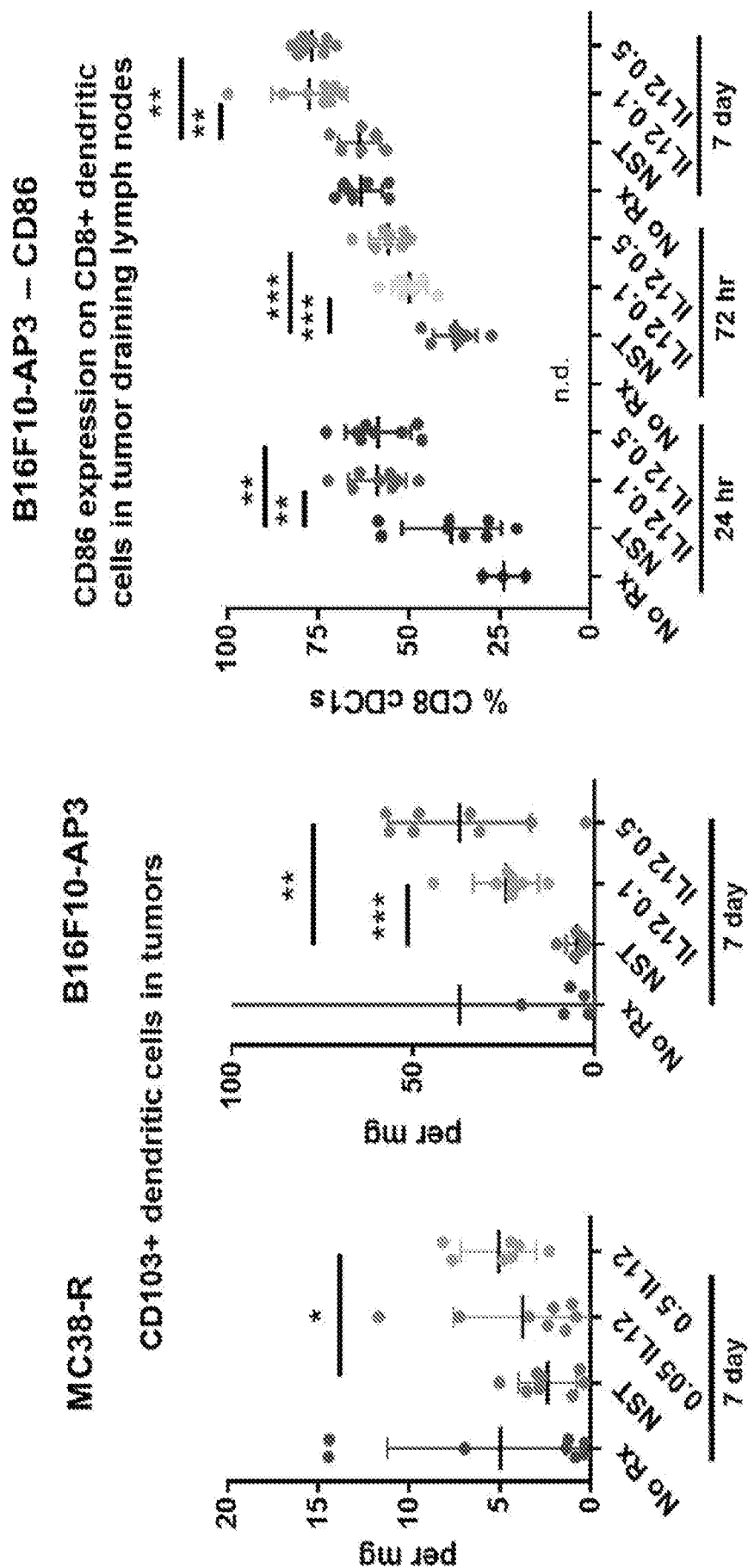


FIG. 33B

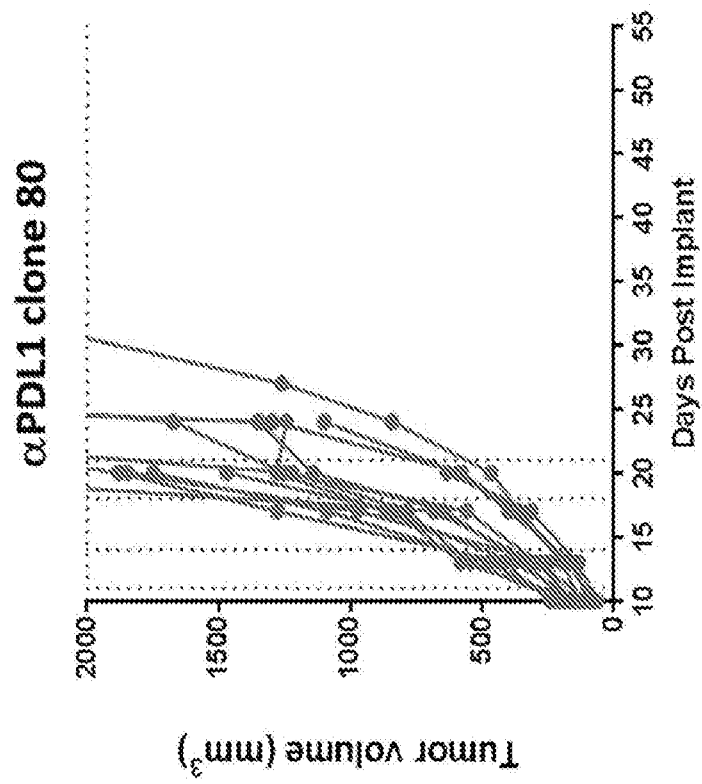


FIG. 33A

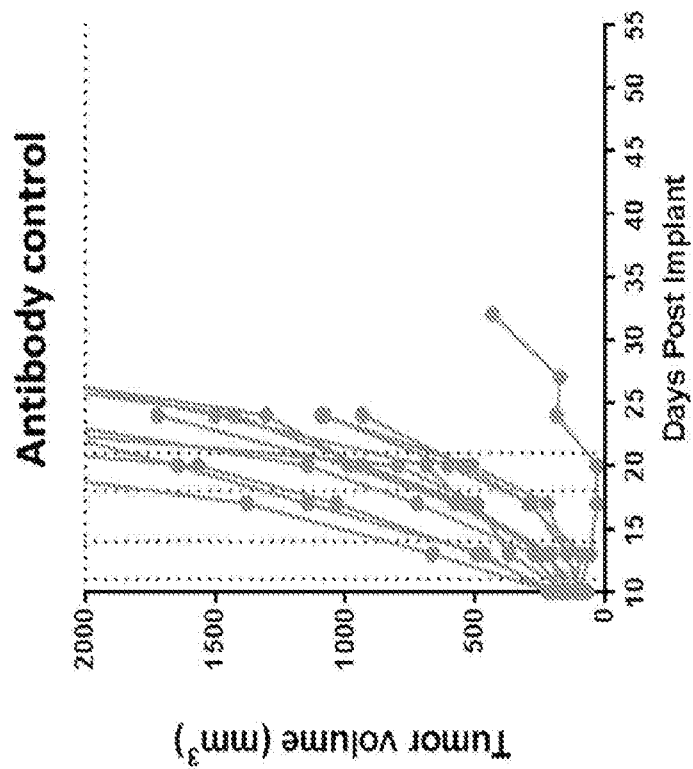


FIG. 33D

