

Table S2. DE-miRNAs from TCGA dataset

miRNA ID	Log fold change	False discovery rate	Tumor	Normal	Delta
Upregulated					
hsa-mir-217	4.237359	0.00095	2742.886	145.6502	11622.59
hsa-mir-224	3.02565	0.027267	349.2049	42.85636	1056.572
hsa-mir-183	2.730561	0.048386	4087.799	617.5533	11161.99
Downregulated					
hsa-mir-142	-1.95755	0.011533	1599.147	6242.973	3130.404
hsa-mir-199b	-1.72226	0.046202	946.2798	3129.33	1629.743
hsa-mir-150	-2.17145	0.002423	471.3316	2131.151	1023.473
hsa-mir-424	-2.72822	1.42E-05	140.6106	936.6654	383.6167
hsa-mir-215	-3.54143	4.24E-10	93.39001	1090.393	330.7346
hsa-mir-3607	-2.83414	7.02E-06	50.00577	358.5205	141.7232