|  |  |  |  |
| --- | --- | --- | --- |
| Table S2. Target genes of AC007255.1 in ESCA (Spearman’s rank correlation coefficient r>0.4 and p<0.001). | | | |
| eRNA symbol | Target gene | Correlation Coefficient *r* | *p*-value |
| AC007255.1 | PRR15 | 0.936 | < 0.001 |
| AC007255.1 | AC007255.2 | 0.816 | < 0.001 |
| AC007255.1 | CHN2 | 0.803 | < 0.001 |
| AC007255.1 | FAM221A | 0.801 | < 0.001 |
| AC007255.1 | PLEKHA6 | 0.795 | < 0.001 |
| AC007255.1 | MYO5C | 0.793 | < 0.001 |
| AC007255.1 | ICA1 | 0.790 | < 0.001 |
| AC007255.1 | AC006042.1 | 0.788 | < 0.001 |
| AC007255.1 | ARFGEF3 | 0.787 | < 0.001 |
| AC007255.1 | TMEM62 | 0.787 | < 0.001 |
| AC007255.1 | HID1 | 0.781 | < 0.001 |
| AC007255.1 | PLS1 | 0.779 | < 0.001 |
| AC007255.1 | AC008870.6 | 0.778 | < 0.001 |
| AC007255.1 | C9orf152 | 0.777 | < 0.001 |
| AC007255.1 | LINC01342 | 0.777 | < 0.001 |
| AC007255.1 | BCL2L15 | 0.776 | < 0.001 |
| AC007255.1 | AC130456.7 | 0.775 | < 0.001 |
| AC007255.1 | PRR15L | 0.773 | < 0.001 |
| AC007255.1 | AC008870.1 | 0.771 | < 0.001 |
| AC007255.1 | TRIM31-AS1 | 0.770 | < 0.001 |
| AC007255.1 | AC127526.2 | 0.768 | < 0.001 |
| AC007255.1 | RHPN2 | 0.767 | < 0.001 |
| AC007255.1 | AL117382.1 | 0.767 | < 0.001 |
| AC007255.1 | GPR160 | 0.765 | < 0.001 |
| AC007255.1 | CLDN3 | 0.764 | < 0.001 |
| AC007255.1 | LONRF3 | 0.764 | < 0.001 |
| AC007255.1 | AKR7L | 0.762 | < 0.001 |
| AC007255.1 | EPCAM | 0.762 | < 0.001 |
| AC007255.1 | AC130456.2 | 0.762 | < 0.001 |
| AC007255.1 | CRACR2B | 0.761 | < 0.001 |
| AC007255.1 | AL592114.3 | 0.761 | < 0.001 |
| AC007255.1 | AKR7A3 | 0.760 | < 0.001 |
| AC007255.1 | TMC5 | 0.760 | < 0.001 |
| AC007255.1 | PLUT | 0.759 | < 0.001 |
| AC007255.1 | TMEM63A | 0.758 | < 0.001 |
| AC007255.1 | TOX3 | 0.758 | < 0.001 |
| AC007255.1 | ARHGAP12 | 0.758 | < 0.001 |
| AC007255.1 | AC007490.1 | 0.757 | < 0.001 |
| AC007255.1 | PIP5K1B | 0.755 | < 0.001 |
| AC007255.1 | TRIM31 | 0.753 | < 0.001 |
| AC007255.1 | LLGL2 | 0.753 | < 0.001 |
| AC007255.1 | DNMBP-AS1 | 0.753 | < 0.001 |
| AC007255.1 | MIR559 | 0.752 | < 0.001 |
| AC007255.1 | AC130456.3 | 0.751 | < 0.001 |
| AC007255.1 | LINC01814 | 0.749 | < 0.001 |
| AC007255.1 | ARHGEF38 | 0.749 | < 0.001 |
| AC007255.1 | MLPH | 0.748 | < 0.001 |
| AC007255.1 | SYTL2 | 0.747 | < 0.001 |
| AC007255.1 | CGN | 0.745 | < 0.001 |
| AC007255.1 | TMEM125 | 0.743 | < 0.001 |
| AC007255.1 | USH1C | 0.743 | < 0.001 |
| AC007255.1 | LINC02747 | 0.742 | < 0.001 |
| AC007255.1 | PLA2G10 | 0.742 | < 0.001 |
| AC007255.1 | HNF4A-AS1 | 0.741 | < 0.001 |
| AC007255.1 | AC004982.2 | 0.741 | < 0.001 |