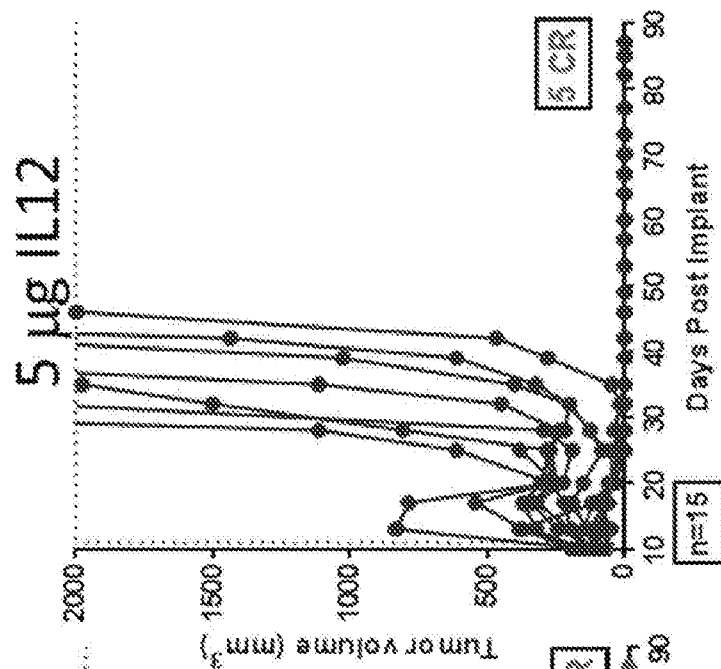


FIG. 33C

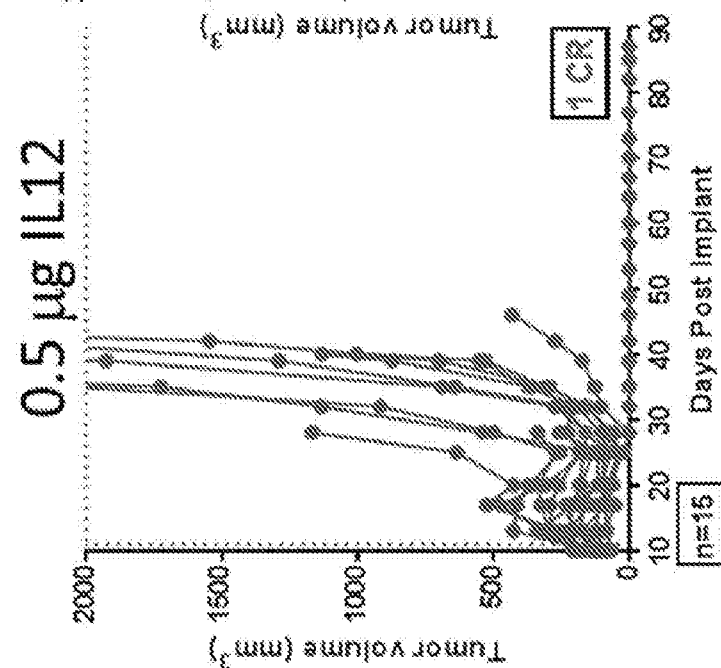


FIG. 33F

FIG. 33E

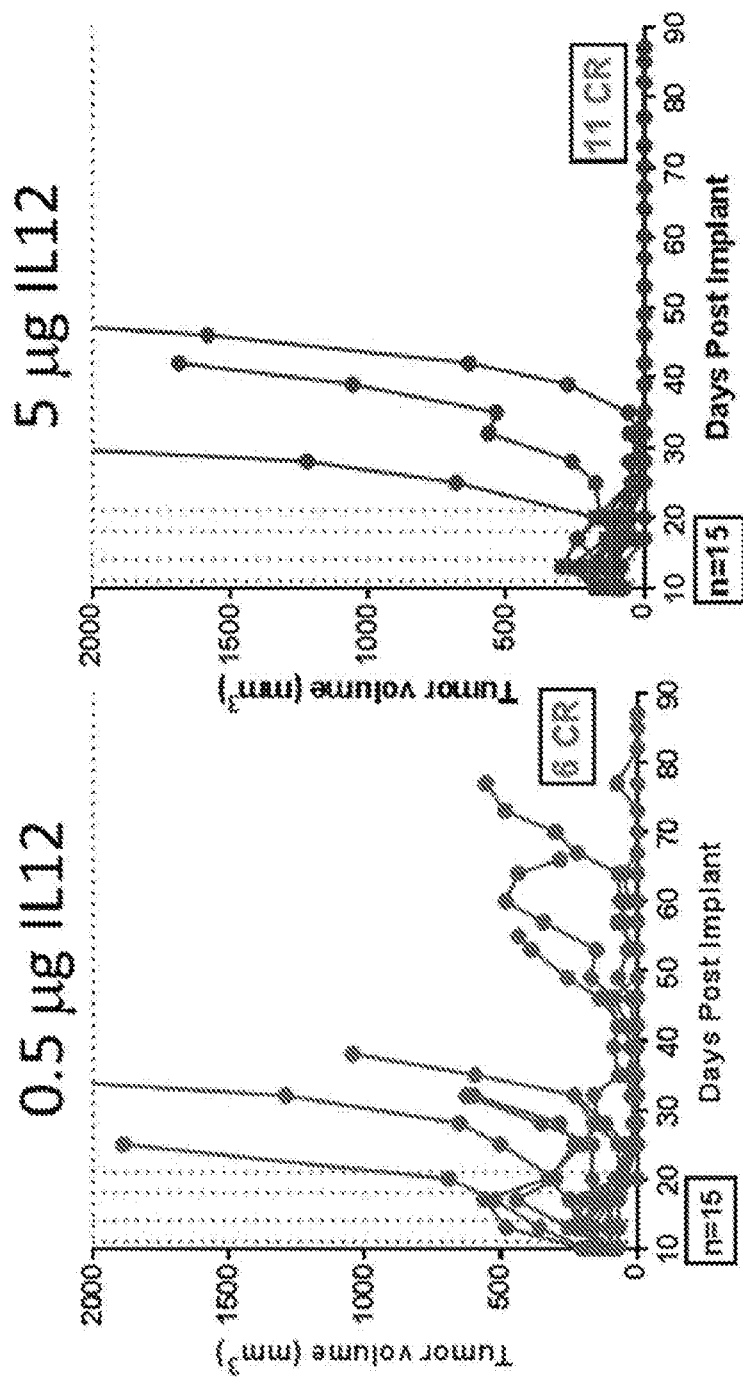


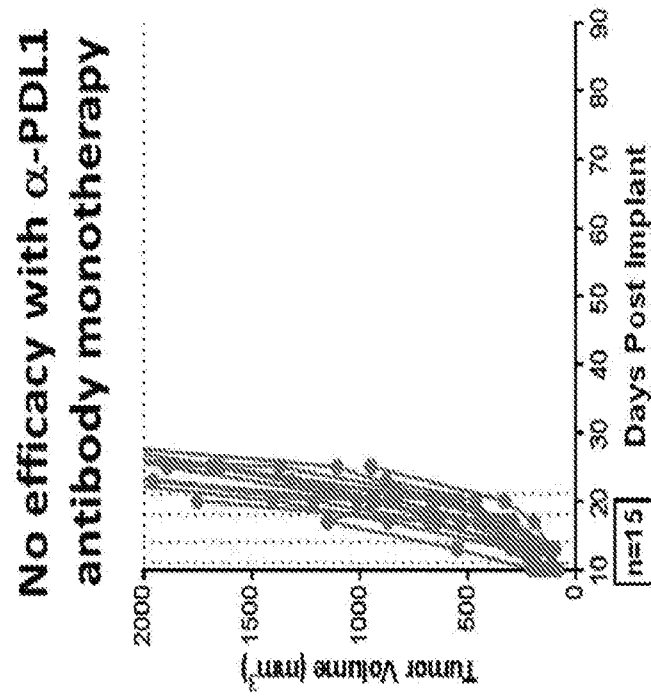
