

```

SP|P56856|CLD18_HUMAN  MSTTTCQVVAFLLSILGLAGCIAATGMDMWSTQDLYDNPVTSVPQYEGLRSCVRQSSGF 60
SP|P56856-2|CLD18_HUMAN MAVTACQGLGFVVSILIGIAGIIAATCMDQWSTQDLYNPNPTAVFNQGLWRSCVRESSGF 60
                        *:.*:** :.*:**:** ** ** *****:**:**:**:**:**:**
                        *

SP|P56856|CLD18_HUMAN  TECRPYFTILGLPAMLQAVRALMIVGIVLGAIGLLVSIFALKCIRIGSMEDSAKANMTLT 120
SP|P56856-2|CLD18_HUMAN TECRGYFTLLGLPAMLQAVRALMIVGIVLGAIGLLVSIFALKCIRIGSMEDSAKANMTLT 120
                        **** **:*:**:**:**:**:**:**:**:**:**:**
                        *

SP|P56856|CLD18_HUMAN  SGIMFIVSGLCAIAGVSVFANMLVINFWMSTANMYTGMGGMVQTVQTRYTFGAALFVGWV 180
SP|P56856-2|CLD18_HUMAN SGIMFIVSGLCAIAGVSVFANMLVINFWMSTANMYTGMGGMVQTVQTRYTFGAALFVGWV 180
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SP|P56856|CLD18_HUMAN  AGGLTLIGGVMMCIACRGLAPEETNYKAVSYHASGHSVAYKPGGFKASTGFGSNTKNKKI 240
SP|P56856-2|CLD18_HUMAN AGGLTLIGGVMMCIACRGLAPEETNYKAVSYHASGHSVAYKPGGFKASTGFGSNTKNKKI 240