| AC007255.1 | MYO15B | 0.741 | < 0.001 |
| AC007255.1 | FA2H | 0.741 | < 0.001 |
| AC007255.1 | PPARG | 0.740 | < 0.001 |
| AC007255.1 | LINC02038 | 0.739 | < 0.001 |
| AC007255.1 | FOXA3 | 0.738 | < 0.001 |
| AC007255.1 | LRRC31 | 0.738 | < 0.001 |
| AC007255.1 | P4HTM | 0.738 | < 0.001 |
| AC007255.1 | ACBD5 | 0.738 | < 0.001 |
| AC007255.1 | AC130456.4 | 0.737 | < 0.001 |
| AC007255.1 | RSPH1 | 0.737 | < 0.001 |
| AC007255.1 | CHDH | 0.737 | < 0.001 |
| AC007255.1 | MIR194-2HG | 0.737 | < 0.001 |
| AC007255.1 | B3GNT3 | 0.736 | < 0.001 |
| AC007255.1 | AC012317.1 | 0.736 | < 0.001 |
| AC007255.1 | TJP3 | 0.736 | < 0.001 |
| AC007255.1 | MYO1A | 0.736 | < 0.001 |
| AC007255.1 | TSPAN3 | 0.735 | < 0.001 |
| AC007255.1 | PDZD3 | 0.735 | < 0.001 |
| AC007255.1 | PLCH1 | 0.735 | < 0.001 |
| AC007255.1 | HNF1A | 0.735 | < 0.001 |
| AC007255.1 | HNF1A-AS1 | 0.734 | < 0.001 |
| AC007255.1 | LINC01978 | 0.733 | < 0.001 |
| AC007255.1 | APOBEC1 | 0.733 | < 0.001 |
| AC007255.1 | AL117382.2 | 0.733 | < 0.001 |
| AC007255.1 | IYD | 0.733 | < 0.001 |
| AC007255.1 | STAMBPL1 | 0.732 | < 0.001 |
| AC007255.1 | SLC37A1 | 0.732 | < 0.001 |
| AC007255.1 | GFPT1 | 0.732 | < 0.001 |
| AC007255.1 | TSPAN12 | 0.731 | < 0.001 |
| AC007255.1 | AC004830.2 | 0.731 | < 0.001 |
| AC007255.1 | MYRF-AS1 | 0.730 | < 0.001 |
| AC007255.1 | CDHR5 | 0.730 | < 0.001 |
| AC007255.1 | R3HDML | 0.730 | < 0.001 |
| AC007255.1 | PPP1R14D | 0.730 | < 0.001 |
| AC007255.1 | HGD | 0.729 | < 0.001 |
| AC007255.1 | AL606489.1 | 0.729 | < 0.001 |
| AC007255.1 | MMP15 | 0.728 | < 0.001 |
| AC007255.1 | MACC1 | 0.728 | < 0.001 |
| AC007255.1 | ERBB3 | 0.728 | < 0.001 |
| AC007255.1 | DTX4 | 0.728 | < 0.001 |
| AC007255.1 | CRYZL2P | 0.728 | < 0.001 |
| AC007255.1 | CLMN | 0.728 | < 0.001 |
| AC007255.1 | MYRF | 0.727 | < 0.001 |
| AC007255.1 | MNX1 | 0.727 | < 0.001 |
| AC007255.1 | GALM | 0.726 | < 0.001 |
| AC007255.1 | SLC26A8 | 0.726 | < 0.001 |
| AC007255.1 | SPIRE2 | 0.725 | < 0.001 |
| AC007255.1 | DDAH1 | 0.725 | < 0.001 |
| AC007255.1 | PRXL2B | 0.725 | < 0.001 |
| AC007255.1 | AGR2 | 0.725 | < 0.001 |
| AC007255.1 | RAB17 | 0.725 | < 0.001 |
| AC007255.1 | TMPRSS2 | 0.725 | < 0.001 |
| AC007255.1 | LINC00543 | 0.724 | < 0.001 |
| AC007255.1 | AC016735.1 | 0.724 | < 0.001 |
| AC007255.1 | NCMAP | 0.724 | < 0.001 |
| AC007255.1 | AVL9 | 0.723 | < 0.001 |
| AC007255.1 | MISP | 0.723 | < 0.001 |
| AC007255.1 | UNC5CL | 0.723 | < 0.001 |
| AC007255.1 | FZD5 | 0.722 | < 0.001 |
| AC007255.1 | GCNT3 | 0.722 | < 0.001 |
| AC007255.1 | HOTTIP | 0.722 | < 0.001 |
| AC007255.1 | AL117382.3 | 0.721 | < 0.001 |
