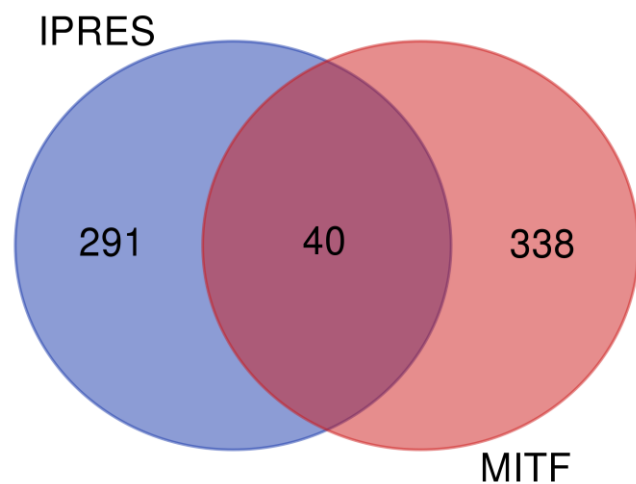
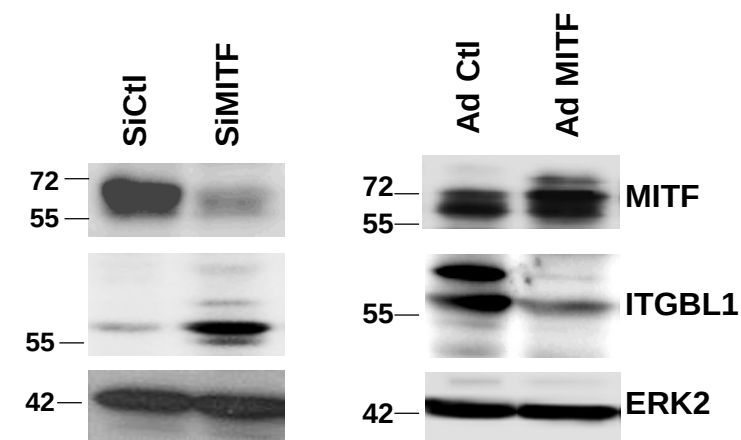
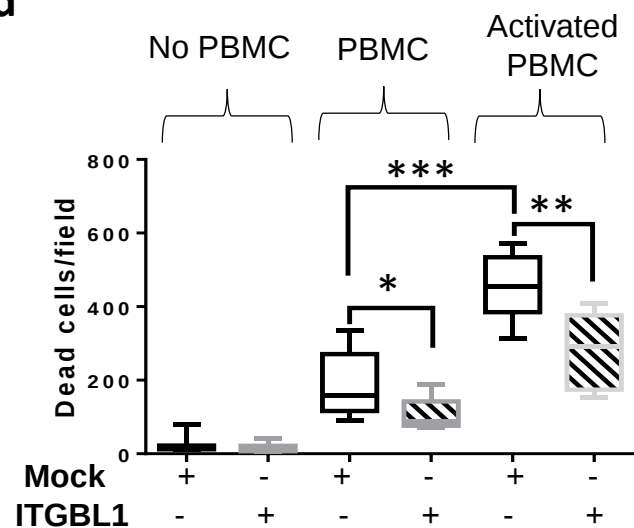