                        *****:**:**:**:**:**:**:**:**:**:**
                        *

SP|P56856|CLD18_HUMAN  YDGGARTEDEVQSYPSKHDYV 261
SP|P56856-2|CLD18_HUMAN YDGGARTEDEVQSYPSKHDYV 261
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```

Fig. 1A

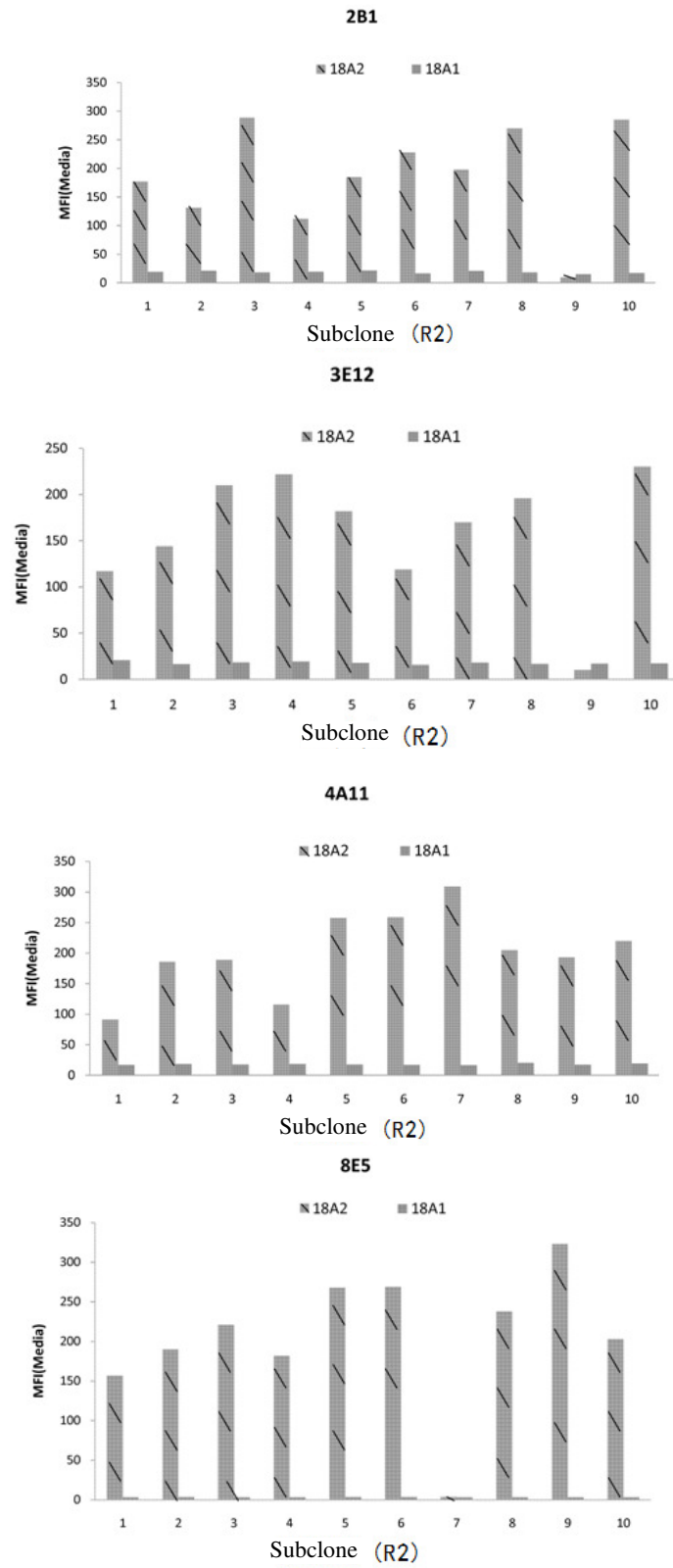


Fig. 1B

	HCDR1										HCDR2																																																											