FIG. 33G

FIG. 34A

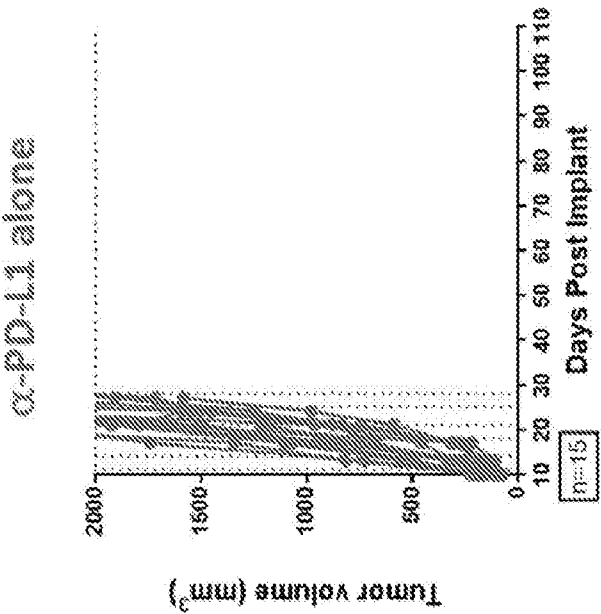


FIG. 34B

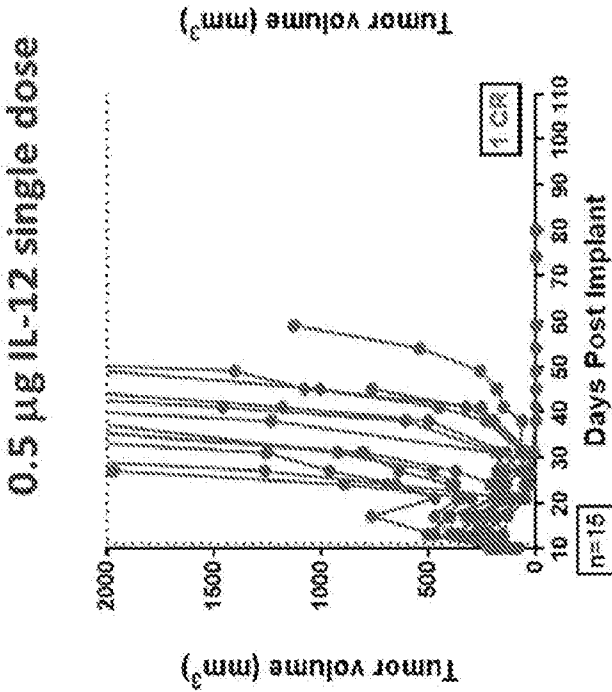


FIG. 34C

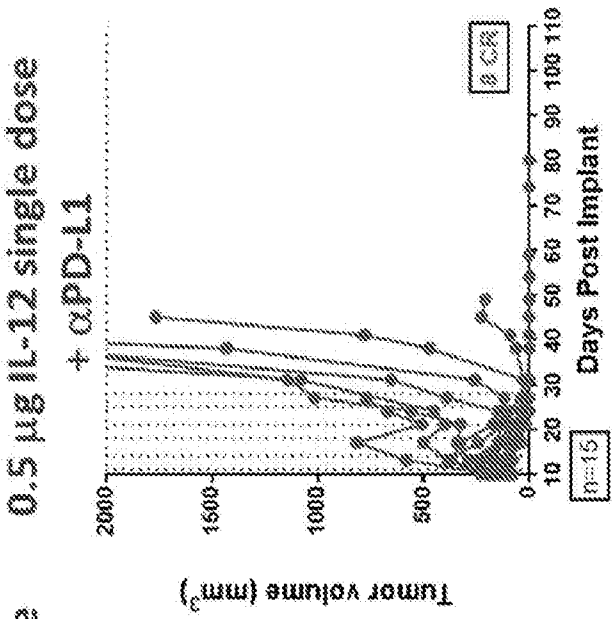