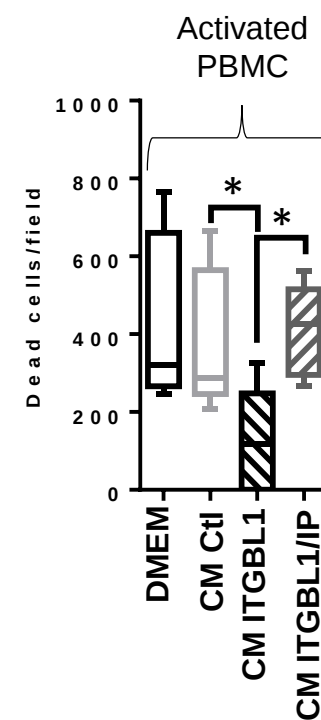
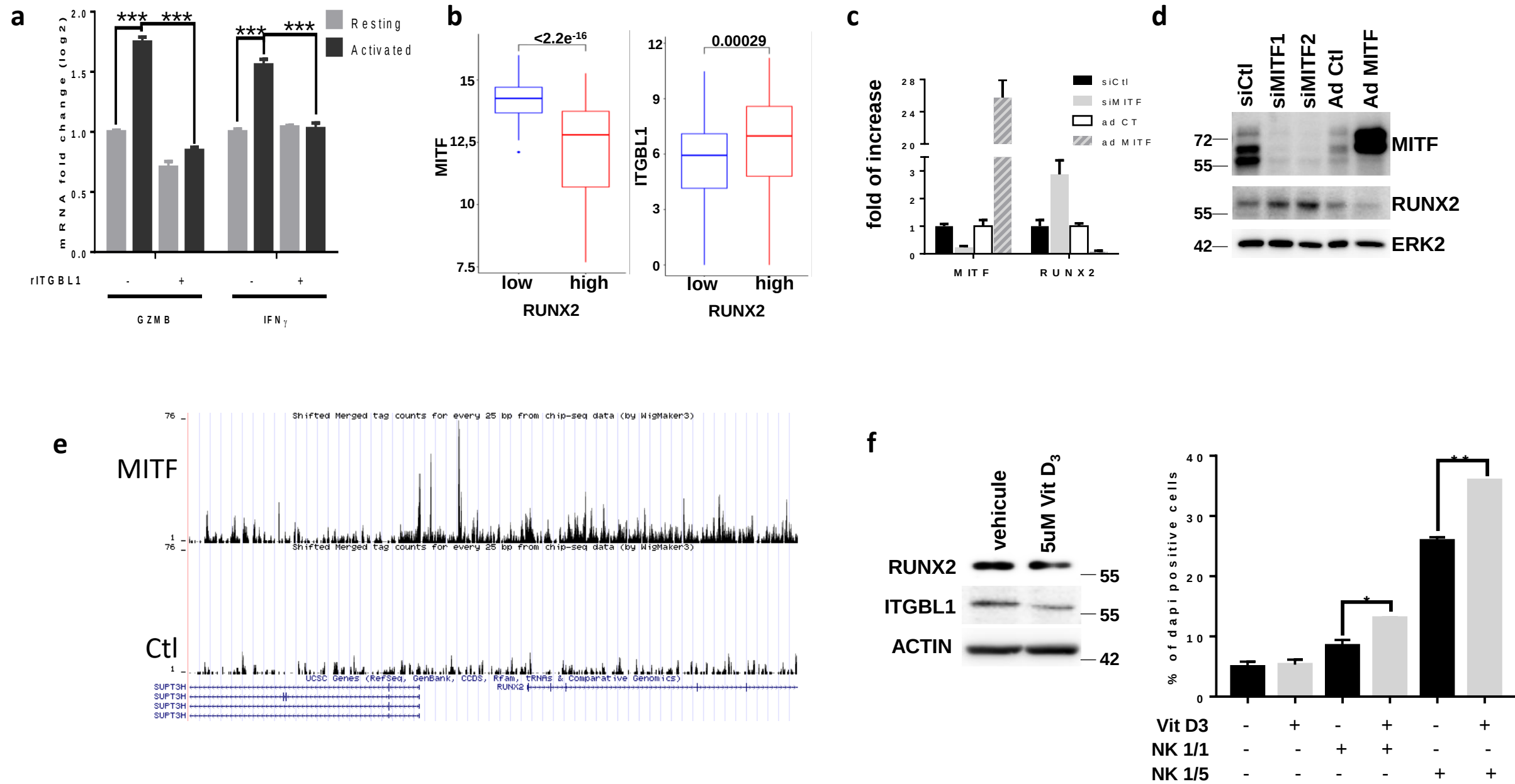