2B1(scFv_Fc)	Q	V	L	Q	L	G	S	G	A	E	L	A	R	P	G	A	S	V	K	M	S	C	K	A	S	G	Y	T	T	S	-	Y	T	M	H	V	W	V	K	Q	R	P	G	Q	G	L	E	W	I	G	Y	I	N	P	S	S	G	Y	I	N	Q	K	F	D	K	A	T			
3E12(scFv_Fc)	Q	V	L	Q	L	G	S	G	A	E	L	A	R	P	G	A	S	V	K	M	S	C	K	A	S	G	Y	T	T	S	-	Y	D	I	N	H	V	W	V	K	Q	R	P	G	Q	G	L	E	W	I	G	W	I	N	P	G	D	S	K	T	K	N	E	K	F	K	G	A	T	
4A1(scFv_Fc)	Q	Q	I	L	V	Q	S	G	P	E	L	K	K	P	E	L	V	K	I	S	C	K	A	S	G	Y	T	T	N	-	G	M	N	W	V	K	Q	A	P	G	K	G	L	K	W	M	G	Y	I	N	T	I	N	T	E	P	T	Y	A	E	E	F	K	G	R	F	A			
0E5(scFv_Fc)	D	V	Q	L	Q	E	S	G	P	D	L	V	K	P	S	Q	S	L	S	L	T	C	T	V	T	G	Y	S	I	T	-	S	G	Y	