**Supplementary Table 2 |** Expression levels and methylation levels of SNORA71B in gliomas (TCGA Database).

|  |  |  |
| --- | --- | --- |
| **Sample** | **Expression level** | **Methylation level** |
| TCGA-06-0125 | 1.963229 | 0.471975 |
| TCGA-06-0152 | 1.343174 | 0.568625 |
| TCGA-06-0171 | 0.558212 | 0.642425 |
| TCGA-06-0190 | 0.876167 | 0.521875 |
| TCGA-06-0210 | 1.258648 | 0.6009 |
| TCGA-06-0211 | 1.305847 | 0.49885 |
| TCGA-06-0221 | 0.733474 | 0.529125 |
| TCGA-06-1804 | 1.086429 | 0.6841 |
| TCGA-06-5408 | 0.27125 | 0.494325 |
| TCGA-06-5410 | 0.40658 | 0.692725 |
| TCGA-06-5411 | 0.544327 | 0.5386 |
| TCGA-06-5412 | 0.276446 | 0.6616 |
| TCGA-06-5413 | 0.363729 | 0.556425 |
| TCGA-06-5414 | 0.529578 | 0.5384 |
| TCGA-06-5416 | 0.86137 | 0.398975 |
| TCGA-06-5417 | 1.778802 | 0.71675 |
| TCGA-06-5418 | 0.872594 | 0.532475 |
| TCGA-06-5856 | 1.279513 | 0.5313 |
| TCGA-06-5858 | 0.581543 | 0.5012 |
| TCGA-06-5859 | 0.412322 | 0.708525 |
| TCGA-12-5295 | 0 | 0.542775 |
| TCGA-12-5299 | 0.492967 | 0.515175 |
| TCGA-14-0736 | 0 | 0.62655 |
| TCGA-14-0781 | 0.521481 | 0.713625 |
| TCGA-14-1034 | 0.379883 | 0.489875 |
| TCGA-14-1402 | 0.673793 | 0.500475 |
| TCGA-15-1444 | 1.014737 | 0.71035 |
| TCGA-19-0957 | 0.48475 | 0.6268 |
| TCGA-19-1389 | 0 | 0.6177 |
| TCGA-19-4065 | 0.72885 | 0.429025 |
| TCGA-19-5960 | 1.29364 | 0.657225 |
| TCGA-26-1442 | 0.847648 | 0.63545 |
| TCGA-26-5132 | 1.2104 | 0.6712 |
| TCGA-26-5133 | 0.855862 | 0.452475 |
| TCGA-26-5134 | 1.406619 | 0.5082 |
| TCGA-26-5135 | 0.480486 | 0.563925 |
| TCGA-26-5136 | 0 | 0.61365 |
| TCGA-26-5139 | 0.857096 | 0.526575 |
| TCGA-28-5204 | 0.213309 | 0.54025 |
| TCGA-28-5207 | 0.416976 | 0.6796 |
| TCGA-28-5208 | 0.368252 | 0.51645 |
| TCGA-28-5209 | 0.500679 | 0.502925 |
| TCGA-28-5213 | 0.417737 | 0.64585 |
| TCGA-28-5215 | 0.392595 | 0.5244 |
| TCGA-28-5216 | 0.730545 | 0.43575 |
| TCGA-28-5218 | 0.580237 | 0.45255 |
| TCGA-28-5220 | 0.218929 | 0.535775 |
| TCGA-32-1980 | 0.526139 | 0.6357 |
| TCGA-32-5222 | 0.350186 | 0.56935 |
| TCGA-41-5651 | 1.296408 | 0.750325 |
| TCGA-76-4925 | 0.916676 | 0.4884 |
| TCGA-76-4926 | 0.604246 | 0.53395 |
| TCGA-76-4927 | 0.5501 | 0.57315 |
| TCGA-76-4928 | 0.240536 | 0.62595 |
| TCGA-76-4929 | 0.796717 | 0.5755 |
| TCGA-76-4931 | 1.348111 | 0.485075 |
| TCGA-76-4932 | 0.366292 | 0.5172 |
| TCGA-CS-4938 | 0.167264 | 0.8068 |
| TCGA-CS-4941 | 0.537784 | 0.63035 |
| TCGA-CS-4942 | 0.356172 | 0.838975 |
| TCGA-CS-4943 | 0 | 0.853475 |
| TCGA-CS-4944 | 0.086447 | 0.7324 |
| TCGA-CS-5390 | 0.709496 | 0.647475 |
| TCGA-CS-5393 | 0.093865 | 0.81055 |
| TCGA-CS-5394 | 0.375954 | 0.844825 |
| TCGA-CS-5395 | 0.170606 | 0.622375 |