| AC007255.1 | ENPP7P11 | 0.721 | < 0.001 |
| AC007255.1 | OSBPL7 | 0.721 | < 0.001 |
| AC007255.1 | SLC41A2 | 0.721 | < 0.001 |
| AC007255.1 | SLC44A3 | 0.721 | < 0.001 |
| AC007255.1 | RNF103 | 0.720 | < 0.001 |
| AC007255.1 | HNF4G | 0.719 | < 0.001 |
| AC007255.1 | GOLT1A | 0.719 | < 0.001 |
| AC007255.1 | PRR26 | 0.719 | < 0.001 |
| AC007255.1 | PDXDC1 | 0.719 | < 0.001 |
| AC007255.1 | RALGPS1 | 0.718 | < 0.001 |
| AC007255.1 | AC106876.1 | 0.718 | < 0.001 |
| AC007255.1 | GPR35 | 0.717 | < 0.001 |
| AC007255.1 | SLC44A4 | 0.717 | < 0.001 |
| AC007255.1 | AC097639.1 | 0.717 | < 0.001 |
| AC007255.1 | KRT8 | 0.717 | < 0.001 |
| AC007255.1 | BCL2L14 | 0.717 | < 0.001 |
| AC007255.1 | AC007009.1 | 0.715 | < 0.001 |
| AC007255.1 | TSPAN1 | 0.715 | < 0.001 |
| AC007255.1 | HNF4A | 0.714 | < 0.001 |
| AC007255.1 | CAPN8 | 0.714 | < 0.001 |
| AC007255.1 | LINC01979 | 0.714 | < 0.001 |
| AC007255.1 | AC020656.2 | 0.713 | < 0.001 |
| AC007255.1 | B3GNT7 | 0.713 | < 0.001 |
| AC007255.1 | FAM83E | 0.713 | < 0.001 |
| AC007255.1 | FMO5 | 0.712 | < 0.001 |
| AC007255.1 | ADCY6 | 0.712 | < 0.001 |
| AC007255.1 | SELENBP1 | 0.712 | < 0.001 |
| AC007255.1 | TMEM82 | 0.712 | < 0.001 |
| AC007255.1 | SH3BGRL2 | 0.711 | < 0.001 |
| AC007255.1 | AL034550.1 | 0.711 | < 0.001 |
| AC007255.1 | CALML4 | 0.711 | < 0.001 |
| AC007255.1 | LINC002481 | 0.710 | < 0.001 |
| AC007255.1 | LRRIQ4 | 0.710 | < 0.001 |
| AC007255.1 | LRRC75A | 0.710 | < 0.001 |
| AC007255.1 | AC004066.2 | 0.710 | < 0.001 |
| AC007255.1 | CHN2-AS1 | 0.710 | < 0.001 |
| AC007255.1 | LINC01594 | 0.709 | < 0.001 |
| AC007255.1 | AC005920.1 | 0.709 | < 0.001 |
| AC007255.1 | TSPAN8 | 0.709 | < 0.001 |
| AC007255.1 | COBL | 0.709 | < 0.001 |
| AC007255.1 | C2orf15 | 0.709 | < 0.001 |
| AC007255.1 | ARHGAP26 | 0.709 | < 0.001 |
| AC007255.1 | GLOD5 | 0.709 | < 0.001 |
| AC007255.1 | SIGIRR | 0.709 | < 0.001 |
| AC007255.1 | AL391832.2 | 0.708 | < 0.001 |
| AC007255.1 | CMTM8 | 0.708 | < 0.001 |
| AC007255.1 | MAP4K3-DT | 0.708 | < 0.001 |
| AC007255.1 | SLC25A25-AS1 | 0.708 | < 0.001 |
| AC007255.1 | RNF186 | 0.708 | < 0.001 |
| AC007255.1 | AC004982.1 | 0.708 | < 0.001 |
| AC007255.1 | CDX2 | 0.707 | < 0.001 |
| AC007255.1 | ACP6 | 0.707 | < 0.001 |
| AC007255.1 | PLEKHS1 | 0.707 | < 0.001 |
| AC007255.1 | BCAR3-AS1 | 0.707 | < 0.001 |
| AC007255.1 | PGPEP1 | 0.707 | < 0.001 |
| AC007255.1 | TMEM139 | 0.707 | < 0.001 |
| AC007255.1 | ARHGEF38-IT1 | 0.707 | < 0.001 |
| AC007255.1 | ANXA4 | 0.707 | < 0.001 |
| AC007255.1 | SMIM31 | 0.706 | < 0.001 |
| AC007255.1 | LINC02313 | 0.706 | < 0.001 |
