

Suppl. Table 2. Log2 Fold Changes (FC) of some genes differentially expressed in BRAFV600E-PTC vs BRAFWT-PTC

	log base 2 FC	AveExpr	t	P.Value	FC
FN1	3.503846518	16.72078701	17.06815105	2.72E-51	11.34391339
ITGA3	1.098755555	13.86520533	15.23524384	5.77E-43	2.141698734
FIGF	-0.818465792	1.375784482	-9.727527552	1.62E-20	-1.763529602
VEGFA	-0.755816866	12.5702292	-9.317513801	4.46E-19	-1.688587414
FLT4	-0.768900806	9.124730001	-8.385426309	5.79E-16	-1.703971028
COL1A1	1.464970136	12.72964536	7.879607575	2.26E-14	2.760577561
FLT1	-0.706200224	11.23411178	-7.710106995	7.42E-14	-1.631501399
PDGFB	-0.470516399	9.1468419	-7.441684995	4.70E-13	-1.385605343
KDR	-0.686536296	10.6454029	-7.0340228	7.03E-12	-1.609414902
THBS2	1.124692302	8.73434242	6.781923802	3.53E-11	2.180550348
ITGB1	0.277196372	12.86294234	6.713470983	5.43E-11	1.211837598
SMAD3	0.310456268	11.11605421	6.679642469	6.71E-11	1.240099833
TGFB2	0.744569111	8.547815665	6.596540016	1.12E-10	1.675473781
SRC	0.279922124	9.980251825	6.035695509	3.19E-09	1.214129344
CSPG4	-0.553701655	8.790612582	-6.021430487	3.46E-09	-1.467847058
FBXW11	0.170641797	10.12396311	4.757978887	2.60E-06	1.12555909
THBS1	0.644020019	10.21206809	4.71987327	3.11E-06	1.562677444
PECAM1	-0.301894147	10.42164559	-4.701309833	3.39E-06	-1.232761872
TGFB1	0.317298219	10.4253761	4.588202581	5.73E-06	1.245994949
THY1	-0.368367743	10.91049017	-4.20727663	3.09E-05	-1.290891496
SMAD2	0.062723724	11.13886159	2.684588622	0.00751564	1.044435734
VEGFC	-0.100769036	9.996714091	-2.20547531	0.027898756	-1.072344929
PDGFA	0.122203965	7.981681542	1.999810592	0.046089401	1.088396306