| TCGA-CS-5396 | 0.254594 | 0.6673 |
| TCGA-CS-5397 | 0.554684 | 0.579975 |
| TCGA-CS-6186 | 0.685655 | 0.718175 |
| TCGA-CS-6188 | 0.462833 | 0.5659 |
| TCGA-CS-6290 | 0.250946 | 0.646325 |
| TCGA-CS-6666 | 0.798532 | 0.811525 |
| TCGA-CS-6667 | 0.293773 | 0.8455 |
| TCGA-CS-6668 | 0.691907 | 0.7899 |
| TCGA-CS-6669 | 0.609518 | 0.7109 |
| TCGA-CS-6670 | 0.24378 | 0.82295 |
| TCGA-DB-5270 | 0.280956 | 0.685425 |
| TCGA-DB-5273 | 0.082014 | 0.7432 |
| TCGA-DB-5274 | 0 | 0.813475 |
| TCGA-DB-5275 | 0.345309 | 0.7469 |
| TCGA-DB-5277 | 0.089263 | 0.775175 |
| TCGA-DB-5278 | 0.767869 | 0.83195 |
| TCGA-DB-5279 | 0.545949 | 0.665075 |
| TCGA-DB-5280 | 0.743537 | 0.864375 |
| TCGA-DB-5281 | 0 | 0.848975 |
| TCGA-DB-A4X9 | 0.353247 | 0.845775 |
| TCGA-DB-A4XA | 0.239188 | 0.74175 |
| TCGA-DB-A4XB | 0.497209 | 0.82715 |
| TCGA-DB-A4XC | 0.4156 | 0.745125 |
| TCGA-DB-A4XD | 0.410369 | 0.867325 |
| TCGA-DB-A4XE | 0.493249 | 0.762575 |
| TCGA-DB-A4XF | 0.471471 | 0.6875 |
| TCGA-DB-A4XG | 0.476162 | 0.75985 |
| TCGA-DB-A4XH | 0.480838 | 0.76265 |
| TCGA-DB-A64L | 1.097083 | 0.792475 |
| TCGA-DB-A64O | 0.413676 | 0.60855 |
| TCGA-DB-A64P | 0.281945 | 0.736975 |
| TCGA-DB-A64Q | 0.539018 | 0.727025 |
| TCGA-DB-A64R | 0.683739 | 0.736325 |
| TCGA-DB-A64S | 0.992645 | 0.82235 |
| TCGA-DB-A64U | 0.687017 | 0.783025 |
| TCGA-DB-A64V | 0.423645 | 0.671775 |
| TCGA-DB-A64W | 1.061869 | 0.714875 |
| TCGA-DB-A64X | 0 | 0.687125 |
| TCGA-DB-A75K | 0.461176 | 0.7761 |
| TCGA-DB-A75L | 0.936724 | 0.706475 |
| TCGA-DB-A75M | 0.738713 | 0.719725 |
| TCGA-DB-A75O | 0.2227 | 0.782 |
| TCGA-DB-A75P | 0.181027 | 0.587725 |
| TCGA-DH-5140 | 0.291641 | 0.498725 |
| TCGA-DH-5141 | 0.476085 | 0.844875 |
| TCGA-DH-5142 | 0.255715 | 0.8374 |
| TCGA-DH-5143 | 0.08437 | 0.7564 |
| TCGA-DH-5144 | 0.361491 | 0.807125 |
| TCGA-DH-A669 | 0.447536 | 0.804175 |
| TCGA-DH-A66B | 0.281314 | 0.773725 |
| TCGA-DH-A66D | 0.086505 | 0.833025 |
| TCGA-DH-A66F | 0.460503 | 0.7527 |
| TCGA-DH-A66G | 0.610109 | 0.705925 |
| TCGA-DH-A7UR | 0.44095 | 0.73535 |
| TCGA-DH-A7US | 0.816666 | 0.7473 |
| TCGA-DH-A7UT | 0.694227 | 0.7295 |
| TCGA-DH-A7UU | 0.797787 | 0.8005 |
| TCGA-DH-A7UV | 1.086305 | 0.74505 |
| TCGA-DU-5847 | 0.347121 | 0.584225 |
| TCGA-DU-5849 | 0.445343 | 0.72845 |
| TCGA-DU-5852 | 0.441138 | 0.517975 |
| TCGA-DU-5853 | 0.87334 | 0.744625 |
| TCGA-DU-5854 | 0.643494 | 0.5949 |
| TCGA-DU-5855 | 0.552923 | 0.72535 |
| TCGA-DU-5870 | 0.499482 | 0.6731 |
| TCGA-DU-5871 | 0.16368 | 0.86265 |
| TCGA-DU-5872 | 0.279174 | 0.589025 |
| TCGA-DU-5874 | 0.448789 | 0.866625 |
| TCGA-DU-6392 | 0.907753 | 0.50615 |
| TCGA-DU-6393 | 0.624926 | 0.846775 |
| TCGA-DU-6394 | 0.520503 | 0.699175 |