| AC007255.1 | AC091588.3 | 0.706 | < 0.001 |
| AC007255.1 | AC091492.1 | 0.706 | < 0.001 |
| AC007255.1 | LINC01843 | 0.705 | < 0.001 |
| AC007255.1 | RASEF | 0.705 | < 0.001 |
| AC007255.1 | BCAS1 | 0.705 | < 0.001 |
| AC007255.1 | GCC2 | 0.705 | < 0.001 |
| AC007255.1 | GOLM1 | 0.705 | < 0.001 |
| AC007255.1 | TBC1D30 | 0.704 | < 0.001 |
| AC007255.1 | TSPAN15 | 0.704 | < 0.001 |
| AC007255.1 | C4orf19 | 0.704 | < 0.001 |
| AC007255.1 | HNF1B | 0.703 | < 0.001 |
| AC007255.1 | AC005037.1 | 0.703 | < 0.001 |
| AC007255.1 | AGR3 | 0.703 | < 0.001 |
| AC007255.1 | GSDMB | 0.703 | < 0.001 |
| AC007255.1 | BNIP5 | 0.702 | < 0.001 |
| AC007255.1 | KCNK5 | 0.702 | < 0.001 |
| AC007255.1 | PDX1 | 0.701 | < 0.001 |
| AC007255.1 | AC129507.3 | 0.701 | < 0.001 |
| AC007255.1 | CHKA | 0.701 | < 0.001 |
| AC007255.1 | PLEKHB1 | 0.701 | < 0.001 |
| AC007255.1 | FUCA1 | 0.700 | < 0.001 |
| AC007255.1 | CFTR | 0.700 | < 0.001 |
| AC007255.1 | AC099521.3 | 0.700 | < 0.001 |
| AC007255.1 | DNAJC22 | 0.700 | < 0.001 |
| AC007255.1 | KCNE3 | 0.699 | < 0.001 |
| AC007255.1 | RUNDC1 | 0.699 | < 0.001 |
| AC007255.1 | MGAT4A | 0.699 | < 0.001 |
| AC007255.1 | C4BPB | 0.699 | < 0.001 |
| AC007255.1 | RAB19 | 0.698 | < 0.001 |
| AC007255.1 | AC254629.1 | 0.698 | < 0.001 |
| AC007255.1 | AC004923.1 | 0.698 | < 0.001 |
| AC007255.1 | SMPDL3B | 0.698 | < 0.001 |
| AC007255.1 | AC090181.3 | 0.697 | < 0.001 |
| AC007255.1 | PRR13 | 0.697 | < 0.001 |
| AC007255.1 | ZNF620 | 0.697 | < 0.001 |
| AC007255.1 | ASL | 0.697 | < 0.001 |
| AC007255.1 | ANKRD40CL | 0.697 | < 0.001 |
| AC007255.1 | ANKS4B | 0.697 | < 0.001 |
| AC007255.1 | RAB11FIP4 | 0.696 | < 0.001 |
| AC007255.1 | TSPAN13 | 0.696 | < 0.001 |
| AC007255.1 | SPATA13 | 0.696 | < 0.001 |
| AC007255.1 | C1GALT1 | 0.696 | < 0.001 |
| AC007255.1 | SLC44A3-AS1 | 0.695 | < 0.001 |
| AC007255.1 | ANKMY2 | 0.694 | < 0.001 |
| AC007255.1 | PIWIL4 | 0.694 | < 0.001 |
| AC007255.1 | GCNT1 | 0.694 | < 0.001 |
| AC007255.1 | ADAP1 | 0.694 | < 0.001 |
| AC007255.1 | AL022322.1 | 0.694 | < 0.001 |
| AC007255.1 | SLC17A4 | 0.694 | < 0.001 |
| AC007255.1 | TM9SF3 | 0.694 | < 0.001 |
| AC007255.1 | RIPK3 | 0.694 | < 0.001 |
| AC007255.1 | PLEKHH1 | 0.694 | < 0.001 |
| AC007255.1 | AC012485.2 | 0.694 | < 0.001 |
| AC007255.1 | ENTPD8 | 0.693 | < 0.001 |
| AC007255.1 | PNKD | 0.693 | < 0.001 |
| AC007255.1 | MUC13 | 0.693 | < 0.001 |
| AC007255.1 | CYP3A5 | 0.693 | < 0.001 |
| AC007255.1 | SSTR1 | 0.693 | < 0.001 |
| AC007255.1 | AL450263.1 | 0.693 | < 0.001 |
| AC007255.1 | METTL7B | 0.693 | < 0.001 |
| AC007255.1 | C2orf72 | 0.692 | < 0.001 |