FIG. 35A

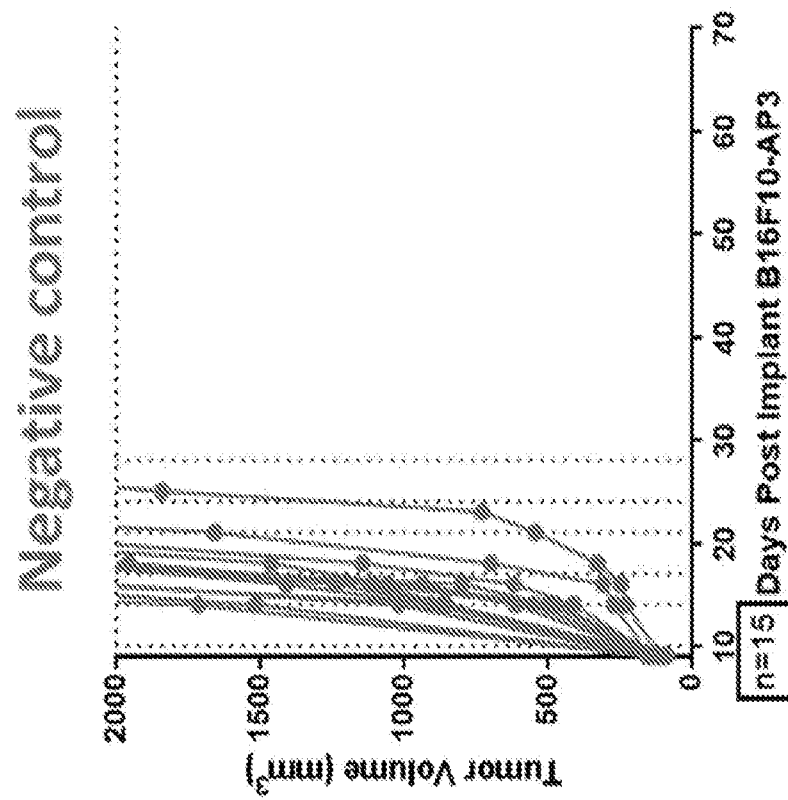


FIG. 35B

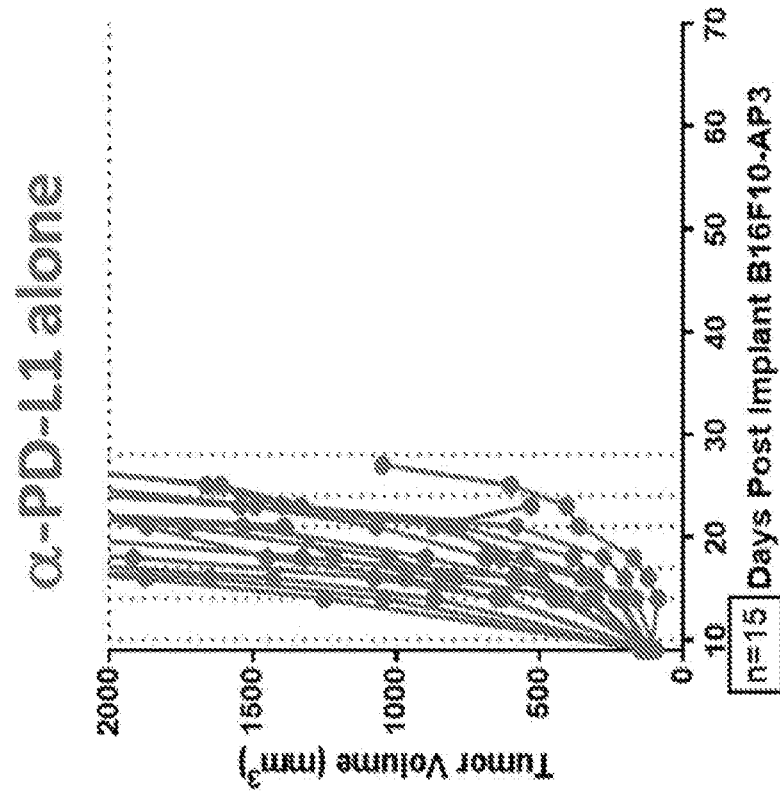


FIG. 35C

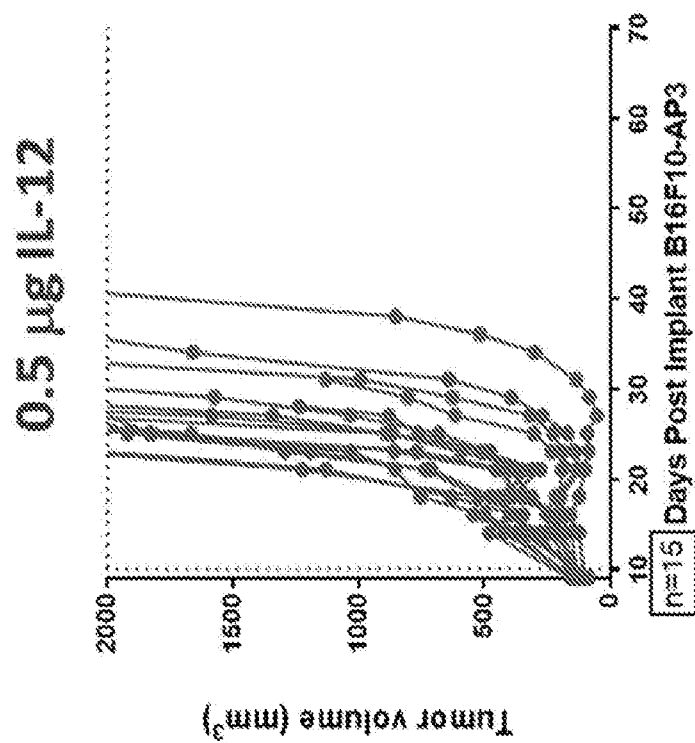


FIG. 35D