N	W	H	W	I	R	Q	F	P	G	N	K	M	E	W	M	G	Y	I	H	-	T	G	S	T	N	Y	N	P	S	L	R	S	R	I	S

	HCDR3																																																	
2B1(scFv_Fc)	L	T	A	D	K	S	S	S	T	A	Y	M	Q	L	S	S	L	T	S	E	D	S	A	V	V	Y	C	A	R	-	-	I	Y	G	N	S	F	A	Y	W	G	Q	G	T	T	V	T	V	S	
3E12(scFv_Fc)	L	T	A	D	K	S	S	S	T	A	Y	M	Q	L	S	S	L	T	S	E	D	S	A	V	V	Y	C	A	R	-	-	G	G	Y	R	D	E	A	F	T	W	G	Q	G	T	T	V	T	V	S
4A1(scFv_Fc)	F	S	L	S	E	T	S	A	T	A	Y	L	I	N	N	L	K	N	E	D	T	A	T	Y	F	C	A	R	-	-	F	S	Y	G	N	S	F	A	Y	W	G	Q	G	T	T	V	T	V	S	
0E5(scFv_Fc)	L	I	R	D	S	K	N	Q	F	L	Q	L	N	S	V	T	I	D	D	T	A	T	Y	Y	C	T	R	-	-	I	Y	G	N	S	F	A	Y	W	G	Q	G	T	S	V	T	V	S			

[illegible]

Fig. 2

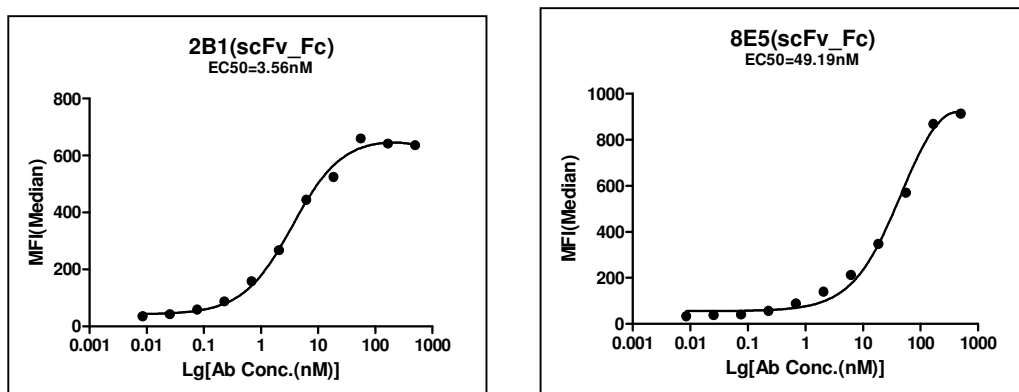


Fig. 3

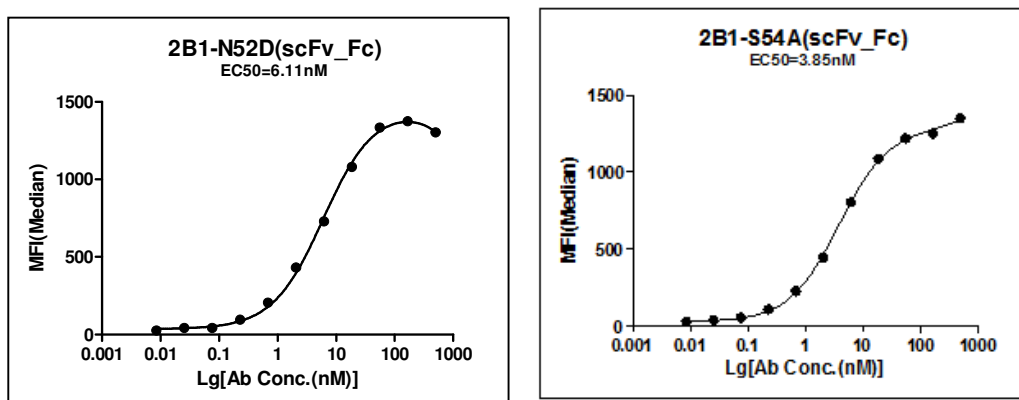


Fig. 4

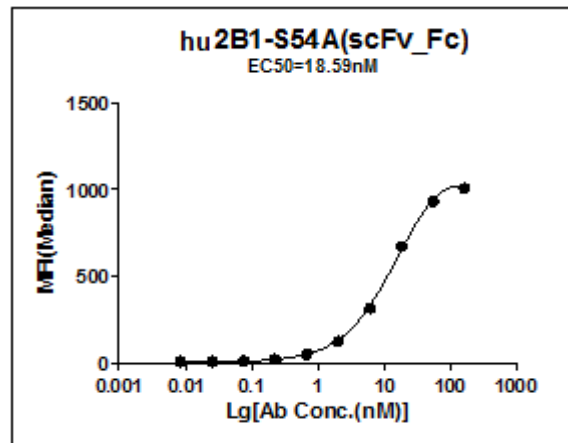


Fig. 5

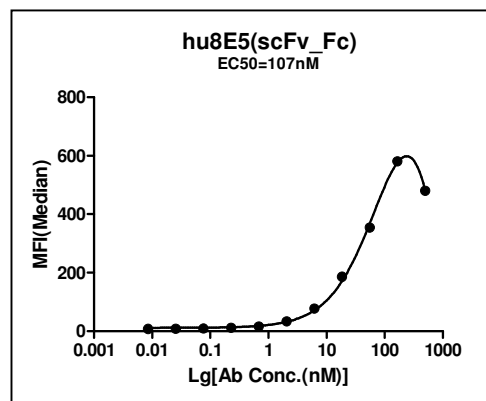


Fig. 6

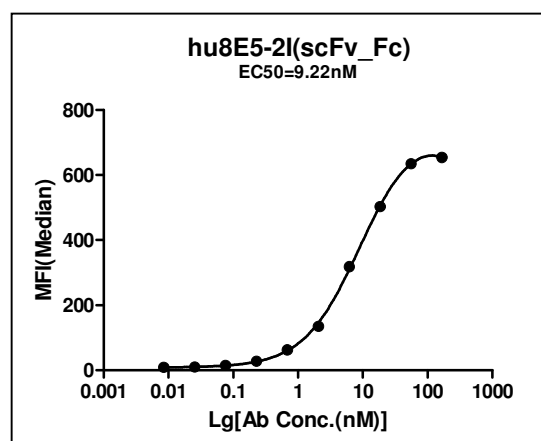


Fig. 7

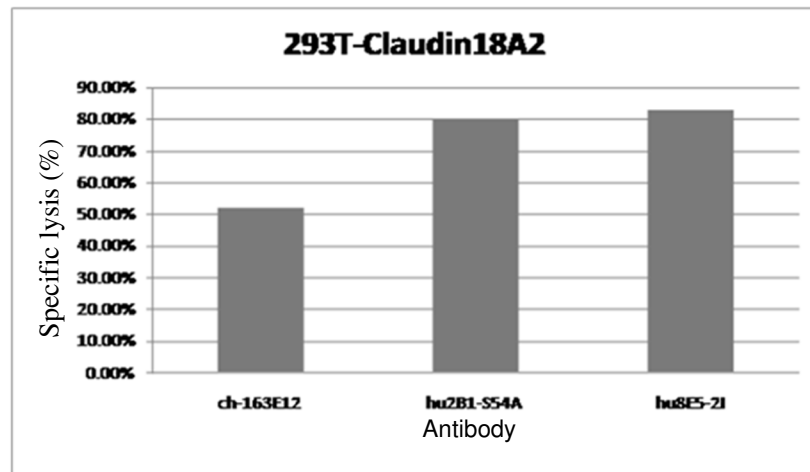


Fig. 8A

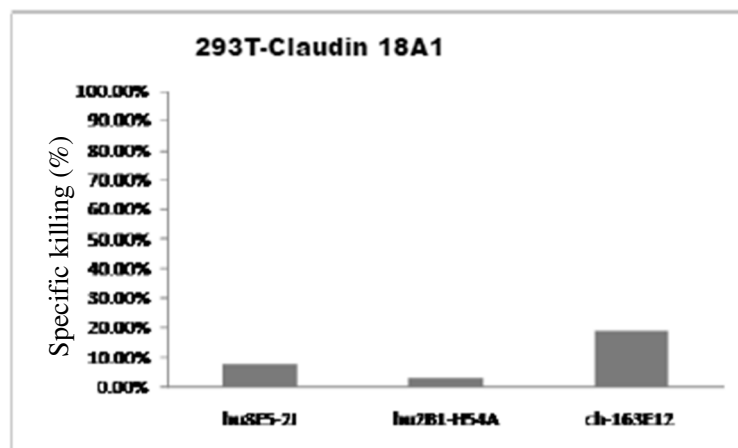