| AC007255.1 | OMP | 0.692 | < 0.001 |
| AC007255.1 | TTLL6 | 0.692 | < 0.001 |
| AC007255.1 | EPS8L3 | 0.692 | < 0.001 |
| AC007255.1 | RPS6KL1 | 0.692 | < 0.001 |
| AC007255.1 | SMIM24 | 0.692 | < 0.001 |
| AC007255.1 | SMIM22 | 0.692 | < 0.001 |
| AC007255.1 | EPHA10 | 0.692 | < 0.001 |
| AC007255.1 | HHLA2 | 0.692 | < 0.001 |
| AC007255.1 | PARD6B | 0.692 | < 0.001 |
| AC007255.1 | CYSTM1 | 0.692 | < 0.001 |
| AC007255.1 | TM4SF5 | 0.692 | < 0.001 |
| AC007255.1 | NOSTRIN | 0.691 | < 0.001 |
| AC007255.1 | MOGAT3 | 0.691 | < 0.001 |
| AC007255.1 | KRT18P15 | 0.691 | < 0.001 |
| AC007255.1 | GORASP1 | 0.691 | < 0.001 |
| AC007255.1 | RN7SL541P | 0.690 | < 0.001 |
| AC007255.1 | SLC39A5 | 0.690 | < 0.001 |
| AC007255.1 | CFAP65 | 0.690 | < 0.001 |
| AC007255.1 | AC116552.1 | 0.689 | < 0.001 |
| AC007255.1 | AC129507.2 | 0.689 | < 0.001 |
| AC007255.1 | AC092868.2 | 0.688 | < 0.001 |
| AC007255.1 | ELF3 | 0.688 | < 0.001 |
| AC007255.1 | CACFD1 | 0.688 | < 0.001 |
| AC007255.1 | SPACA4 | 0.688 | < 0.001 |
| AC007255.1 | FAM47E | 0.688 | < 0.001 |
| AC007255.1 | SULT1C2 | 0.686 | < 0.001 |
| AC007255.1 | ADCY6-DT | 0.686 | < 0.001 |
| AC007255.1 | ACSM3 | 0.686 | < 0.001 |
| AC007255.1 | CEMIP2 | 0.686 | < 0.001 |
| AC007255.1 | VILL | 0.686 | < 0.001 |
| AC007255.1 | RBM47 | 0.686 | < 0.001 |
| AC007255.1 | EVX1 | 0.685 | < 0.001 |
| AC007255.1 | C1orf115 | 0.685 | < 0.001 |
| AC007255.1 | AL121895.1 | 0.685 | < 0.001 |
| AC007255.1 | CA13 | 0.685 | < 0.001 |
| AC007255.1 | TTC38 | 0.685 | < 0.001 |
| AC007255.1 | SMIM6 | 0.685 | < 0.001 |
| AC007255.1 | SLC3A1 | 0.684 | < 0.001 |
| AC007255.1 | GAL3ST1 | 0.684 | < 0.001 |
| AC007255.1 | KAZALD1 | 0.684 | < 0.001 |
| AC007255.1 | AC008147.1 | 0.683 | < 0.001 |
| AC007255.1 | CAPN5 | 0.683 | < 0.001 |
| AC007255.1 | AC009065.3 | 0.683 | < 0.001 |
| AC007255.1 | VIL1 | 0.683 | < 0.001 |
| AC007255.1 | LINC00628 | 0.683 | < 0.001 |
| AC007255.1 | AC097493.2 | 0.683 | < 0.001 |
| AC007255.1 | FER1L4 | 0.683 | < 0.001 |
| AC007255.1 | ZSCAN16 | 0.683 | < 0.001 |
| AC007255.1 | DEPTOR | 0.683 | < 0.001 |
| AC007255.1 | AC114488.1 | 0.683 | < 0.001 |
| AC007255.1 | SMAD6 | 0.682 | < 0.001 |
| AC007255.1 | SLC6A20 | 0.682 | < 0.001 |
| AC007255.1 | EXOC6 | 0.682 | < 0.001 |
| AC007255.1 | PHACTR2-AS1 | 0.682 | < 0.001 |
| AC007255.1 | AL591895.1 | 0.682 | < 0.001 |
| AC007255.1 | MUC3A | 0.682 | < 0.001 |
| AC007255.1 | FAM81A | 0.681 | < 0.001 |
| AC007255.1 | AL359075.1 | 0.681 | < 0.001 |
| AC007255.1 | AC008040.2 | 0.681 | < 0.001 |
| AC007255.1 | NOL4L | 0.681 | < 0.001 |
| AC007255.1 | FMN1 | 0.681 | < 0.001 |
