

Statistics						
Name	Events	% Parent	% Grandparent	% Total	PE-A Mean	PE-A Median
1:P2	8,755	99.97	87.55	87.55	17,195	14,400

Figure S1

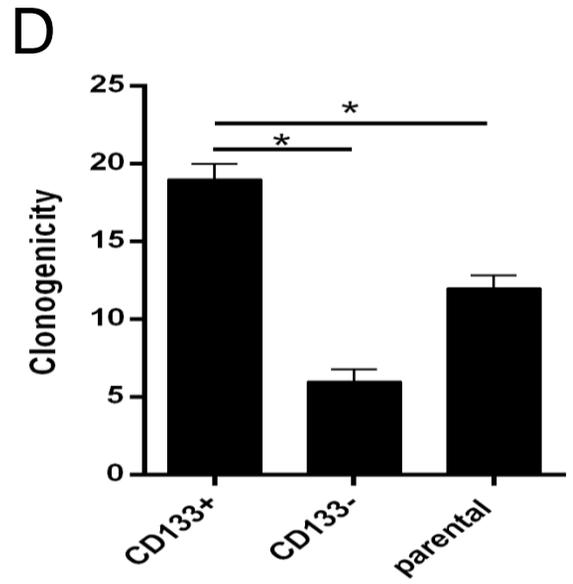
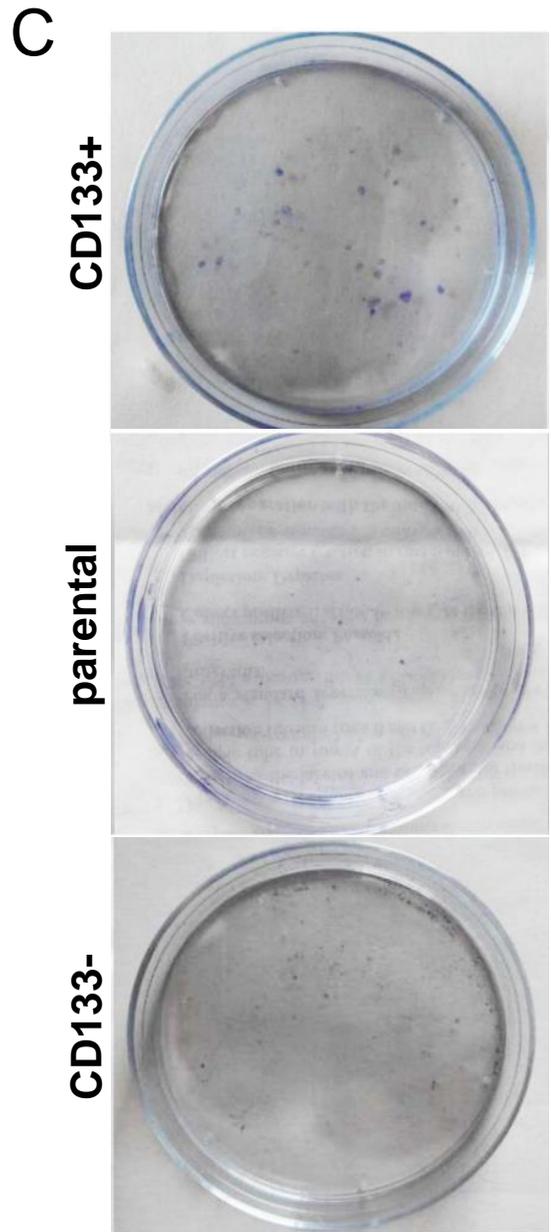
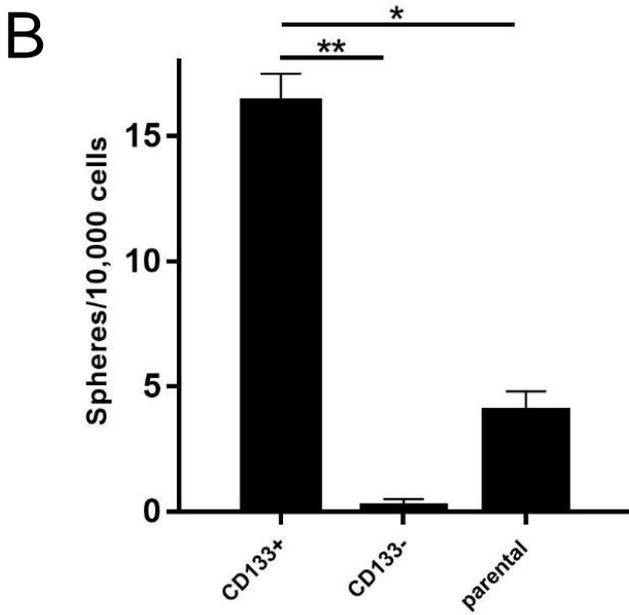
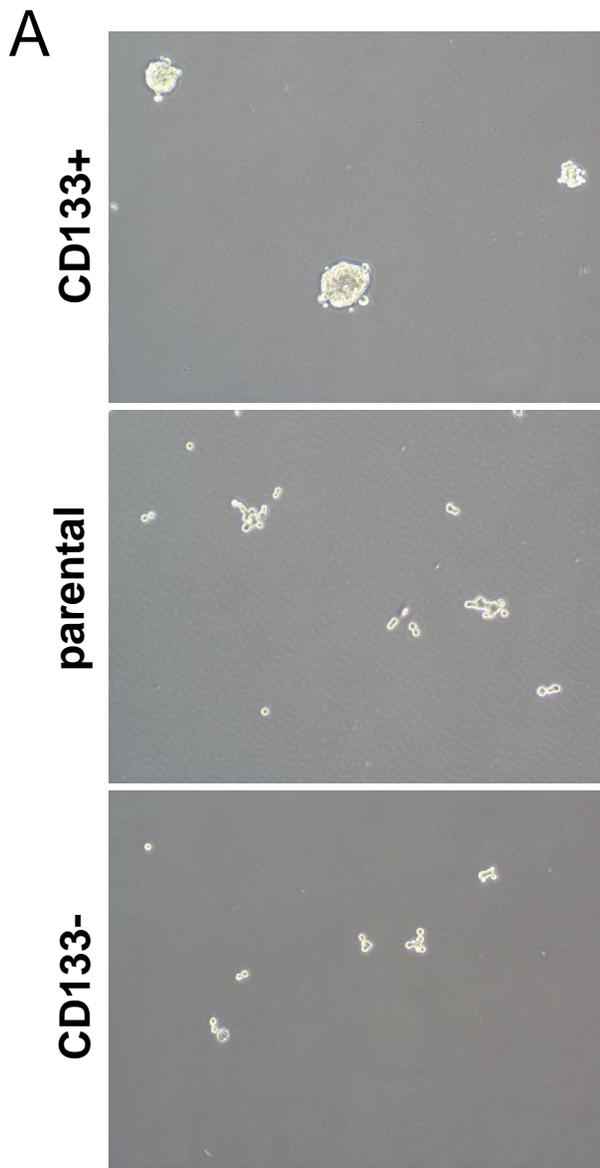