| TCGA-DU-6395 | 0.22251 | 0.834875 |
| TCGA-DU-6396 | 0.621056 | 0.742975 |
| TCGA-DU-6397 | 0.301957 | 0.66675 |
| TCGA-DU-6399 | 0.548371 | 0.788925 |
| TCGA-DU-6400 | 0.688545 | 0.837375 |
| TCGA-DU-6401 | 0.165839 | 0.85585 |
| TCGA-DU-6402 | 0.403973 | 0.5802 |
| TCGA-DU-6403 | 0.319171 | 0.553675 |
| TCGA-DU-6404 | 0.707812 | 0.572975 |
| TCGA-DU-6405 | 0.690027 | 0.574775 |
| TCGA-DU-6406 | 0.575808 | 0.5074 |
| TCGA-DU-6407 | 0 | 0.7718 |
| TCGA-DU-6408 | 0.411771 | 0.796725 |
| TCGA-DU-6410 | 0.765633 | 0.732425 |
| TCGA-DU-6542 | 1.104217 | 0.695425 |
| TCGA-DU-7006 | 0.605845 | 0.5529 |
| TCGA-DU-7007 | 0.28866 | 0.70175 |
| TCGA-DU-7008 | 0.24413 | 0.879 |
| TCGA-DU-7009 | 0.545656 | 0.81925 |
| TCGA-DU-7010 | 0.073286 | 0.7005 |
| TCGA-DU-7011 | 0.426209 | 0.8023 |
| TCGA-DU-7012 | 0.287882 | 0.65415 |
| TCGA-DU-7013 | 0.27905 | 0.5768 |
| TCGA-DU-7014 | 0.130724 | 0.80525 |
| TCGA-DU-7015 | 0.193821 | 0.743025 |
| TCGA-DU-7018 | 0.227273 | 0.71105 |
| TCGA-DU-7019 | 0.076968 | 0.761075 |
| TCGA-DU-7290 | 0.234458 | 0.6719 |
| TCGA-DU-7292 | 0.458753 | 0.549375 |
| TCGA-DU-7294 | 0.805964 | 0.798525 |
| TCGA-DU-7298 | 0.518652 | 0.829 |
| TCGA-DU-7299 | 0.360418 | 0.76525 |
| TCGA-DU-7300 | 0.358216 | 0.713625 |
| TCGA-DU-7301 | 0.160999 | 0.84915 |
| TCGA-DU-7302 | 0.318233 | 0.707675 |
| TCGA-DU-7304 | 0.153899 | 0.534125 |
| TCGA-DU-7306 | 0.449892 | 0.725875 |
| TCGA-DU-7309 | 0.384729 | 0.7987 |
| TCGA-DU-8158 | 0.384519 | 0.531975 |
| TCGA-DU-8161 | 0.438629 | 0.648725 |
| TCGA-DU-8162 | 0.28641 | 0.656875 |
| TCGA-DU-8163 | 0.356824 | 0.783375 |
| TCGA-DU-8164 | 0.672813 | 0.7585 |
| TCGA-DU-8165 | 0.993939 | 0.598725 |
| TCGA-DU-8166 | 0.155368 | 0.876825 |
| TCGA-DU-8167 | 0.504096 | 0.717275 |
| TCGA-DU-8168 | 0.93428 | 0.7654 |
| TCGA-DU-A5TP | 0.696905 | 0.744825 |
| TCGA-DU-A5TR | 0.21445 | 0.683675 |
| TCGA-DU-A5TS | 0 | 0.74065 |
| TCGA-DU-A5TT | 0.518541 | 0.4947 |
| TCGA-DU-A5TU | 0.203277 | 0.757475 |
| TCGA-DU-A5TW | 0.857724 | 0.6942 |
| TCGA-DU-A5TY | 1.142289 | 0.5561 |
| TCGA-DU-A6S2 | 0.512979 | 0.7251 |
| TCGA-DU-A6S3 | 0.467096 | 0.7514 |
| TCGA-DU-A6S6 | 0.399511 | 0.800275 |
| TCGA-DU-A6S7 | 0.780171 | 0.832625 |
| TCGA-DU-A6S8 | 0.54788 | 0.7965 |
| TCGA-DU-A76K | 0.389338 | 0.556675 |
| TCGA-DU-A76L | 0.687637 | 0.559425 |
| TCGA-DU-A76O | 0.524112 | 0.777 |
| TCGA-DU-A76R | 0.837344 | 0.7931 |
| TCGA-DU-A7T6 | 0.582882 | 0.4533 |
| TCGA-DU-A7T8 | 0.45225 | 0.838925 |
| TCGA-DU-A7TA | 0.439646 | 0.7792 |
| TCGA-DU-A7TB | 0.527179 | 0.489475 |
| TCGA-DU-A7TC | 0.25662 | 0.783625 |
| TCGA-DU-A7TD | 1.115241 | 0.5743 |
| TCGA-DU-A7TG | 0.532744 | 0.6809 |
| TCGA-DU-A7TI | 0.100818 | 0.814675 |
| TCGA-DU-A7TJ | 1.343213 | 0.417425 |