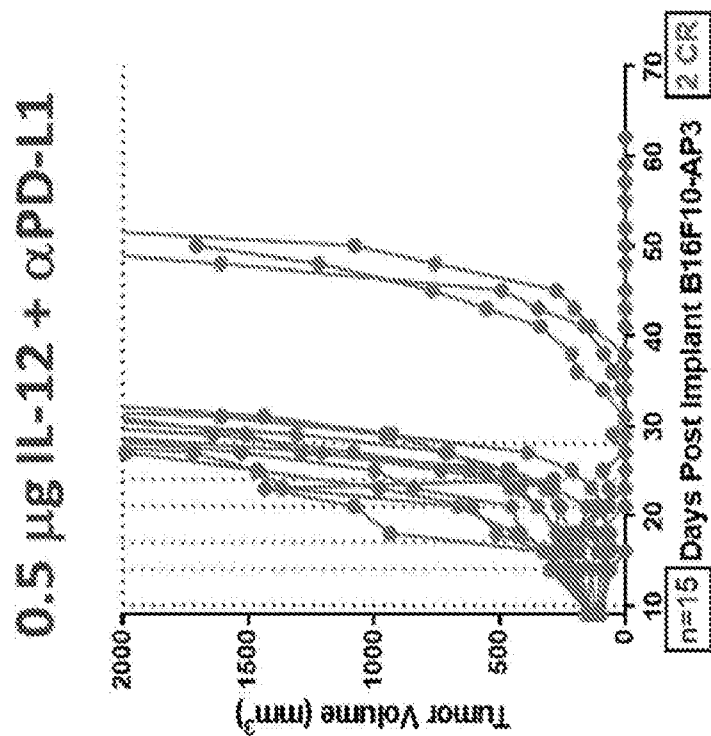


FIG. 36A

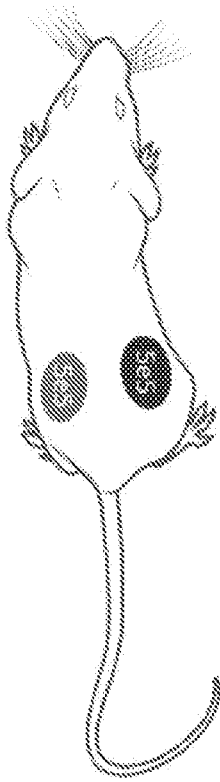


FIG. 36B

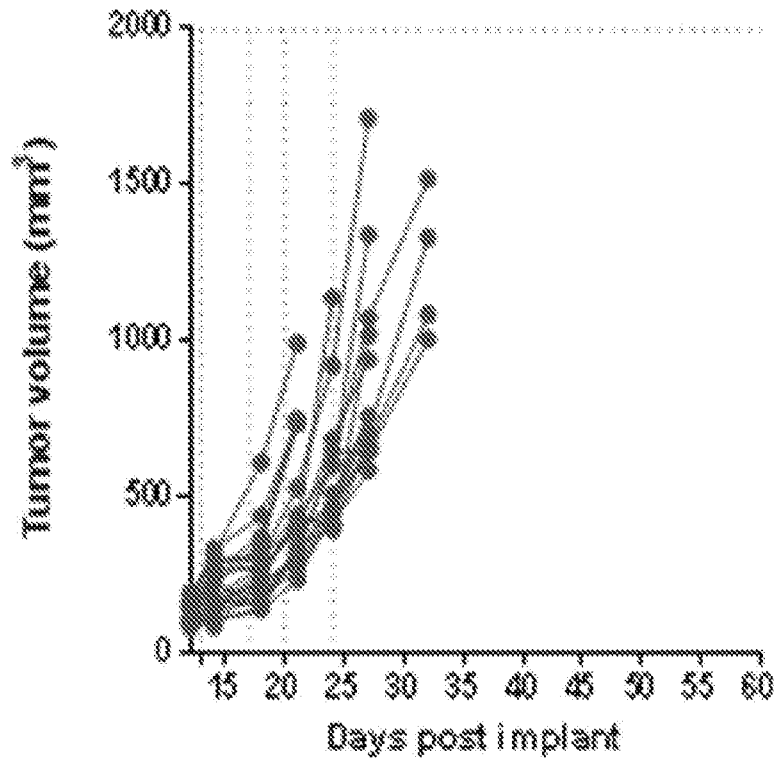


FIG. 36C