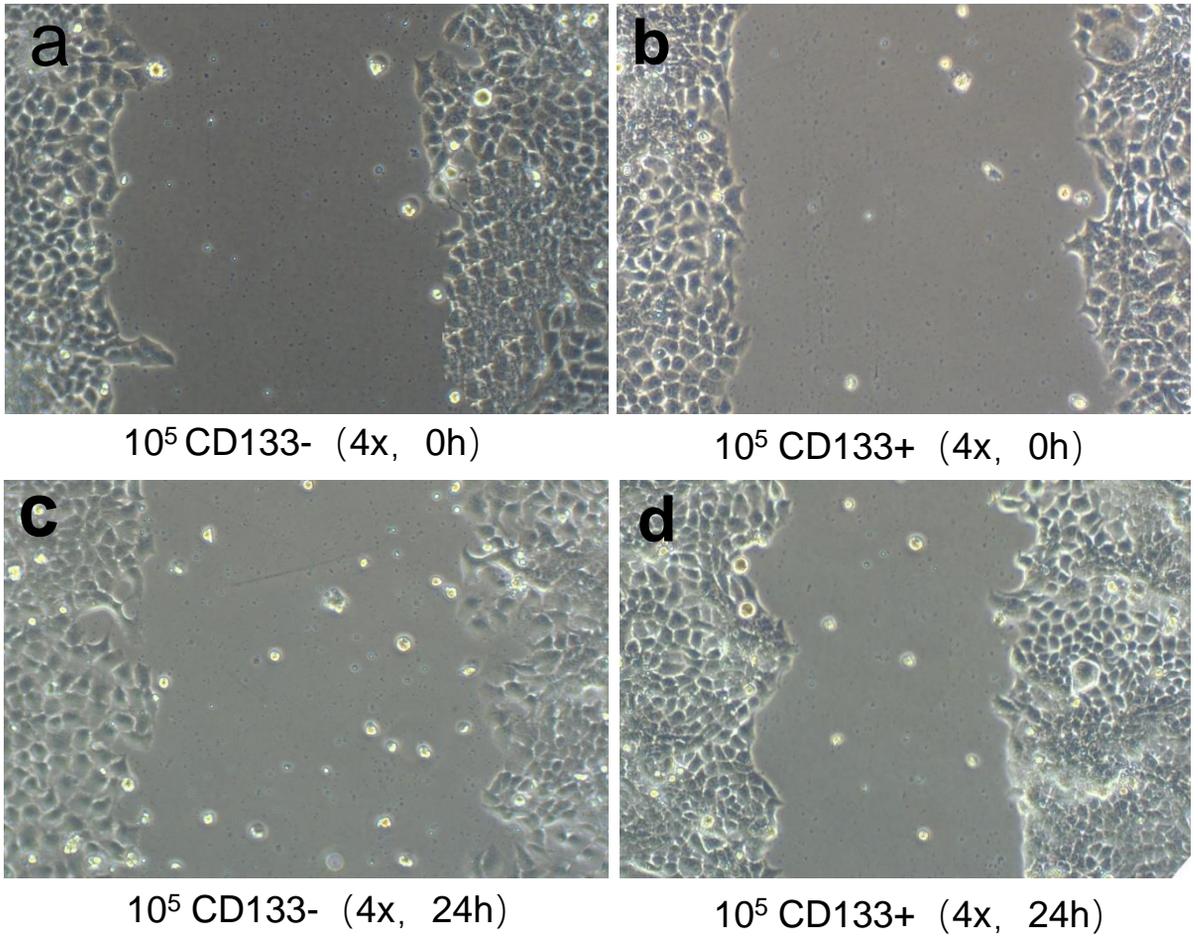
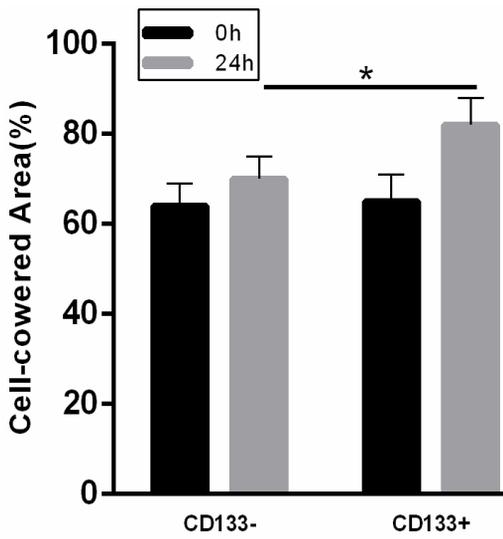
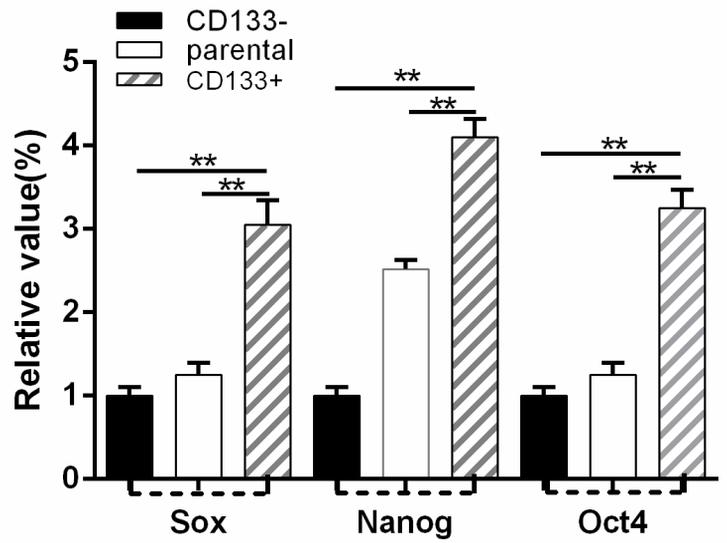


