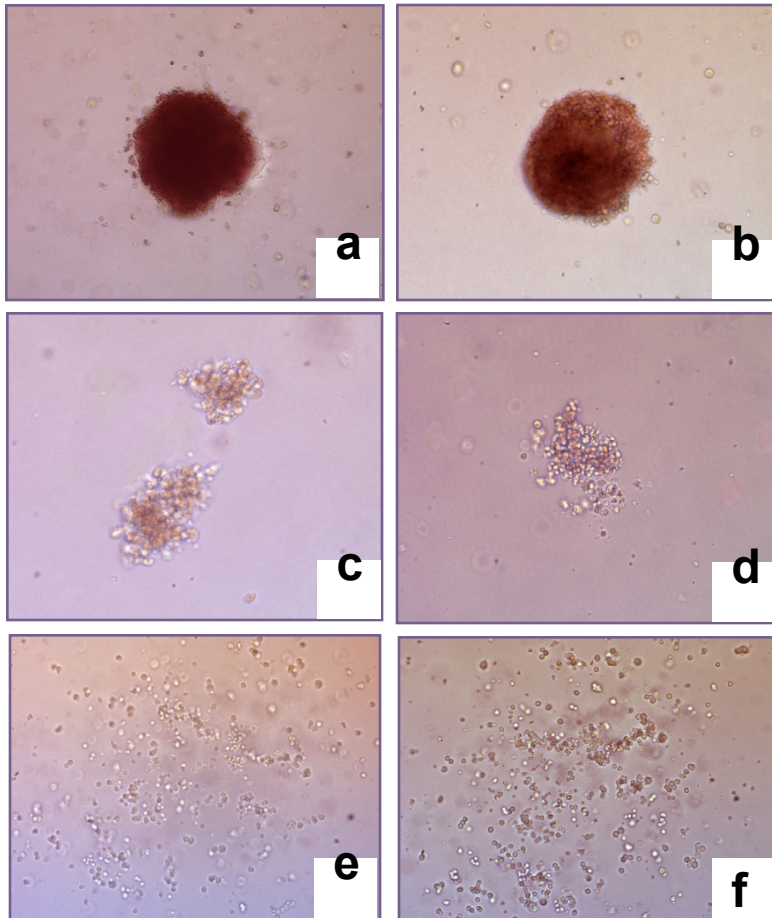
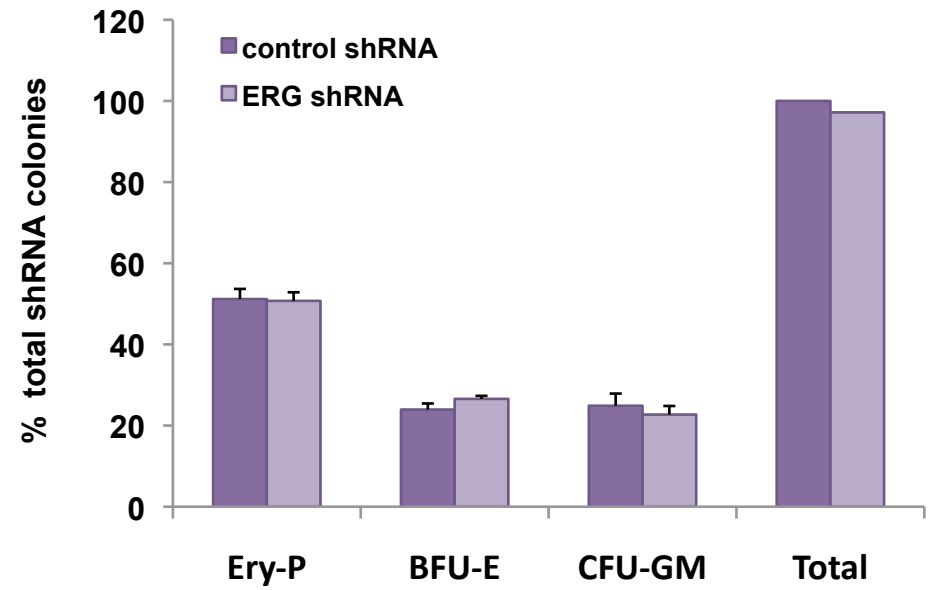


A**B****C**

colonies	Ery-P	BFU-E	CFU-GM	Total #
control shRNA	54.5±3.54	25.5±3.54	26.5±2.12	106.5
ERG shRNA	52.5±3.54	28±0.71	24±2.12	104.5

Average # of colonies/plate (n=3)