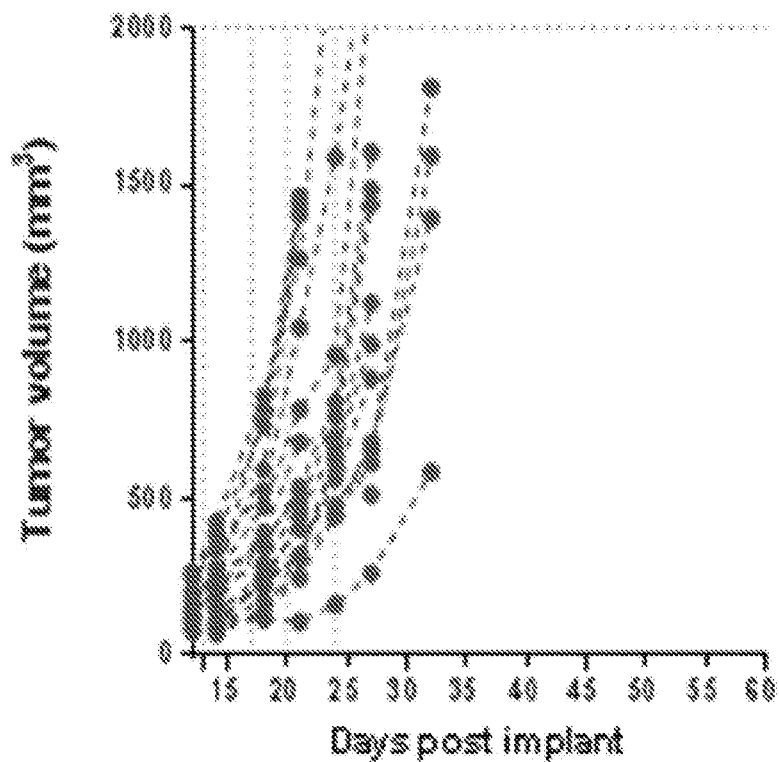


FIG. 36D

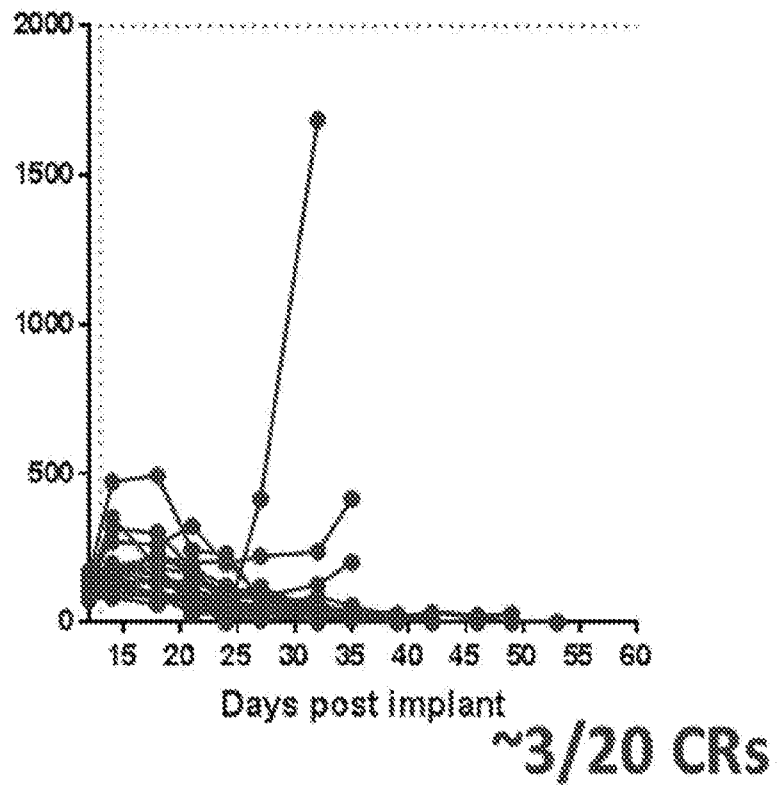


FIG. 36E

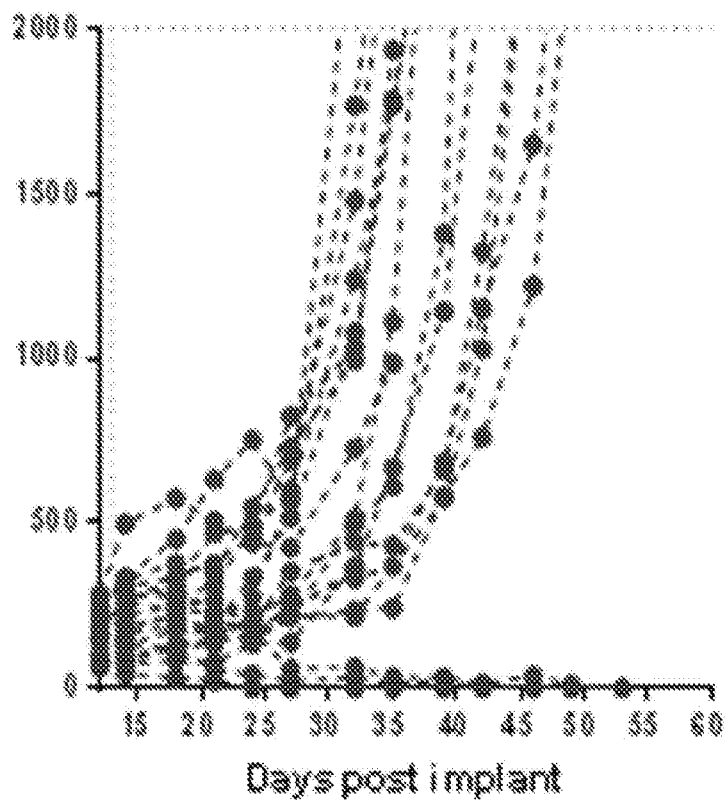
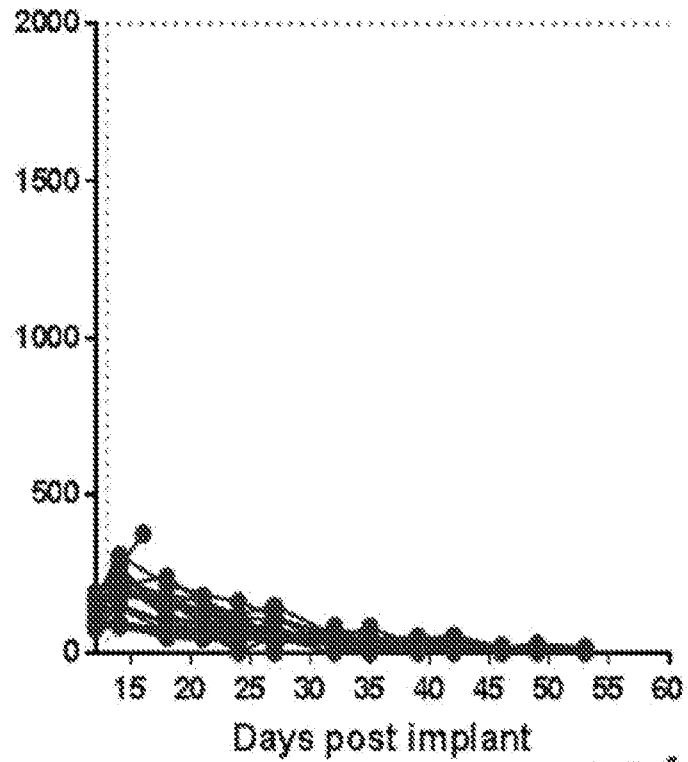