Figure S2

A**B****C**

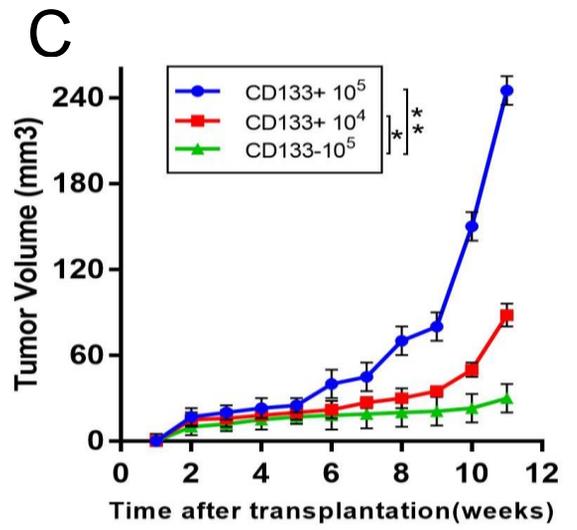
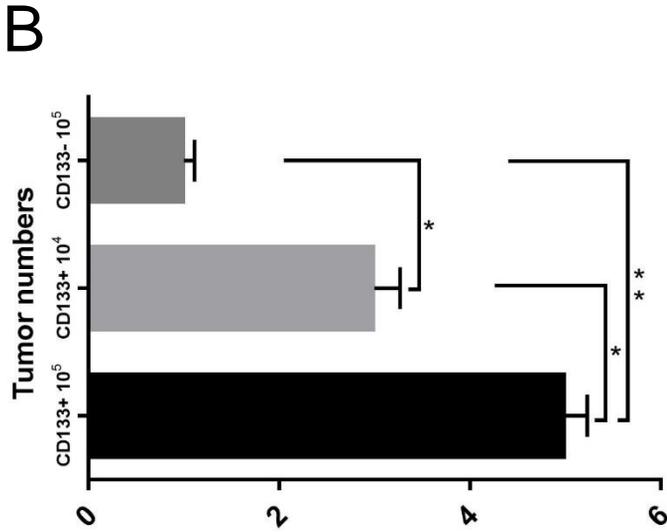
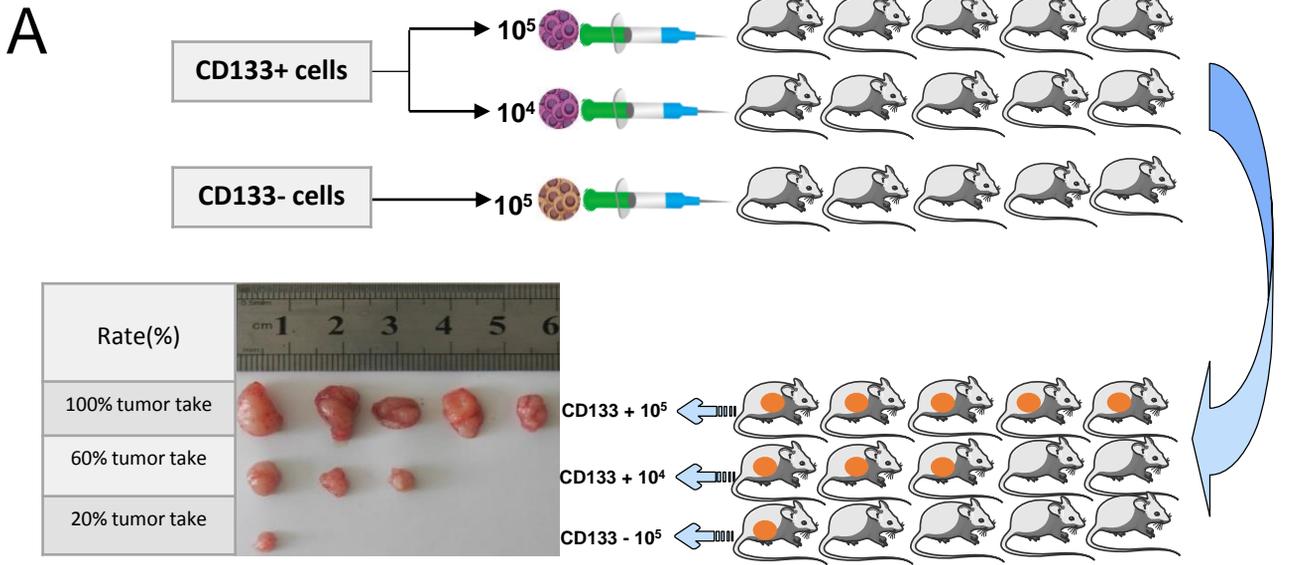