| AC007255.1 | BTNL8 | 0.681 | < 0.001 |
| AC007255.1 | LINC01091 | 0.681 | < 0.001 |
| AC007255.1 | SPDEF | 0.680 | < 0.001 |
| AC007255.1 | SGK2 | 0.680 | < 0.001 |
| AC007255.1 | SPATA6L | 0.680 | < 0.001 |
| AC007255.1 | LGALS4 | 0.680 | < 0.001 |
| AC007255.1 | AC090181.2 | 0.680 | < 0.001 |
| AC007255.1 | AC008521.1 | 0.680 | < 0.001 |
| AC007255.1 | IL17RB | 0.680 | < 0.001 |
| AC007255.1 | TRIM10 | 0.680 | < 0.001 |
| AC007255.1 | PPP1R1B | 0.680 | < 0.001 |
| AC007255.1 | ABCC6P1 | 0.680 | < 0.001 |
| AC007255.1 | MGAM2 | 0.679 | < 0.001 |
| AC007255.1 | SEMA4G | 0.679 | < 0.001 |
| AC007255.1 | POC1B | 0.679 | < 0.001 |
| AC007255.1 | ALDOB | 0.679 | < 0.001 |
| AC007255.1 | OCLN | 0.679 | < 0.001 |
| AC007255.1 | MS4A8 | 0.679 | < 0.001 |
| AC007255.1 | NANOGP4 | 0.679 | < 0.001 |
| AC007255.1 | AC127521.1 | 0.679 | < 0.001 |
| AC007255.1 | CCNI2 | 0.679 | < 0.001 |
| AC007255.1 | ACY3 | 0.679 | < 0.001 |
| AC007255.1 | AC073957.3 | 0.678 | < 0.001 |
| AC007255.1 | TRBV29OR9-2 | 0.678 | < 0.001 |
| AC007255.1 | KRTAP5-AS1 | 0.677 | < 0.001 |
| AC007255.1 | AC009065.6 | 0.677 | < 0.001 |
| AC007255.1 | GPRC5A | 0.677 | < 0.001 |
| AC007255.1 | SFMBT1 | 0.677 | < 0.001 |
| AC007255.1 | SLC29A2 | 0.677 | < 0.001 |
| AC007255.1 | KCNJ3 | 0.676 | < 0.001 |
| AC007255.1 | GPD1 | 0.676 | < 0.001 |
| AC007255.1 | ENPP4 | 0.676 | < 0.001 |
| AC007255.1 | AC079775.1 | 0.676 | < 0.001 |
| AC007255.1 | MIR3646 | 0.676 | < 0.001 |
| AC007255.1 | PTPRH | 0.675 | < 0.001 |
| AC007255.1 | AL078459.1 | 0.675 | < 0.001 |
| AC007255.1 | PAFAH2 | 0.675 | < 0.001 |
| AC007255.1 | HDHD3 | 0.675 | < 0.001 |
| AC007255.1 | CLDN23 | 0.675 | < 0.001 |
| AC007255.1 | MFSD9 | 0.675 | < 0.001 |
| AC007255.1 | REPS2 | 0.675 | < 0.001 |
| AC007255.1 | ABHD11 | 0.675 | < 0.001 |
| AC007255.1 | RAD17P1 | 0.675 | < 0.001 |
| AC007255.1 | TMPRSS3 | 0.674 | < 0.001 |
| AC007255.1 | NEK3 | 0.674 | < 0.001 |
| AC007255.1 | SYBU | 0.674 | < 0.001 |
| AC007255.1 | TRIM15 | 0.674 | < 0.001 |
| AC007255.1 | DDC | 0.674 | < 0.001 |
| AC007255.1 | ALPI | 0.674 | < 0.001 |
| AC007255.1 | ALG1L5P | 0.674 | < 0.001 |
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| AC007255.1 | SOWAHA | 0.673 | < 0.001 |
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| AC007255.1 | TMEM53 | 0.669 | < 0.001 |
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| AC007255.1 | AL133410.1 | 0.668 | < 0.001 |
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| AC007255.1 | GALNT4 | 0.668 | < 0.001 |
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| AC007255.1 | CLDN7 | 0.623 | < 0.001 |