FIG. 36F



~16/20 CRs

FIG. 36G

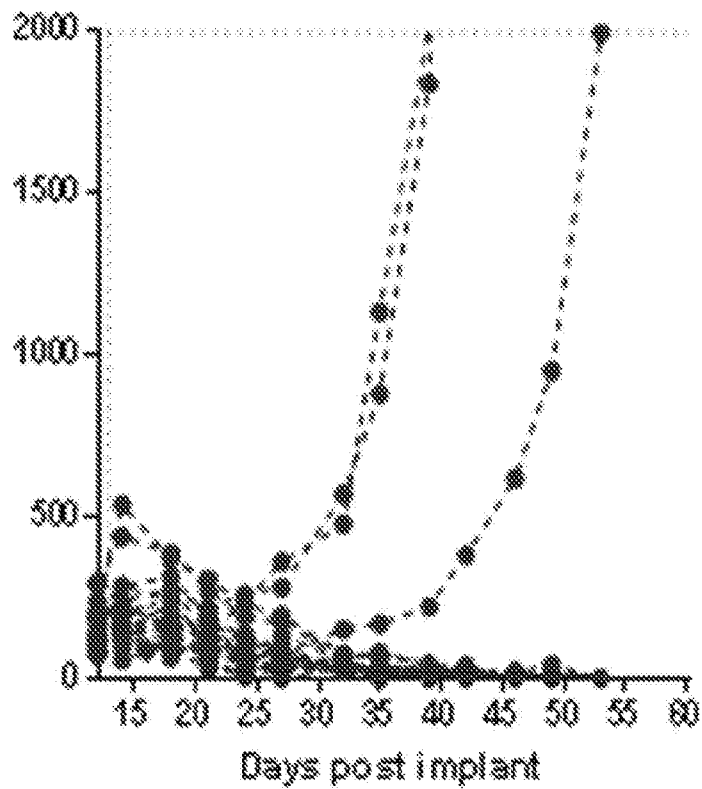


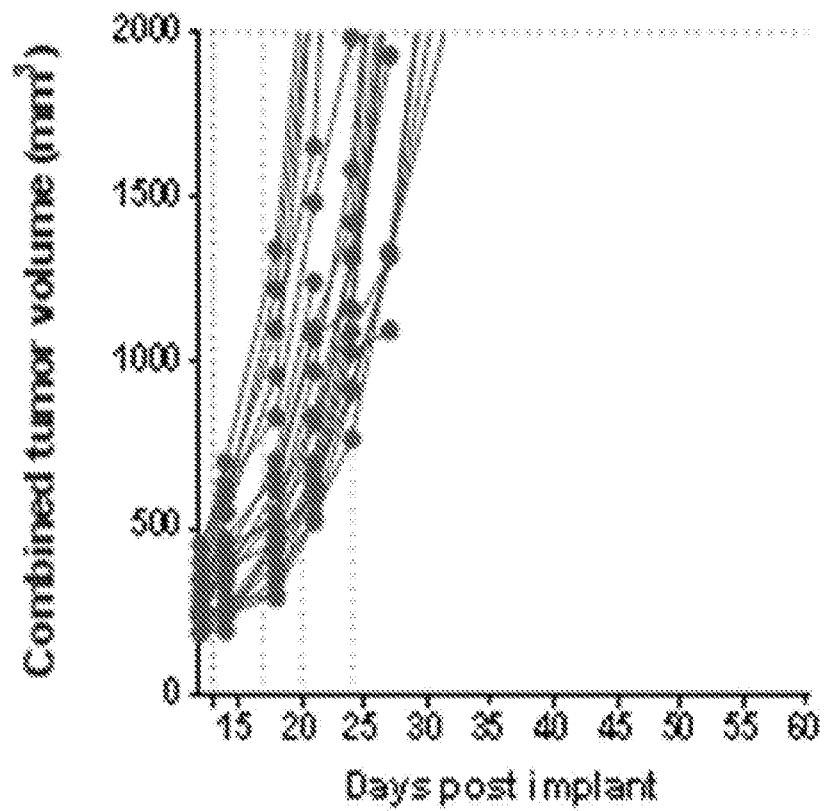
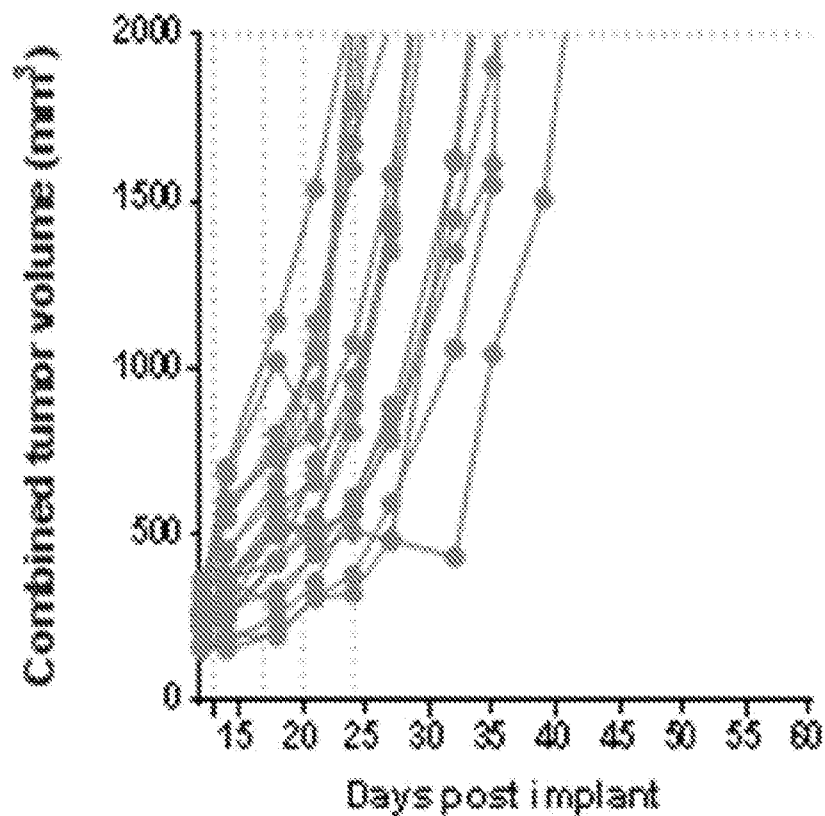
FIG. 37A**FIG. 37B**