Figure S4

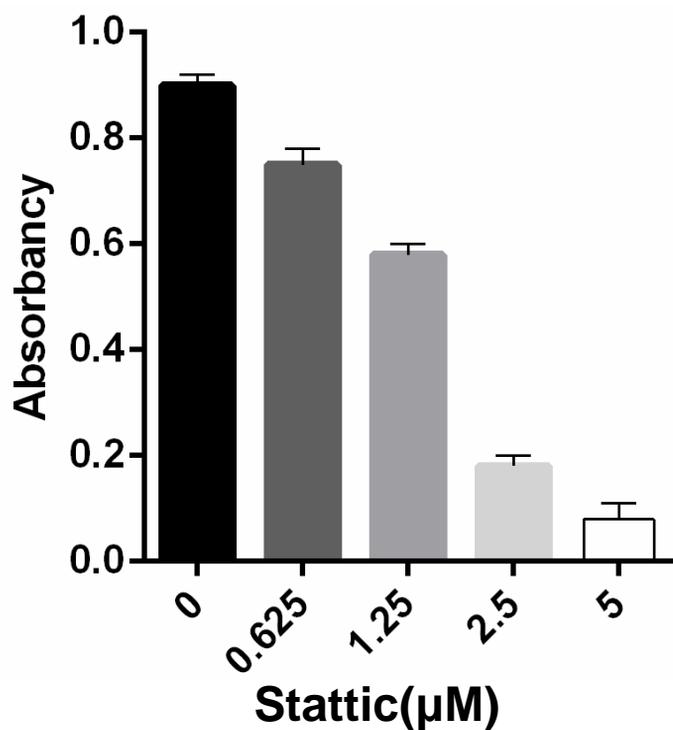
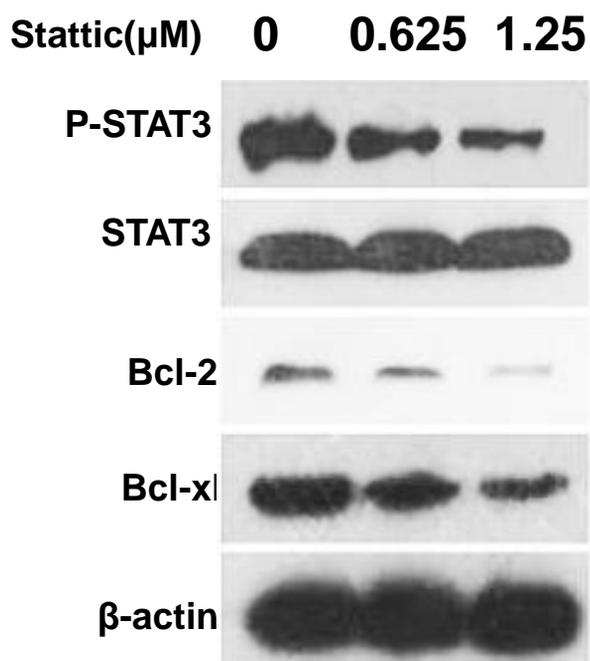
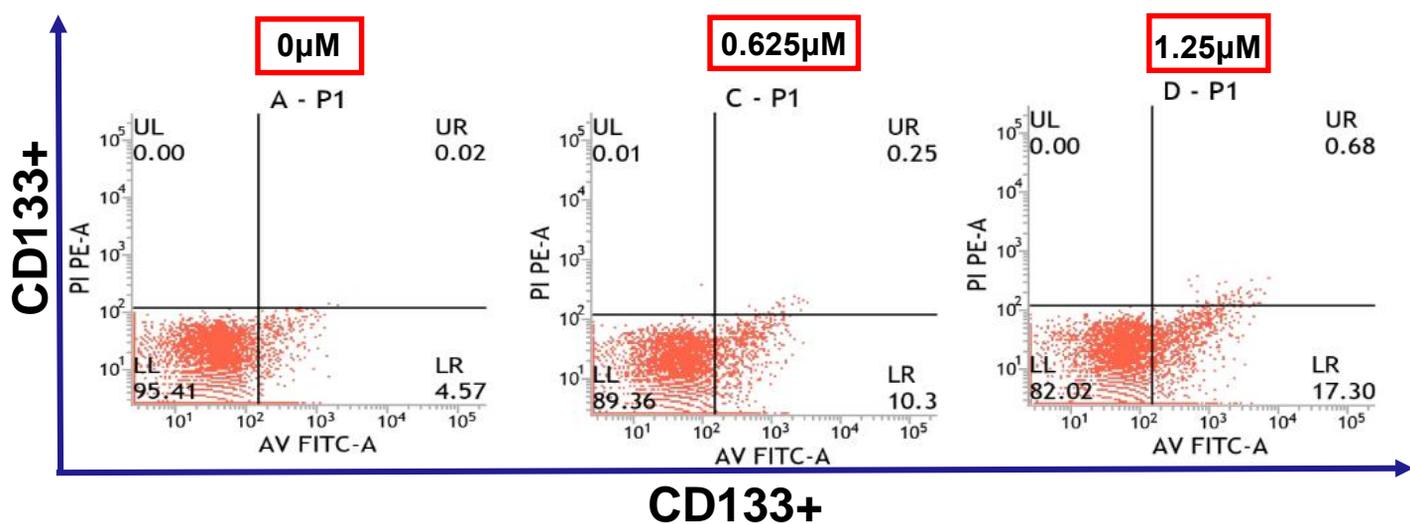
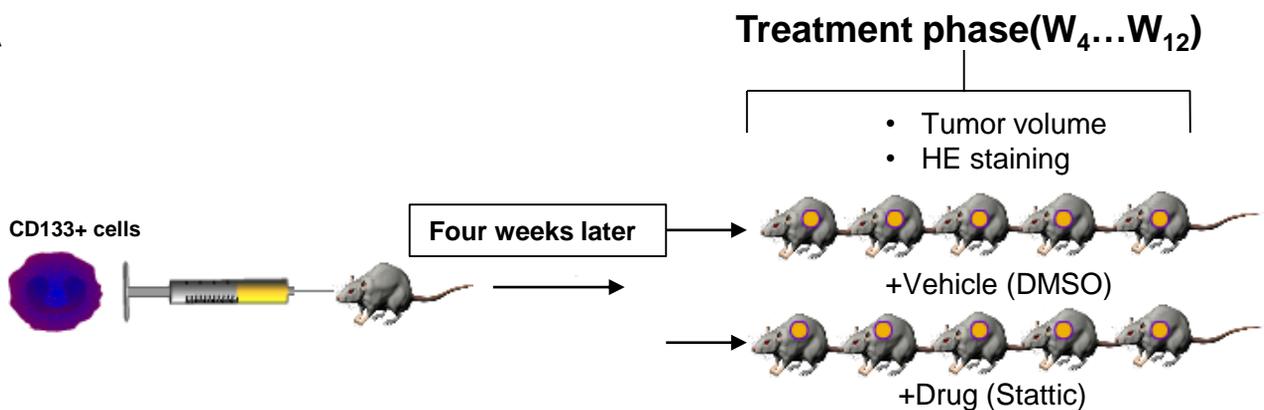
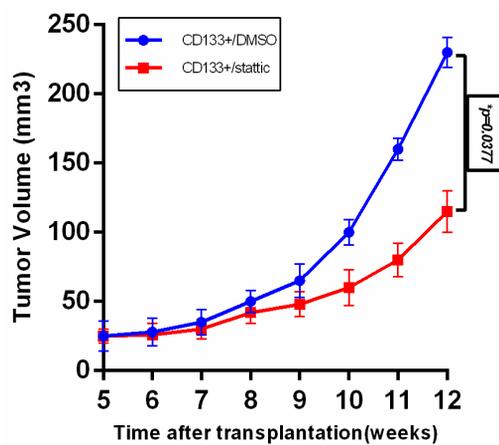