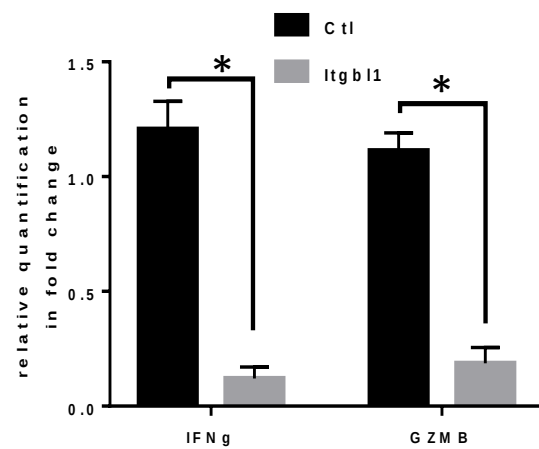
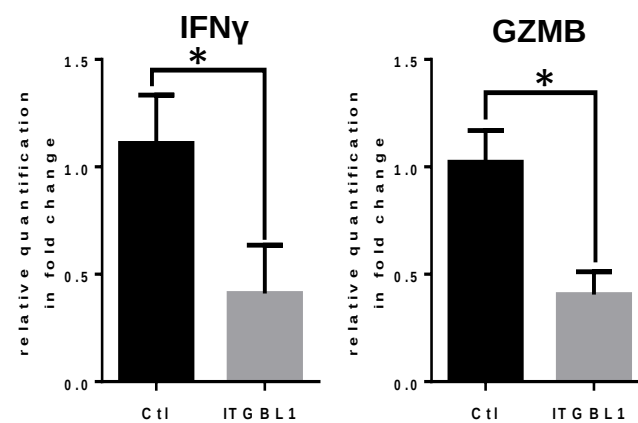
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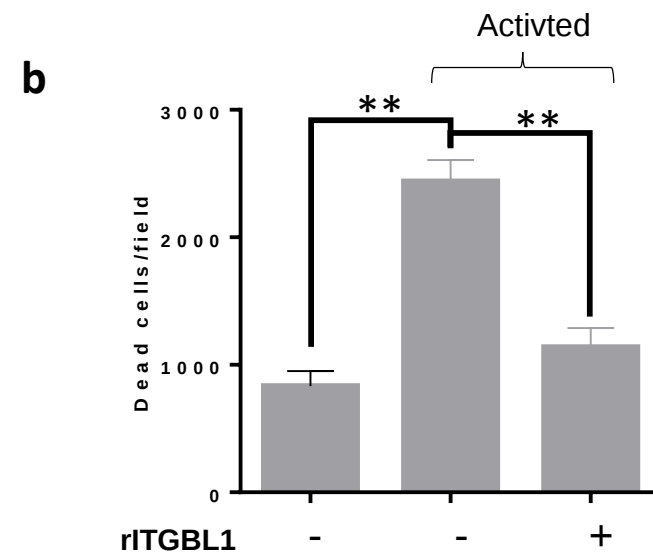
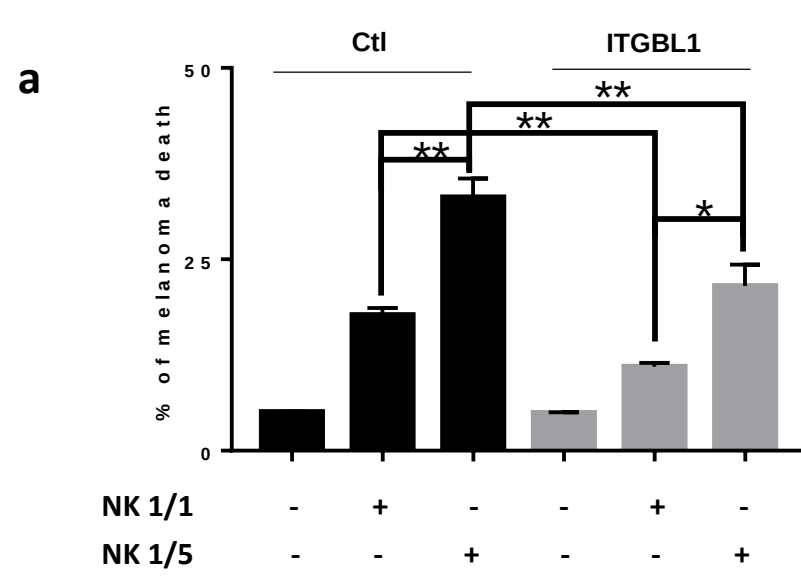
Gene	MITF			IPRES	
	Log FC	P-value		Log FC	P-value
INHBA	4,3198	5,99E-09		2,6198	1,48E-02
FBLN1	1,3935	7,17E-06		2,5892	2,20E-03
COL12A1	2,4904	8,14E-06		2,3880	4,64E-02
PXDN	5,7379	2,84E-12		1,7923	5,18E-03
COL6A1	2,2295	4,17E-07		1,7475	1,95E-02
VCAN	3,7303	3,03E-06		1,7074	1,29E-02
COL6A2	3,3997	4,45E-09		1,6218	1,48E-02
FBN2	3,2584	4,63E-07		1,5996	3,72E-03
F3	2,4917	9,32E-07		1,5700	5,18E-03
OLFML2B	2,2038	7,72E-06		1,5142	1,95E-02
IGFBP6	3,7212	9,47E-09		1,4760	4,64E-02
ESM1	2,8882	8,98E-06		1,4639	4,13E-02
ITGBL1	3,1480	1,50E-05		1,4087	4,13E-02
SERPINE1	4,7582	8,27E-09		1,3778	2,53E-02
LOXL2	3,0618	1,23E-08		1,3461	2,22E-02
STC2	3,6063	1,63E-06		1,2594	1,26E-03
VEGFC	4,5247	1,91E-08		1,2444	1,12E-02