| TCGA-E1-5302 | 0.166365 | 0.799225 |
| TCGA-E1-5303 | 0.179499 | 0.770775 |
| TCGA-E1-5304 | 0 | 0.866475 |
| TCGA-E1-5305 | 0.296139 | 0.8546 |
| TCGA-E1-5307 | 0.090395 | 0.85445 |
| TCGA-E1-5311 | 0.072249 | 0.8389 |
| TCGA-E1-5318 | 0.087243 | 0.85305 |
| TCGA-E1-5319 | 0.094325 | 0.839175 |
| TCGA-E1-5322 | 0.142773 | 0.816 |
| TCGA-E1-A7YD | 0.702265 | 0.478475 |
| TCGA-E1-A7YE | 0.078302 | 0.434025 |
| TCGA-E1-A7YH | 0.399496 | 0.7382 |
| TCGA-E1-A7YI | 0.601536 | 0.528925 |
| TCGA-E1-A7YJ | 1.064214 | 0.544125 |
| TCGA-E1-A7YK | 0.47168 | 0.6494 |
| TCGA-E1-A7YL | 0.909732 | 0.68615 |
| TCGA-E1-A7YM | 1.030201 | 0.668025 |
| TCGA-E1-A7YN | 0.526918 | 0.607275 |
| TCGA-E1-A7YO | 0.432381 | 0.657 |
| TCGA-E1-A7YQ | 0.798588 | 0.675325 |
| TCGA-E1-A7YS | 0.875662 | 0.57285 |
| TCGA-E1-A7YU | 0.685573 | 0.705975 |
| TCGA-E1-A7YV | 1.038251 | 0.653525 |
| TCGA-E1-A7YW | 0.367076 | 0.76845 |
| TCGA-E1-A7YY | 0.50586 | 0.645575 |
| TCGA-E1-A7Z2 | 0.659128 | 0.56955 |
| TCGA-E1-A7Z3 | 0.310526 | 0.7763 |
| TCGA-E1-A7Z4 | 0.414602 | 0.71835 |
| TCGA-E1-A7Z6 | 0.194014 | 0.69145 |
| TCGA-EZ-7264 | 0.80703 | 0.784575 |
| TCGA-F6-A8O3 | 0.308232 | 0.768825 |
| TCGA-F6-A8O4 | 0.745937 | 0.7645 |
| TCGA-FG-5962 | 0.471663 | 0.79135 |
| TCGA-FG-5963 | 0.93421 | 0.655575 |
| TCGA-FG-5964 | 0.095471 | 0.742175 |
| TCGA-FG-5965 | 0.385257 | 0.610363 |
| TCGA-FG-6688 | 0.612257 | 0.695075 |
| TCGA-FG-6689 | 0.275156 | 0.72045 |
| TCGA-FG-6690 | 0.182017 | 0.6765 |
| TCGA-FG-6691 | 0.166336 | 0.789825 |
| TCGA-FG-6692 | 0.420215 | 0.465925 |
| TCGA-FG-7634 | 0.830323 | 0.71285 |
| TCGA-FG-7636 | 0.097062 | 0.822125 |
| TCGA-FG-7637 | 0.504454 | 0.79195 |
| TCGA-FG-7638 | 0.094402 | 0.769775 |
| TCGA-FG-7641 | 0.286933 | 0.77845 |
| TCGA-FG-7643 | 0.361228 | 0.742075 |
| TCGA-FG-8181 | 0.164733 | 0.636475 |
| TCGA-FG-8182 | 0.573749 | 0.814975 |
| TCGA-FG-8185 | 0.486847 | 0.838075 |
| TCGA-FG-8186 | 0.381167 | 0.835975 |
| TCGA-FG-8187 | 0.410271 | 0.727175 |
| TCGA-FG-8188 | 0.424005 | 0.815325 |
| TCGA-FG-8189 | 0.452086 | 0.620375 |
| TCGA-FG-8191 | 0.081445 | 0.6763 |
| TCGA-FG-A4MT | 0.963682 | 0.801975 |
| TCGA-FG-A4MU | 0.585837 | 0.48695 |
| TCGA-FG-A4MW | 0.757094 | 0.557125 |
| TCGA-FG-A4MX | 0.354476 | 0.75595 |
| TCGA-FG-A4MY | 0.150621 | 0.758925 |
| TCGA-FG-A60J | 0.093653 | 0.798025 |
| TCGA-FG-A60K | 0.44183 | 0.795375 |
| TCGA-FG-A60L | 0.567946 | 0.731775 |
| TCGA-FG-A6IZ | 0 | 0.8246 |
| TCGA-FG-A6J1 | 0.112419 | 0.79295 |
| TCGA-FG-A6J3 | 0 | 0.634375 |
| TCGA-FG-A70Y | 0.180728 | 0.74375 |
| TCGA-FG-A70Z | 0.07929 | 0.5536 |
| TCGA-FG-A710 | 0.687756 | 0.7472 |
| TCGA-FG-A711 | 0.097218 | 0.73155 |
