Table 4 - Differentially expressed in Glioblastoma multiforme (GBM)

| Gene Symbol | Median (tumor) | Median (normal) | Log2 (Fold Change) | P-value (adj) |
| --- | --- | --- | --- | --- |
| Down |  |  |  |  |
| *TPPP* | 5.000 | 122.947 | -4.369 | 5.73e-104 |
| *AP3B2* | 14.100 | 91.829 | -2.620 | 1.64e-78 |
| *NAGPA* | 8.740 | 32.819 | -1.796 | 2.48e-63 |
| *SLC13A5* | 0.500 | 4.210 | -1.796 | 2.58e-44 |
| *IDS* | 67.991 | 229.586 | -1.741 | 3.97e-50 |
| *MAPK10* | 28.680 | 88.769 | -1.597 | 1.23e-44 |
| *ATP6V0A1* | 63.782 | 172.421 | -1.421 | 1.92e-44 |
| *SLC17A6* | 0.140 | 1.990 | -1.391 | 6.93e-39 |
| *ATP6V1H* | 46.122 | 121.591 | -1.379 | 7.46e-45 |
| *ABCA2* | 63.241 | 164.677 | -1.367 | 1.32e-34 |

Up

| Gene Symbol | Median (tumor) | Median (normal) | Log2 (Fold Change) | P-value (adj) |
| --- | --- | --- | --- | --- |
| *GLB1* | 38.861 | 9.970 | 1.861 | 9.11e-123 |
| *NAGLU* | 29.900 | 6.010 | 2.140 | 4.07e-126 |
| *DNASE2* | 53.671 | 9.830 | 2.336 | 1.04e-148 |
| *GNS* | 52.980 | 9.270 | 2.394 | 3.10e-122 |
| *NAGA* | 23.760 | 3.530 | 2.450 | 1.85e-142 |
| *MAN2B1* | 46.701 | 7.190 | 2.542 | 4.33e-119 |
| *NPC2* | 313.603 | 31.710 | 3.266 | 1.06e-132 |
| *CTSC* | 59.169 | 3.330 | 3.797 | 9.25e-134 |
| *CTSS* | 54.719 | 2.020 | 4.206 | 1.26e-128 |
| *CD68* | 130.879 | 2.150 | 5.388 | 7.83e-137 |