c**d****e**

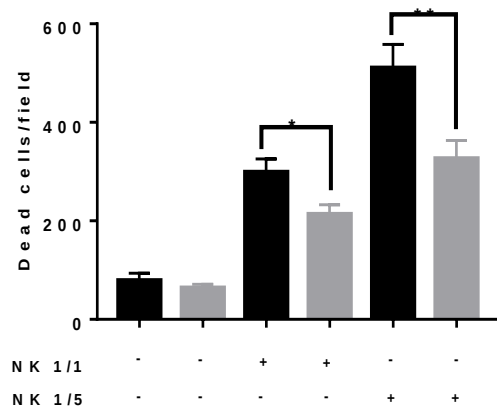


Sup figure 2

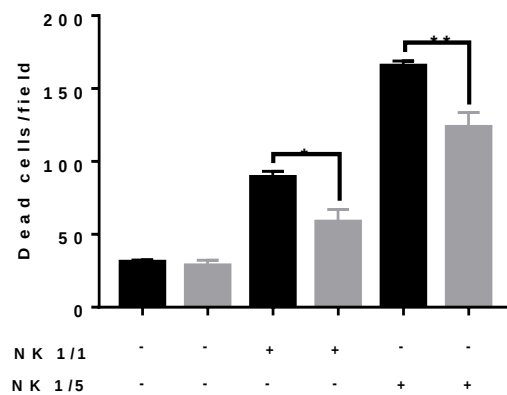
a**b**



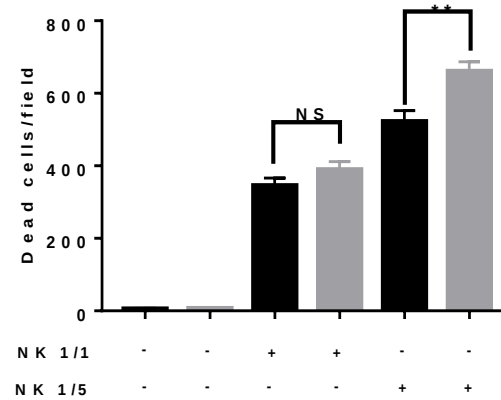
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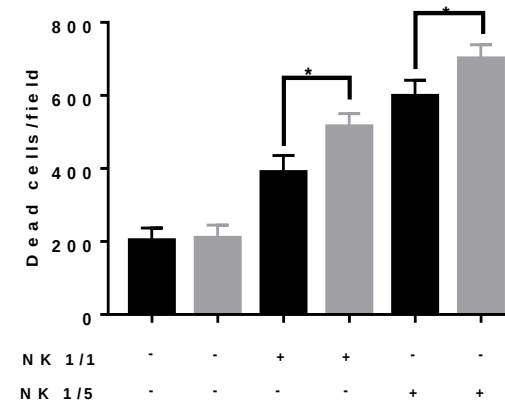
C-13-08



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