| TCGA-FG-A713 | 0.408322 | 0.715925 |
| TCGA-FG-A87N | 0.547649 | 0.48755 |
| TCGA-FG-A87Q | 0.669748 | 0.60535 |
| TCGA-FN-7833 | 0.757872 | 0.8693 |
| TCGA-HT-7467 | 0.245818 | 0.7555 |
| TCGA-HT-7468 | 0.823343 | 0.817825 |
| TCGA-HT-7469 | 1.096785 | 0.76775 |
| TCGA-HT-7470 | 0.336399 | 0.770775 |
| TCGA-HT-7471 | 0.669589 | 0.772475 |
| TCGA-HT-7472 | 0.586823 | 0.748775 |
| TCGA-HT-7473 | 0.316546 | 0.762225 |
| TCGA-HT-7474 | 0.282011 | 0.79155 |
| TCGA-HT-7475 | 0.077709 | 0.855275 |
| TCGA-HT-7476 | 0.395082 | 0.768575 |
| TCGA-HT-7477 | 0.772669 | 0.73325 |
| TCGA-HT-7478 | 0.518485 | 0.69675 |
| TCGA-HT-7479 | 0.29938 | 0.8072 |
| TCGA-HT-7480 | 0.987001 | 0.811125 |
| TCGA-HT-7481 | 0.824039 | 0.773475 |
| TCGA-HT-7482 | 0.242251 | 0.835375 |
| TCGA-HT-7483 | 0.318662 | 0.807125 |
| TCGA-HT-7485 | 0.078947 | 0.8629 |
| TCGA-HT-7601 | 0.310647 | 0.67665 |
| TCGA-HT-7602 | 0.237899 | 0.8088 |
| TCGA-HT-7603 | 0.077769 | 0.7445 |
| TCGA-HT-7604 | 0 | 0.82655 |
| TCGA-HT-7605 | 0.065933 | 0.79225 |
| TCGA-HT-7606 | 0.556569 | 0.7465 |
| TCGA-HT-7607 | 0.522571 | 0.648575 |
| TCGA-HT-7608 | 0.148177 | 0.76305 |
| TCGA-HT-7609 | 0 | 0.828275 |
| TCGA-HT-7610 | 0.23463 | 0.790375 |
| TCGA-HT-7611 | 0.619405 | 0.7666 |
| TCGA-HT-7616 | 0.547299 | 0.61985 |
| TCGA-HT-7620 | 0.576024 | 0.75305 |
| TCGA-HT-7676 | 0.417254 | 0.7756 |
| TCGA-HT-7677 | 0.994853 | 0.798925 |
| TCGA-HT-7680 | 0 | 0.5938 |
| TCGA-HT-7681 | 0.301164 | 0.724475 |
| TCGA-HT-7684 | 0 | 0.790575 |
| TCGA-HT-7686 | 0 | 0.7245 |
| TCGA-HT-7687 | 0.649456 | 0.747675 |
| TCGA-HT-7688 | 0 | 0.7253 |
| TCGA-HT-7689 | 0.36895 | 0.71545 |
| TCGA-HT-7690 | 0.145356 | 0.83225 |
| TCGA-HT-7691 | 0.148198 | 0.585475 |
| TCGA-HT-7692 | 0.663156 | 0.768375 |
| TCGA-HT-7693 | 0 | 0.846475 |
| TCGA-HT-7694 | 0.222382 | 0.77435 |
| TCGA-HT-7695 | 0.121383 | 0.8198 |
| TCGA-HT-7854 | 0.534183 | 0.72245 |
| TCGA-HT-7855 | 0 | 0.85005 |
| TCGA-HT-7856 | 0.205487 | 0.6902 |
| TCGA-HT-7857 | 0.558393 | 0.569325 |
| TCGA-HT-7858 | 0.823759 | 0.69475 |
| TCGA-HT-7860 | 0.806869 | 0.547825 |
| TCGA-HT-7873 | 0.649695 | 0.8431 |
| TCGA-HT-7874 | 0.151456 | 0.666675 |
| TCGA-HT-7875 | 0.458405 | 0.769325 |
| TCGA-HT-7877 | 0.603783 | 0.74455 |
| TCGA-HT-7879 | 0.852351 | 0.77365 |
| TCGA-HT-7880 | 0.13355 | 0.700925 |
| TCGA-HT-7881 | 0.222683 | 0.746875 |
| TCGA-HT-7882 | 1.138275 | 0.5394 |
| TCGA-HT-7884 | 0.983088 | 0.823325 |
| TCGA-HT-7902 | 0.925257 | 0.705625 |
| TCGA-HT-8010 | 0.166412 | 0.6914 |
| TCGA-HT-8011 | 0.554465 | 0.4573 |
| TCGA-HT-8012 | 0.460824 | 0.79455 |
| TCGA-HT-8013 | 0.075509 | 0.840675 |
| TCGA-HT-8015 | 0.100746 | 0.61545 |
| TCGA-HT-8018 | 0.150572 | 0.664725 |