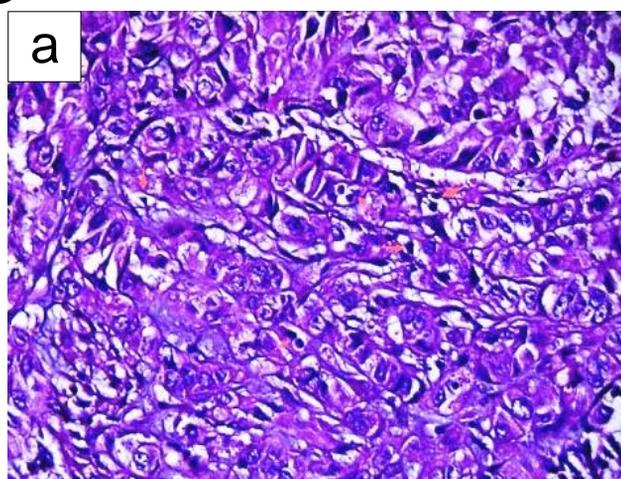
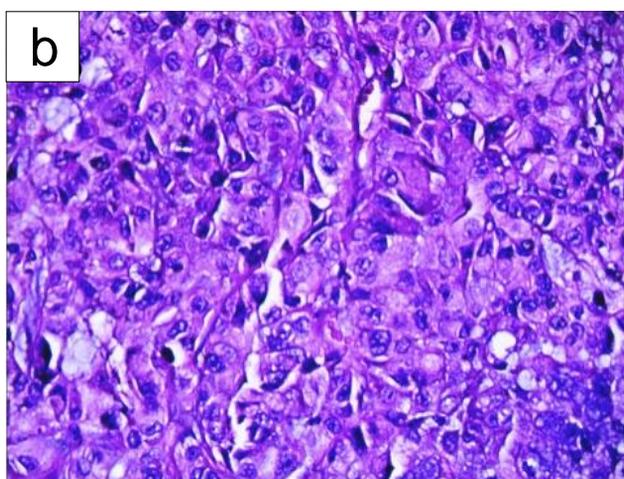
A**C****B**

Figure S5

A**B****C****D**

CD133+/Stattic



CD133+/DMSO

Gene Name	Sequence(5'-3')	Tm	Amplicon length(bp)
SOX2	F:CTGAGCGCCCTGCAGTACAA	64.8	82
	R:GCGAGTAGGACATGCTGTAGGTG	63.3	
NANOG	F:CCTGTGATTTGTGGCCTGA	64.4	168
	R:CTCTGCAGAAGTGGGTGTTTG	62.1	
OCT	F:GAGGAAGCTGACAACAATGAAAATC	62.4	109
	R:GGTTGCCTCTCACTCGGTTC	62.2	
GADPH	F:GCACCGTCAAGGCTGAGAAC	63.3	138
	R:TGGTGAAGACGCCAGTGGA	64.0	