

**A****TCGA-DD-A39Y-01**

	Gene expression	Gene score	Gene weight	Score x weight
RALA	Above median	0	3.6386383	0
PLOD2	Above median	0	4.571733	0
PPT1	Above median	0	3.6378715	0
CTSC	Above median	0	2.929253	0
ETV5	Above median	0	3.15026	0
ALDOA	Above median	0	3.5781527	0
ARPC2	Above median	0	3.4750159	0
SLC38A1	Above median	0	4.9902163	0
TM4SF1	Above median	0	3.6512656	0
PFKFB3	Above median	0	4.242654	0
RGN	below median	0	3.4342885	0
EPHX2	below median	0	4.14249	0
IQGAP2	below median	0	3.2472568	0
SLC22A7	below median	0	3.2640932	0
CYP27A1	below median	0	3.2004652	0
COBLL1	below median	0	3.4549387	0
CYP4F12	below median	0	4.133354	0
APOC3	below median	0	3.7753599	0
CDO1	below median	0	3.1609652	0
PAH	below median	0	3.727815	0
ITPR2	below median	0	3.3972855	0
SERPINA10	below median	0	3.569119	0
SLC2A2	below median	0	2.9978447	0

mHPS 0

**B****TCGA-LG-A9QC-01**

	Gene expression	Gene score	Gene weight	Score x weight
RALA	below median	1	3.6386383	3.6386383
PLOD2	below median	1	4.571733	4.571733
PPT1	below median	1	3.6378715	3.6378715
CTSC	Above median	0	2.929253	0
ETV5	below median	1	3.15026	3.15026
ALDOA	below median	1	3.5781527	3.5781527
ARPC2	below median	1	3.4750159	3.4750159
SLC38A1	below median	1	4.9902163	4.9902163
TM4SF1	below median	1	3.6512656	3.6512656
PFKFB3	below median	1	4.242654	4.242654
RGN	Above median	1	3.4342885	3.4342885
EPHX2	Above median	1	4.14249	4.14249
IQGAP2	Above median	1	3.2472568	3.2472568
SLC22A7	below median	0	3.2640932	0
CYP27A1	Above median	1	3.2004652	3.2004652
COBLL1	Above median	1	3.4549387	3.4549387
CYP4F12	Above median	1	4.133354	4.133354
APOC3	Above median	1	3.7753599	3.7753599
CDO1	Above median	1	3.1609652	3.1609652
PAH	Above median	1	3.727815	3.727815
ITPR2	Above median	1	3.3972855	3.3972855
SERPINA10	Above median	1	3.569119	3.569119
SLC2A2	Above median	1	2.9978447	2.9978447

mHPS 77.17699