| TCGA-HT-8019 | 0.562759 | 0.63305 |
| TCGA-HT-8104 | 0.205195 | 0.52865 |
| TCGA-HT-8105 | 0.666514 | 0.7424 |
| TCGA-HT-8107 | 0.394086 | 0.6254 |
| TCGA-HT-8108 | 0.252977 | 0.820125 |
| TCGA-HT-8109 | 0.499685 | 0.733375 |
| TCGA-HT-8110 | 0.448906 | 0.6008 |
| TCGA-HT-8111 | 0.351241 | 0.705525 |
| TCGA-HT-8113 | 0.349585 | 0.697025 |
| TCGA-HT-8114 | 0.522815 | 0.692025 |
| TCGA-HT-8558 | 0.07927 | 0.691725 |
| TCGA-HT-8563 | 0.469029 | 0.628 |
| TCGA-HT-8564 | 1.429961 | 0.56385 |
| TCGA-HT-A4DS | 0 | 0.548025 |
| TCGA-HT-A4DV | 0.349482 | 0.7629 |
| TCGA-HT-A5R5 | 0.11211 | 0.7689 |
| TCGA-HT-A5R7 | 0.316836 | 0.633825 |
| TCGA-HT-A5R9 | 0.993534 | 0.79585 |
| TCGA-HT-A5RA | 0.437942 | 0.423725 |
| TCGA-HT-A5RB | 1.187034 | 0.811975 |
| TCGA-HT-A5RC | 0.387431 | 0.5398 |
| TCGA-HT-A614 | 0.224967 | 0.77775 |
| TCGA-HT-A615 | 0.29263 | 0.7758 |
| TCGA-HT-A616 | 0.109763 | 0.737475 |
| TCGA-HT-A617 | 0.30674 | 0.6483 |
| TCGA-HT-A618 | 0.299617 | 0.657675 |
| TCGA-HT-A619 | 1.096516 | 0.829175 |
| TCGA-HT-A61A | 0.302212 | 0.732825 |
| TCGA-HT-A61B | 0 | 0.725825 |
| TCGA-HT-A61C | 0.518462 | 0.502 |
| TCGA-HT-A74H | 0 | 0.599075 |
| TCGA-HT-A74J | 0.6328 | 0.730325 |
| TCGA-HT-A74K | 0 | 0.643475 |
| TCGA-HT-A74L | 0.802287 | 0.711775 |
| TCGA-HT-A74O | 0.102071 | 0.7049 |
| TCGA-HW-7486 | 0.293771 | 0.773275 |
| TCGA-HW-7487 | 0.342261 | 0.729325 |
| TCGA-HW-7489 | 0.080944 | 0.760025 |
| TCGA-HW-7490 | 0.093012 | 0.818725 |
| TCGA-HW-7491 | 0.44091 | 0.78015 |
| TCGA-HW-7493 | 0 | 0.750825 |
| TCGA-HW-7495 | 0.486354 | 0.82915 |
| TCGA-HW-8319 | 0.623337 | 0.709425 |
| TCGA-HW-8320 | 0.561204 | 0.82395 |
| TCGA-HW-8321 | 0.46223 | 0.737825 |
| TCGA-HW-8322 | 0.697705 | 0.819475 |
| TCGA-HW-A5KJ | 0.099422 | 0.7638 |
| TCGA-HW-A5KK | 0.719452 | 0.6265 |
| TCGA-HW-A5KL | 0.345491 | 0.847175 |
| TCGA-HW-A5KM | 0.066477 | 0.6615 |
| TCGA-IK-7675 | 0.648311 | 0.8559 |
| TCGA-IK-8125 | 0.323482 | 0.74575 |
| TCGA-KT-A74X | 0.369813 | 0.7751 |
| TCGA-KT-A7W1 | 1.133811 | 0.501075 |
| TCGA-P5-A5ET | 0.095226 | 0.84475 |
| TCGA-P5-A5EU | 0.567964 | 0.66305 |
| TCGA-P5-A5EV | 0.499394 | 0.7648 |
| TCGA-P5-A5EW | 0 | 0.857325 |
| TCGA-P5-A5EX | 0 | 0.54545 |
| TCGA-P5-A5EY | 0.36907 | 0.56655 |
| TCGA-P5-A5EZ | 0.71099 | 0.6654 |
| TCGA-P5-A5F0 | 1.223134 | 0.7152 |
| TCGA-P5-A5F1 | 0 | 0.822375 |
| TCGA-P5-A5F2 | 1.075412 | 0.636925 |
| TCGA-P5-A5F4 | 0.568683 | 0.723875 |
| TCGA-P5-A5F6 | 0.784809 | 0.6682 |
| TCGA-P5-A72U | 0.09863 | 0.571575 |
| TCGA-P5-A72W | 0 | 0.8417 |
| TCGA-P5-A72X | 0.828254 | 0.688575 |
| TCGA-P5-A72Z | 0.836295 | 0.66685 |
| TCGA-P5-A730 | 0.47567 | 0.782625 |
| TCGA-P5-A731 | 0.425681 | 0.758875 |
| TCGA-P5-A733 | 0 | 0.742275 |