FIG. 37C

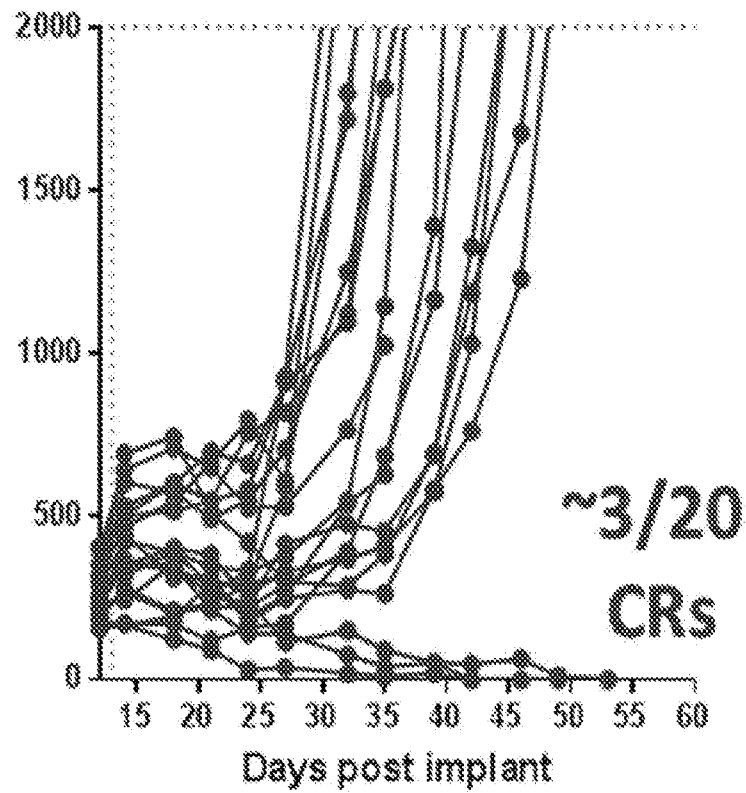


FIG. 37D

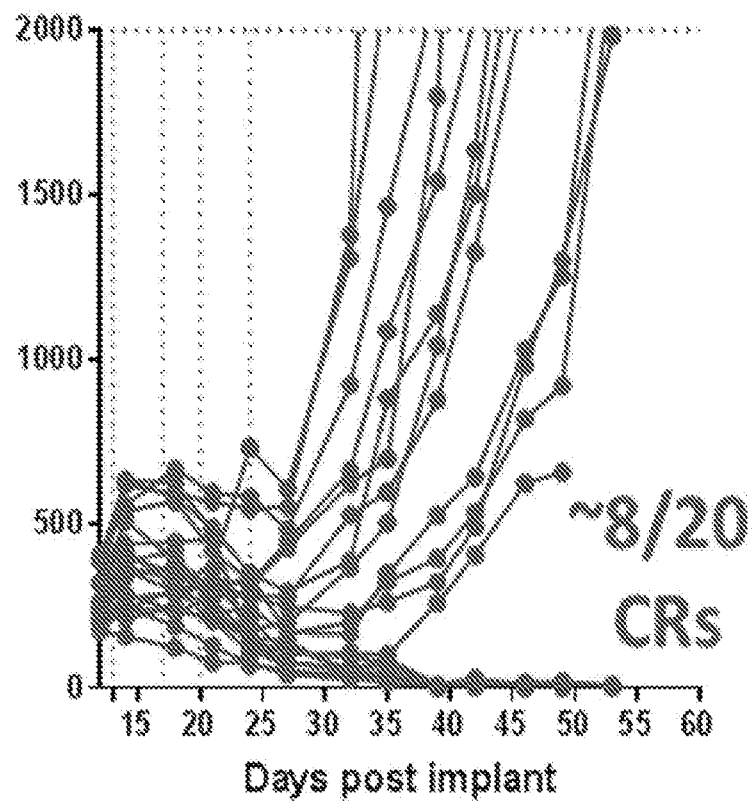


FIG. 37E

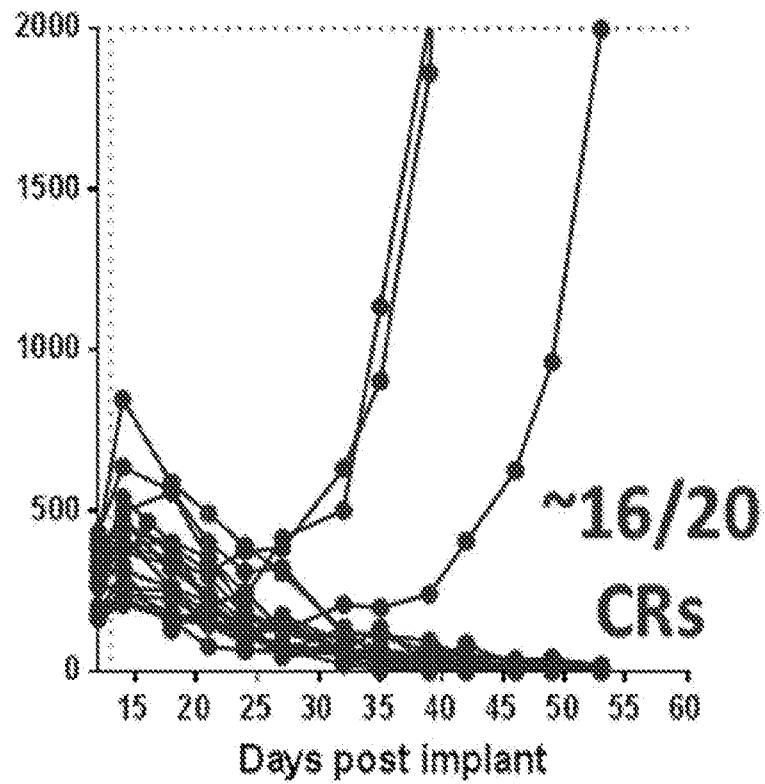


FIG. 37F

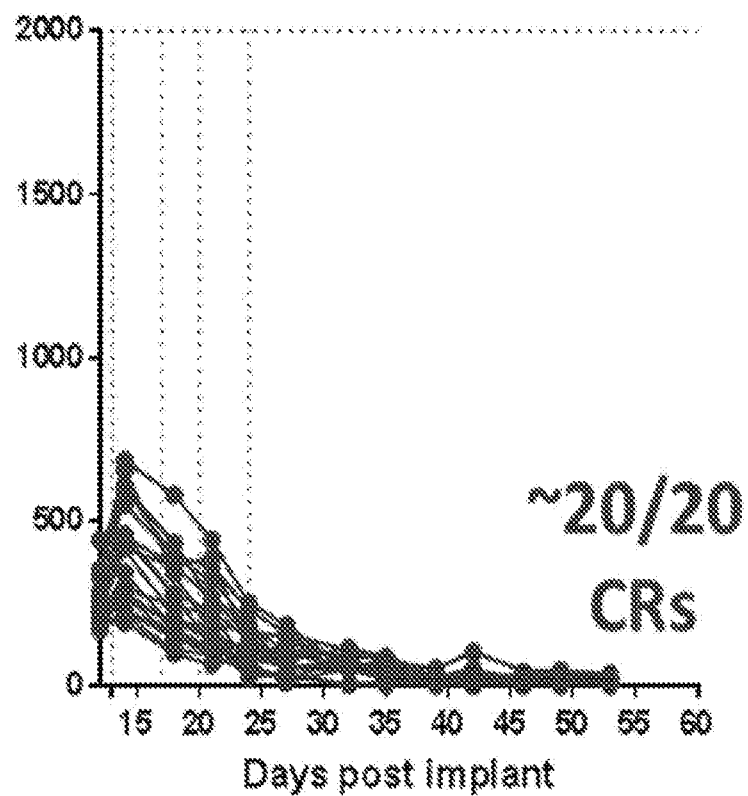


FIG. 38

In vitro and in vivo hIL-12 expression comparison