| AC007255.1 | CDHR2 | 0.623 | < 0.001 |
| AC007255.1 | NGLY1 | 0.623 | < 0.001 |
| AC007255.1 | ANXA13 | 0.623 | < 0.001 |
| AC007255.1 | MSH5-SAPCD1 | 0.623 | < 0.001 |
| AC007255.1 | TC2N | 0.622 | < 0.001 |
| AC007255.1 | SPINK4 | 0.622 | < 0.001 |
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| AC007255.1 | GALNT7 | 0.622 | < 0.001 |
| AC007255.1 | ASS1 | 0.622 | < 0.001 |
| AC007255.1 | CCL15-CCL14 | 0.622 | < 0.001 |
| AC007255.1 | SNORC | 0.622 | < 0.001 |
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| AC007255.1 | RGL3 | 0.621 | < 0.001 |
| AC007255.1 | GUCY2C | 0.621 | < 0.001 |
| AC007255.1 | AL031320.2 | 0.621 | < 0.001 |
| AC007255.1 | BEND7 | 0.621 | < 0.001 |
| AC007255.1 | AOC1 | 0.621 | < 0.001 |
| AC007255.1 | SLC25A20 | 0.621 | < 0.001 |
| AC007255.1 | RRBP1 | 0.621 | < 0.001 |
| AC007255.1 | HES6 | 0.621 | < 0.001 |
| AC007255.1 | PGAP6 | 0.621 | < 0.001 |
| AC007255.1 | CASP10 | 0.621 | < 0.001 |
| AC007255.1 | SNRPCP5 | 0.621 | < 0.001 |
| AC007255.1 | GMDS | 0.620 | < 0.001 |
| AC007255.1 | KIF13B | 0.620 | < 0.001 |
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| AC007255.1 | CHMP2B | 0.620 | < 0.001 |
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| AC007255.1 | C6orf99 | 0.619 | < 0.001 |
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| AC007255.1 | PKDCC | 0.619 | < 0.001 |
| AC007255.1 | ZG16B | 0.619 | < 0.001 |
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| AC007255.1 | KRT18P11 | 0.618 | < 0.001 |
| AC007255.1 | KCNJ11 | 0.617 | < 0.001 |
| AC007255.1 | KRT8P45 | 0.617 | < 0.001 |
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| AC007255.1 | CMC1 | 0.616 | < 0.001 |
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| AC007255.1 | SPINK1 | 0.614 | < 0.001 |
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| AC007255.1 | AC007637.1 | 0.614 | < 0.001 |
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| AC007255.1 | IFT20 | 0.585 | < 0.001 |
| AC007255.1 | MYH14 | 0.585 | < 0.001 |
| AC007255.1 | LYZ | 0.584 | < 0.001 |
| AC007255.1 | AC090505.1 | 0.584 | < 0.001 |
| AC007255.1 | AC004594.1 | 0.584 | < 0.001 |
| AC007255.1 | SLC17A9 | 0.584 | < 0.001 |
| AC007255.1 | AL513174.1 | 0.584 | < 0.001 |
| AC007255.1 | AC087588.2 | 0.584 | < 0.001 |
| AC007255.1 | SPRY1 | 0.584 | < 0.001 |
| AC007255.1 | PRELID3B | 0.584 | < 0.001 |
| AC007255.1 | NPIPB14P | 0.583 | < 0.001 |
| AC007255.1 | PNRC2 | 0.583 | < 0.001 |
| AC007255.1 | CAMK2N1 | 0.583 | < 0.001 |
| AC007255.1 | CT83 | 0.583 | < 0.001 |
| AC007255.1 | AC137630.1 | 0.583 | < 0.001 |