| TCGA-P5-A735 | 0.364807 | 0.83965 |
| TCGA-P5-A736 | 0 | 0.658525 |
| TCGA-P5-A737 | 0.372313 | 0.704175 |
| TCGA-P5-A77W | 1.40675 | 0.798875 |
| TCGA-P5-A77X | 0.407194 | 0.743475 |
| TCGA-P5-A780 | 0.574228 | 0.73245 |
| TCGA-P5-A781 | 0.534795 | 0.7668 |
| TCGA-QH-A65R | 0.179257 | 0.77375 |
| TCGA-QH-A65S | 0.347626 | 0.74215 |
| TCGA-QH-A65V | 0.205403 | 0.789925 |
| TCGA-QH-A65X | 0 | 0.732675 |
| TCGA-QH-A65Z | 0.382041 | 0.793475 |
| TCGA-QH-A6CS | 0.204592 | 0.6236 |
| TCGA-QH-A6CU | 0.492018 | 0.690375 |
| TCGA-QH-A6CV | 0.639622 | 0.725725 |
| TCGA-QH-A6CW | 0.206608 | 0.841125 |
| TCGA-QH-A6CX | 0 | 0.644375 |
| TCGA-QH-A6CY | 0.090169 | 0.767875 |
| TCGA-QH-A6CZ | 0.272179 | 0.814075 |
| TCGA-QH-A6X3 | 0.506503 | 0.817275 |
| TCGA-QH-A6X4 | 0.469911 | 0.852375 |
| TCGA-QH-A6X5 | 0.666774 | 0.75075 |
| TCGA-QH-A6X8 | 0.778557 | 0.828475 |
| TCGA-QH-A6X9 | 0.846833 | 0.69845 |
| TCGA-QH-A6XA | 0.192432 | 0.757075 |
| TCGA-QH-A6XC | 0.268987 | 0.6092 |
| TCGA-QH-A86X | 0.719423 | 0.77495 |
| TCGA-QH-A870 | 0.377135 | 0.850175 |
| TCGA-R8-A6MK | 0.822515 | 0.80735 |
| TCGA-R8-A6ML | 0.243088 | 0.791575 |
| TCGA-R8-A6MO | 0.555829 | 0.796075 |
| TCGA-R8-A73M | 0.239342 | 0.781475 |
| TCGA-RY-A83X | 0.616708 | 0.8121 |
| TCGA-RY-A83Y | 0.70731 | 0.623275 |
| TCGA-RY-A83Z | 0 | 0.7656 |
| TCGA-RY-A840 | 0.531076 | 0.74415 |
| TCGA-RY-A843 | 0.372879 | 0.812725 |
| TCGA-RY-A845 | 0.089486 | 0.837675 |
| TCGA-RY-A847 | 0.112764 | 0.7037 |
| TCGA-S9-A6TS | 0.829949 | 0.7606 |
| TCGA-S9-A6TU | 0.523196 | 0.8086 |
| TCGA-S9-A6TV | 0.14258 | 0.794375 |
| TCGA-S9-A6TW | 0.513554 | 0.697475 |
| TCGA-S9-A6TX | 0.474727 | 0.720725 |
| TCGA-S9-A6TY | 0.881123 | 0.6894 |
| TCGA-S9-A6TZ | 0.459002 | 0.824125 |
| TCGA-S9-A6U0 | 0.122415 | 0.585475 |
| TCGA-S9-A6U1 | 1.018377 | 0.817525 |
| TCGA-S9-A6U2 | 0.198342 | 0.771525 |
| TCGA-S9-A6U5 | 0.986039 | 0.7399 |
| TCGA-S9-A6U6 | 0.351774 | 0.774225 |
| TCGA-S9-A6U8 | 0.911378 | 0.8096 |
| TCGA-S9-A6U9 | 0.191892 | 0.75485 |
| TCGA-S9-A6UA | 0.778501 | 0.5612 |
| TCGA-S9-A6UB | 1.069363 | 0.7798 |
| TCGA-S9-A6WD | 0.851255 | 0.6976 |
| TCGA-S9-A6WE | 0.369284 | 0.80555 |
| TCGA-S9-A6WG | 1.106193 | 0.7769 |
| TCGA-S9-A6WH | 0.24171 | 0.830925 |
| TCGA-S9-A6WI | 0.149142 | 0.6867 |
| TCGA-S9-A6WL | 0.876301 | 0.694875 |
| TCGA-S9-A6WM | 0.240311 | 0.488325 |
| TCGA-S9-A6WN | 0.346696 | 0.588375 |
| TCGA-S9-A6WO | 0.20395 | 0.8568 |
| TCGA-S9-A6WP | 0.886845 | 0.74745 |
| TCGA-S9-A6WQ | 0.266493 | 0.677575 |
| TCGA-S9-A7IQ | 0.47731 | 0.70635 |
| TCGA-S9-A7IS | 0.490788 | 0.606775 |
| TCGA-S9-A7IX | 0.480168 | 0.636475 |
| TCGA-S9-A7IY | 0.520206 | 0.668725 |
| TCGA-S9-A7IZ | 0.505066 | 0.64185 |