Fig. 8B

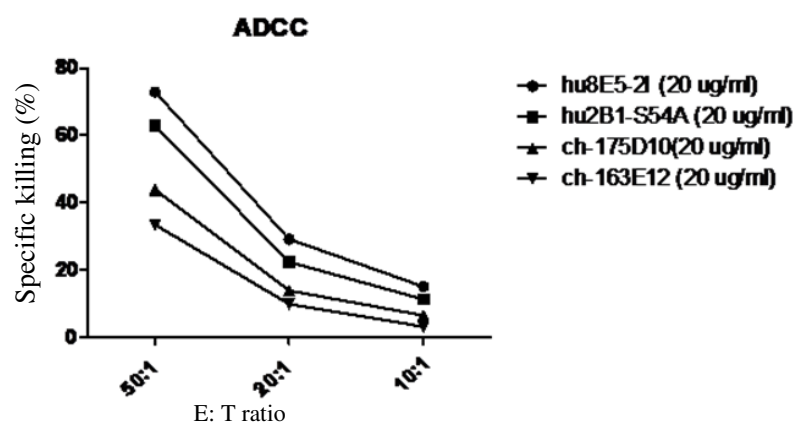


Fig. 9

Gastric cancer **PDX-GA0006**

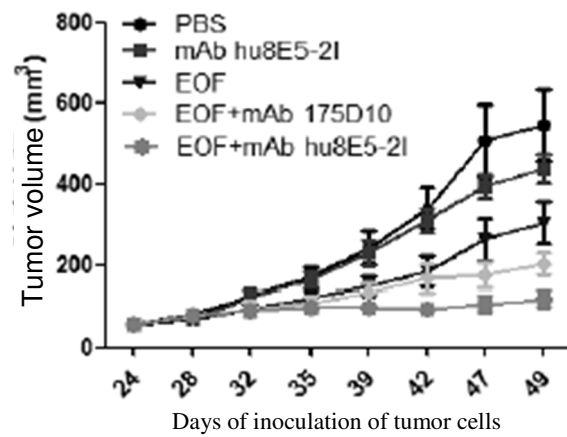


Fig. 10

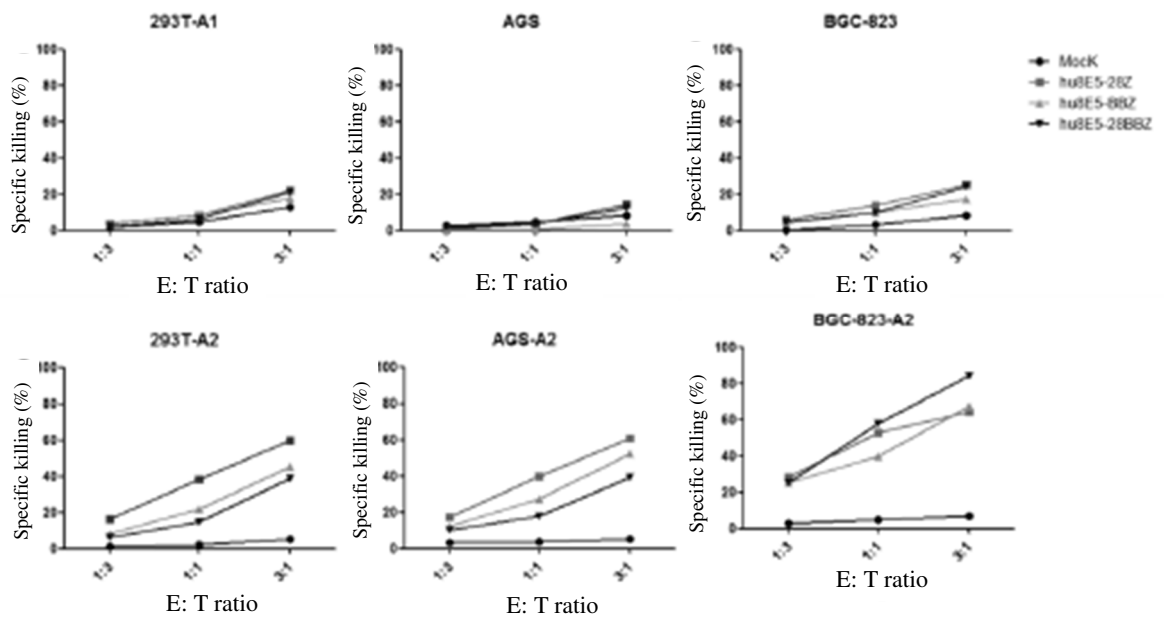


Fig. 11

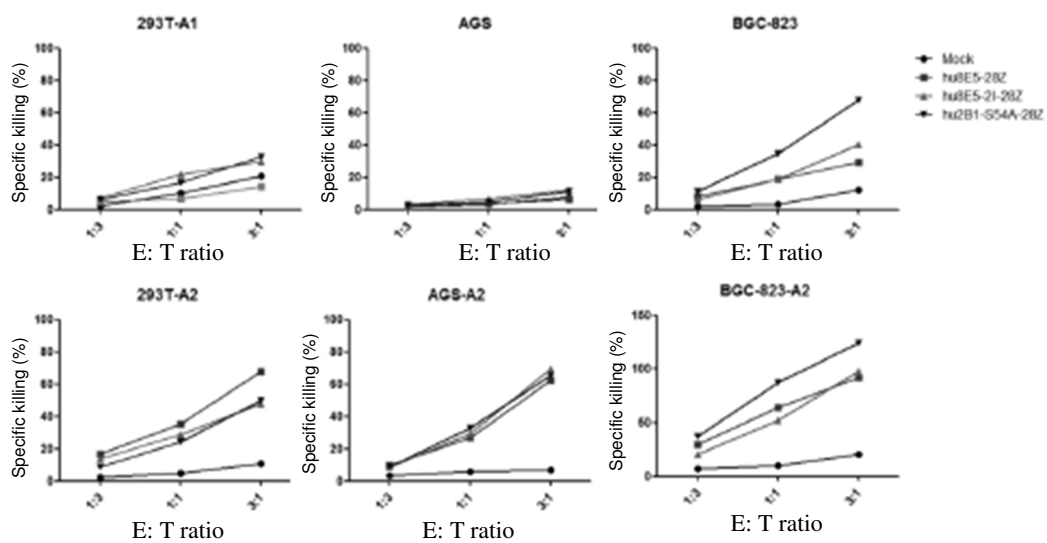


Fig. 12

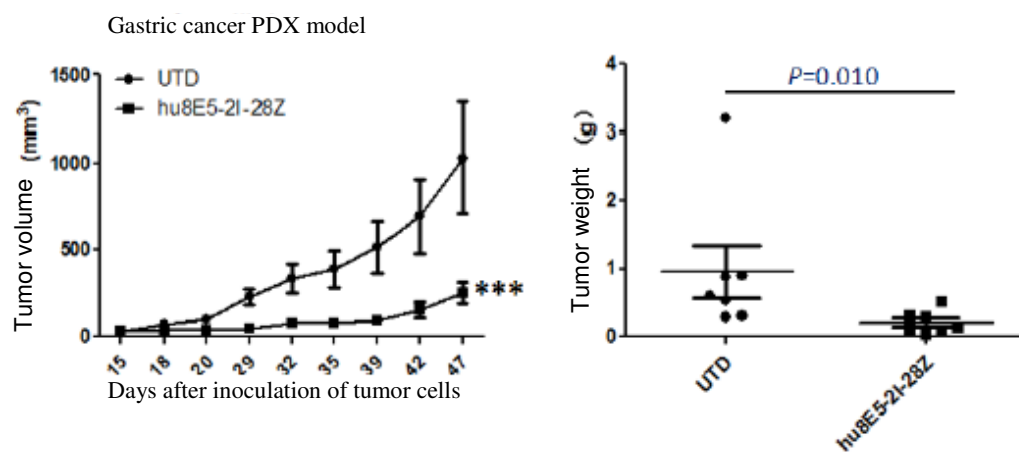


Fig. 13A

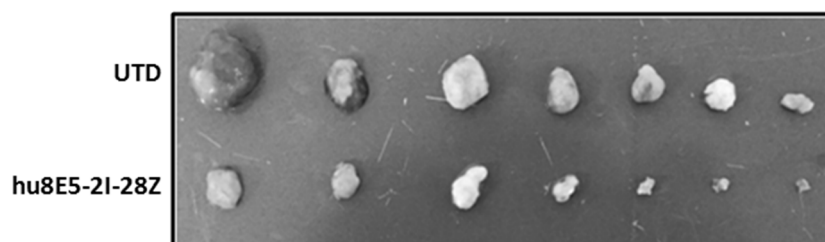


Fig. 13B

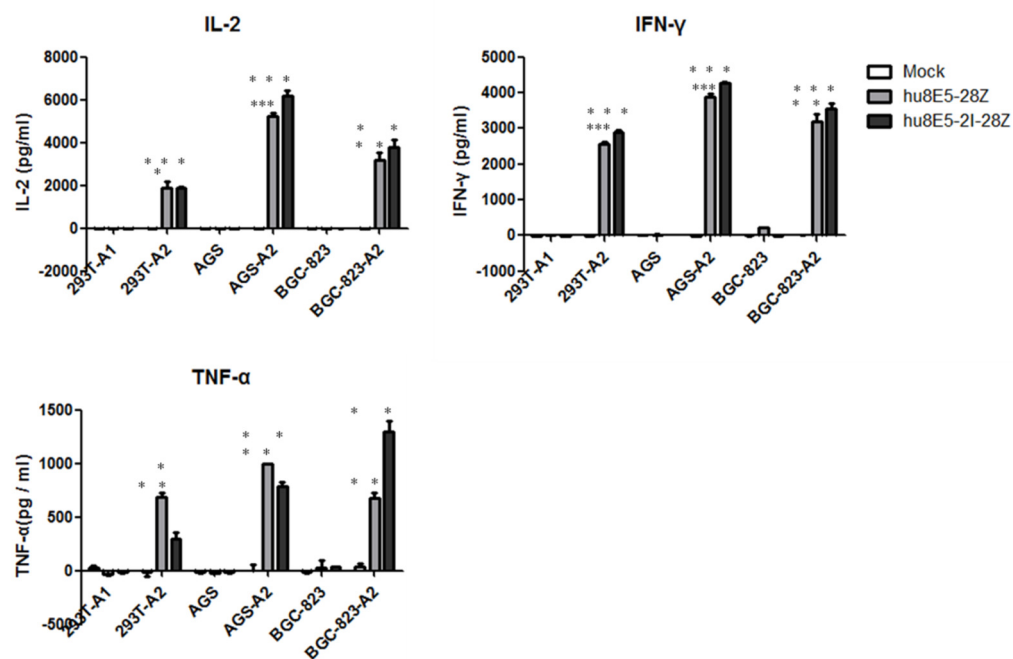


Fig. 14

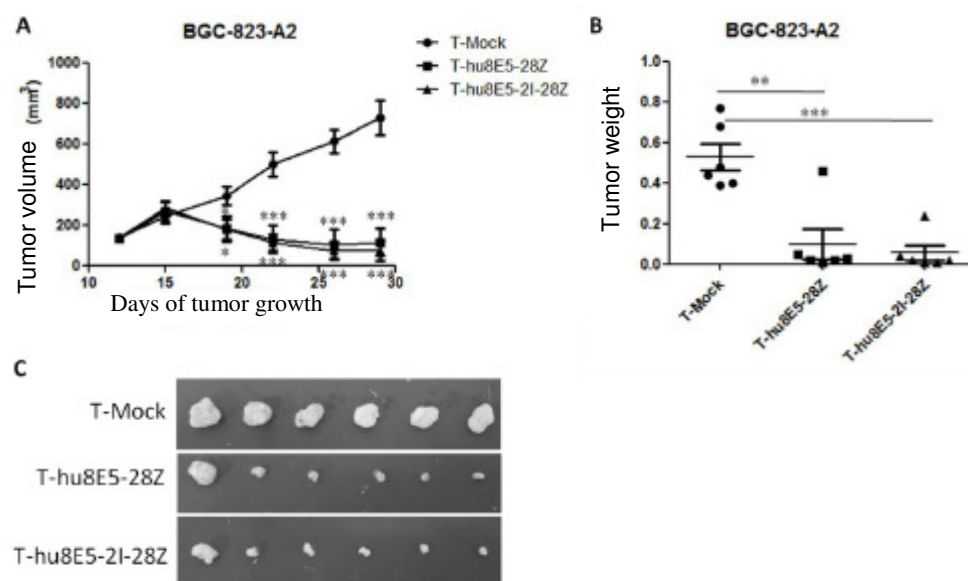


Fig. 15

