

Suppl. Table 4. Log2 Fold Changes (FC) of some genes differentially expressed in PTC with BRAFV600E and hTERT mutation vs BRAFWT-PTC

	log base 2 FC	AveExpr	t	P.Value	FC
FN1	3.96557226	16.72078701	8.365631193	6.70E-16	15.62270385
ITGA3	1.174065803	13.86520533	7.050050164	6.34E-12	2.256467192
VEGFA	-1.226656415	12.5702292	-6.548741432	1.51E-10	-2.340239876
COL1A1	2.707736566	12.72964536	6.307157417	6.51E-10	6.532958911
FLT4	-1.27026472	9.124730001	-5.999299034	3.93E-09	-2.412058204
KDR	-1.295654761	10.6454029	-5.748850397	1.61E-08	-2.45488384
THBS2	2.189221655	8.73434242	5.716896922	1.92E-08	4.560593724
FLT1	-1.003574098	11.23411178	-4.744969669	2.76E-06	-2.004960894
FIGF	-0.848506092	1.375784482	-4.367252926	1.55E-05	-1.800635404
THBS1	1.298426293	10.21206809	4.120972486	4.45E-05	2.4596044
PDGFB	-0.569556193	9.1468419	-3.901077156	0.000109576	-1.484066967
SRC	0.353283545	9.980251825	3.298868091	0.001043635	1.277464805
ITGB1	0.293857573	12.86294234	3.082108375	0.002174802	1.225913829
TGFB1	0.431832536	10.4253761	2.704218341	0.007092145	1.348945944
VEGFC	-0.236300956	9.996714091	-2.239711118	0.025571327	-1.177968496
TGFB2	0.579122933	8.547815665	2.221945448	0.026757151	1.493940751