| TCGA-S9-A7J0 | 1.485482 | 0.8059 |
| TCGA-S9-A7J1 | 1.02454 | 0.797525 |
| TCGA-S9-A7J2 | 0.73869 | 0.8537 |
| TCGA-S9-A7J3 | 0.92139 | 0.7424 |
| TCGA-S9-A7QW | 0.294467 | 0.8565 |
| TCGA-S9-A7QX | 0.453562 | 0.699 |
| TCGA-S9-A7QY | 1.203758 | 0.73365 |
| TCGA-S9-A7QZ | 0.789896 | 0.8015 |
| TCGA-S9-A7R1 | 0.80714 | 0.7836 |
| TCGA-S9-A7R2 | 0.111609 | 0.6668 |
| TCGA-S9-A7R3 | 0.308828 | 0.8154 |
| TCGA-S9-A7R4 | 0.940721 | 0.809025 |
| TCGA-S9-A7R7 | 0.449101 | 0.693475 |
| TCGA-S9-A7R8 | 0.829075 | 0.787625 |
| TCGA-S9-A89V | 0.892352 | 0.411525 |
| TCGA-S9-A89Z | 0.425055 | 0.71535 |
| TCGA-TM-A7C3 | 1.244668 | 0.594525 |
| TCGA-TM-A7C4 | 0.307612 | 0.747 |
| TCGA-TM-A7C5 | 0.656119 | 0.8526 |
| TCGA-TM-A7CA | 0.493287 | 0.767925 |
| TCGA-TM-A7CF | 0.721865 | 0.79185 |
| TCGA-TM-A84B | 0.460673 | 0.762075 |
| TCGA-TM-A84C | 0.776212 | 0.4779 |
| TCGA-TM-A84F | 0 | 0.742325 |
| TCGA-TM-A84G | 0.939151 | 0.819375 |
| TCGA-TM-A84H | 1.430201 | 0.70975 |
| TCGA-TM-A84I | 1.365172 | 0.6245 |
| TCGA-TM-A84J | 0.628169 | 0.3113 |
| TCGA-TM-A84L | 0.638778 | 0.75995 |
| TCGA-TM-A84M | 1.356413 | 0.589525 |
| TCGA-TM-A84O | 0.54281 | 0.80095 |
| TCGA-TM-A84Q | 0.447008 | 0.74995 |
| TCGA-TM-A84R | 0.342479 | 0.70215 |
| TCGA-TM-A84S | 0.530008 | 0.6565 |
| TCGA-TM-A84T | 0.552457 | 0.849525 |
| TCGA-TQ-A7RF | 0.486568 | 0.759875 |
| TCGA-TQ-A7RG | 0.655143 | 0.796275 |
| TCGA-TQ-A7RH | 0.827099 | 0.74505 |
| TCGA-TQ-A7RI | 1.078853 | 0.68235 |
| TCGA-TQ-A7RJ | 0.566312 | 0.74015 |
| TCGA-TQ-A7RK | 0.680726 | 0.875563 |
| TCGA-TQ-A7RM | 1.499321 | 0.828975 |
| TCGA-TQ-A7RN | 1.007178 | 0.761725 |
| TCGA-TQ-A7RO | 0.703934 | 0.79205 |
| TCGA-TQ-A7RP | 0.586959 | 0.6632 |
| TCGA-TQ-A7RQ | 0.957556 | 0.672875 |
| TCGA-TQ-A7RR | 0.89181 | 0.726575 |
| TCGA-TQ-A7RS | 0.336686 | 0.751275 |
| TCGA-TQ-A7RV | 1.033472 | 0.6935 |
| TCGA-TQ-A7RW | 0.692519 | 0.658 |
| TCGA-TQ-A8XE | 0.730128 | 0.730075 |
| TCGA-VM-A8C8 | 0.68912 | 0.655675 |
| TCGA-VM-A8C9 | 0 | 0.58695 |
| TCGA-VM-A8CA | 0 | 0.706925 |
| TCGA-VM-A8CB | 0.630474 | 0.7048 |
| TCGA-VM-A8CD | 0.58569 | 0.5746 |
| TCGA-VM-A8CE | 0.851864 | 0.767475 |
| TCGA-VM-A8CF | 0 | 0.7802 |
| TCGA-VM-A8CH | 0.713529 | 0.80365 |
| TCGA-VV-A829 | 0.275497 | 0.5097 |
| TCGA-VV-A86M | 0.577223 | 0.770525 |
| TCGA-VW-A7QS | 1.067979 | 0.677325 |
| TCGA-VW-A8FI | 0.127871 | 0.561175 |
| TCGA-W9-A837 | 0.322106 | 0.74855 |
| TCGA-WH-A86K | 0.844136 | 0.8117 |
| TCGA-WY-A858 | 0.762775 | 0.7213 |
| TCGA-WY-A859 | 0.484804 | 0.74025 |
| TCGA-WY-A85A | 0.84041 | 0.7813 |
| TCGA-WY-A85B | 0.61092 | 0.832 |
| TCGA-WY-A85C | 0.535432 | 0.764575 |
| TCGA-WY-A85D | 0.112428 | 0.8107 |
| TCGA-WY-A85E | 0.400487 | 0.77125 |