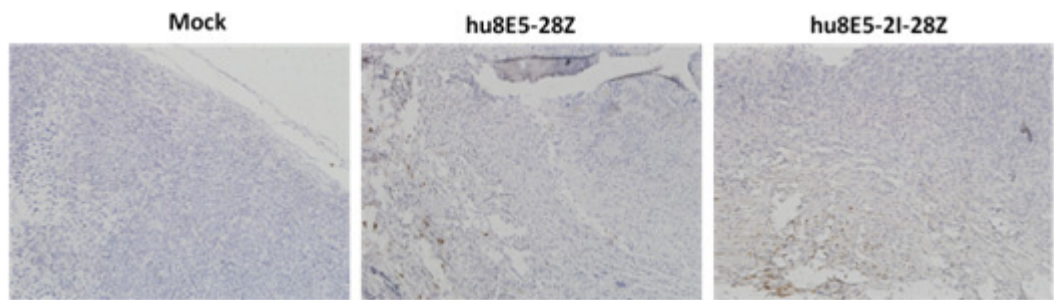


Fig. 16

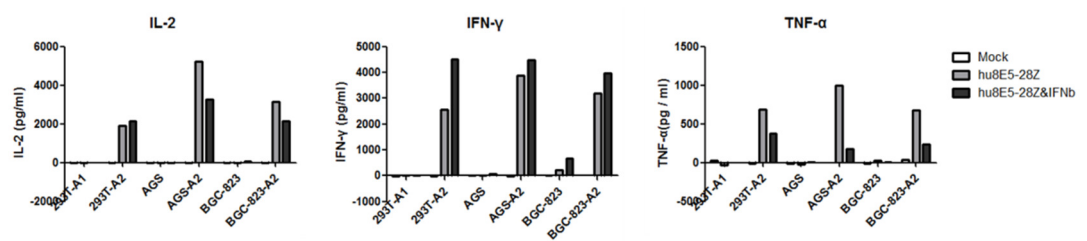


Fig. 17A

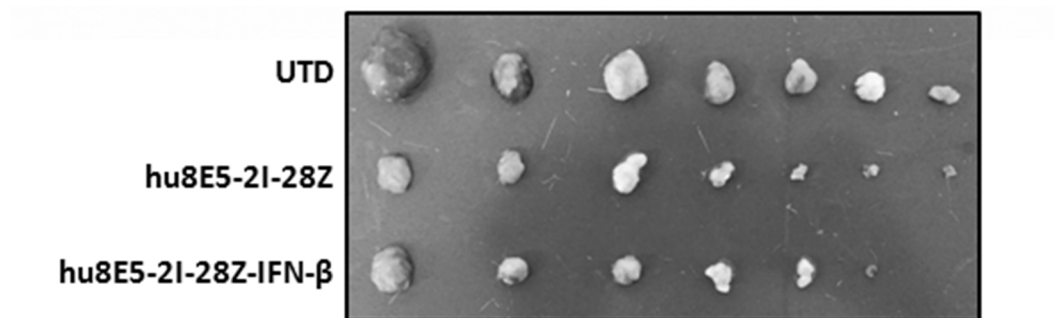


Fig. 17B

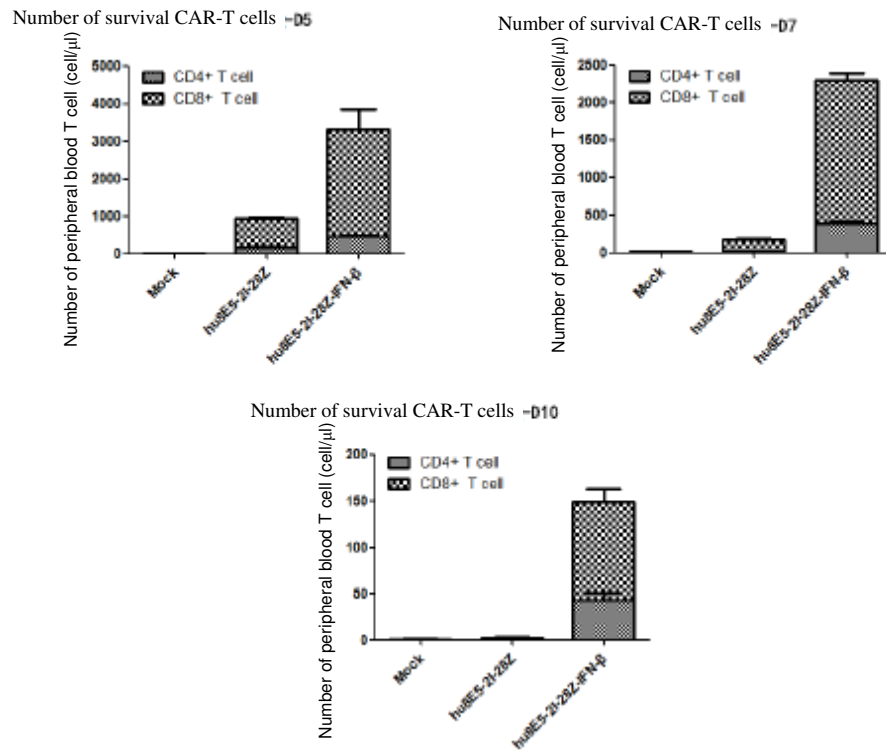


Fig. 17C

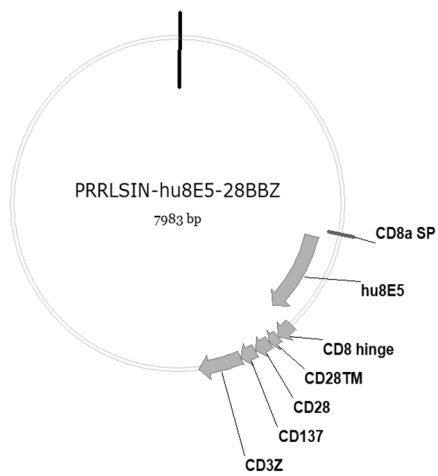


Fig. 18A

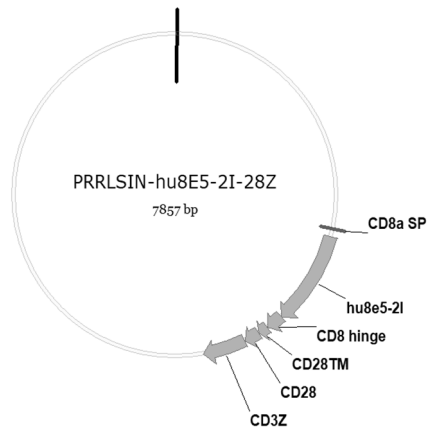


Fig. 18B

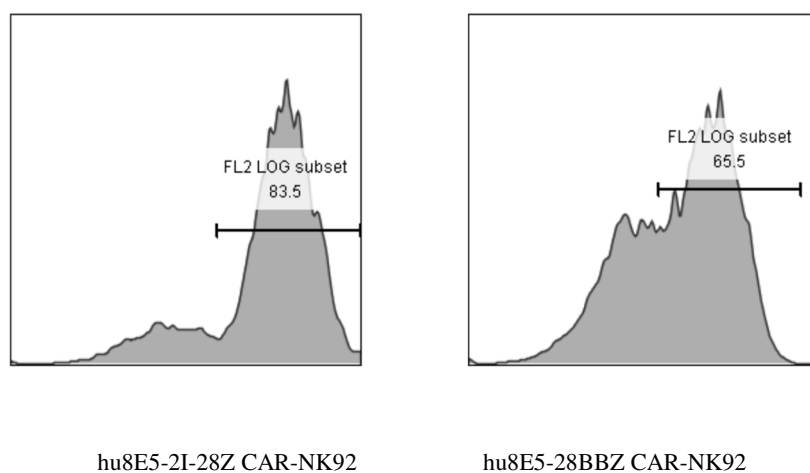


Fig. 19

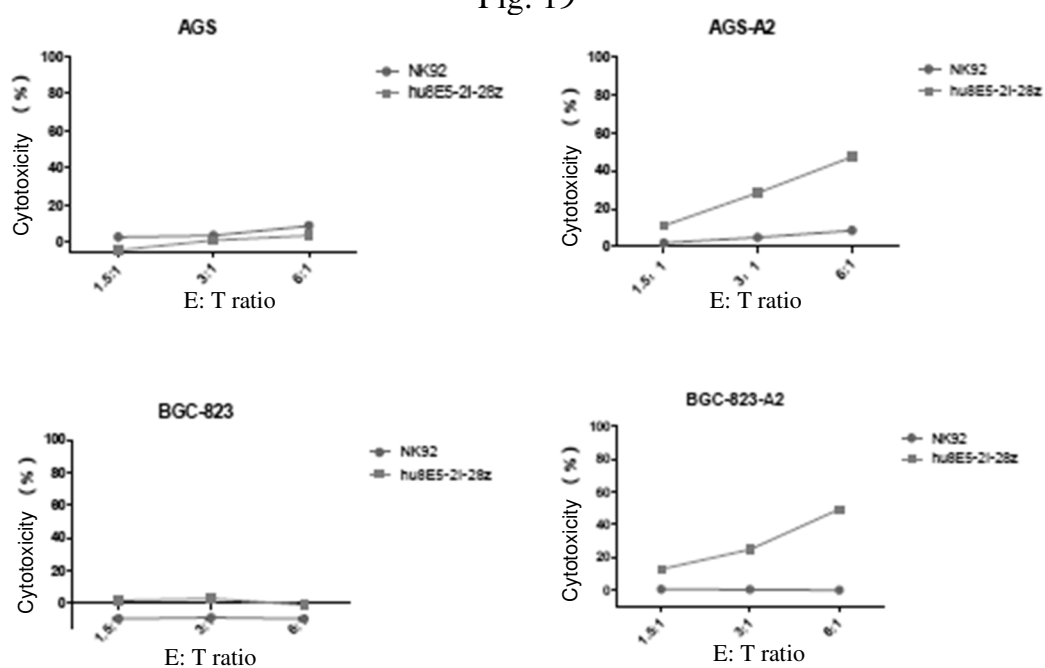


Fig. 20

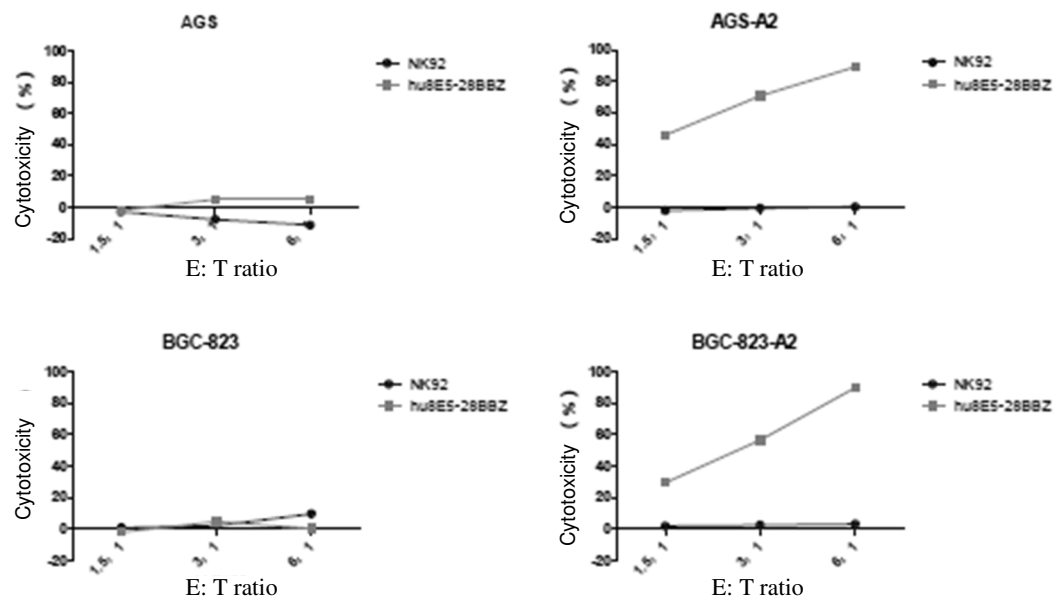


Fig. 21