| AC007255.1 | LINC02535 | 0.583 | < 0.001 |
| AC007255.1 | AGT | 0.583 | < 0.001 |
| AC007255.1 | MAGIX | 0.583 | < 0.001 |
| AC007255.1 | CCDC198 | 0.583 | < 0.001 |
| AC007255.1 | AC010487.1 | 0.583 | < 0.001 |
| AC007255.1 | DYRK2 | 0.583 | < 0.001 |
| AC007255.1 | PTMAP3 | 0.583 | < 0.001 |
| AC007255.1 | SEPHS2 | 0.583 | < 0.001 |
| AC007255.1 | LINC02453 | 0.582 | < 0.001 |
| AC007255.1 | KCTD14 | 0.582 | < 0.001 |
| AC007255.1 | CALR4P | 0.582 | < 0.001 |
| AC007255.1 | CEACAM18 | 0.582 | < 0.001 |
| AC007255.1 | NSUN7 | 0.582 | < 0.001 |
| AC007255.1 | EIF3J-DT | 0.582 | < 0.001 |
| AC007255.1 | AC005041.2 | 0.582 | < 0.001 |
| AC007255.1 | AC010976.2 | 0.582 | < 0.001 |
| AC007255.1 | RNF125 | 0.581 | < 0.001 |
| AC007255.1 | AC091544.5 | 0.581 | < 0.001 |
| AC007255.1 | LRATD2 | 0.581 | < 0.001 |
| AC007255.1 | GYG2 | 0.581 | < 0.001 |
| AC007255.1 | BAIAP2L2 | 0.581 | < 0.001 |
| AC007255.1 | SLC9A8 | 0.581 | < 0.001 |
| AC007255.1 | GAU1 | 0.581 | < 0.001 |
| AC007255.1 | LGALS3 | 0.581 | < 0.001 |
| AC007255.1 | ZMIZ2 | 0.581 | < 0.001 |
| AC007255.1 | MICU1 | 0.581 | < 0.001 |
| AC007255.1 | MIR4728 | 0.581 | < 0.001 |
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| AC007255.1 | TMC7 | 0.580 | < 0.001 |
| AC007255.1 | SLC18B1 | 0.580 | < 0.001 |
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| AC007255.1 | PLLP | 0.580 | < 0.001 |
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| AC007255.1 | AZGP1 | 0.579 | < 0.001 |
| AC007255.1 | AL590438.1 | 0.579 | < 0.001 |
| AC007255.1 | SLC43A1 | 0.579 | < 0.001 |
| AC007255.1 | C4BPA | 0.579 | < 0.001 |
| AC007255.1 | TMEM229A | 0.579 | < 0.001 |
| AC007255.1 | KRT7 | 0.579 | < 0.001 |
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| AC007255.1 | NEK6 | 0.579 | < 0.001 |
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| AC007255.1 | NEK8 | 0.577 | < 0.001 |
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| AC007255.1 | ZBED3 | 0.577 | < 0.001 |
| AC007255.1 | GMPPB | 0.577 | < 0.001 |
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| AC007255.1 | AC012354.3 | 0.577 | < 0.001 |
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| AC007255.1 | CCT8P1 | 0.577 | < 0.001 |
| AC007255.1 | SLC6A19 | 0.577 | < 0.001 |
| AC007255.1 | ZNF75D | 0.577 | < 0.001 |
| AC007255.1 | AC007272.1 | 0.577 | < 0.001 |
| AC007255.1 | AL355353.2 | 0.577 | < 0.001 |
| AC007255.1 | SUSD1 | 0.576 | < 0.001 |
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| AC007255.1 | AL691482.2 | 0.576 | < 0.001 |
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| AC007255.1 | MAN2A2 | 0.576 | < 0.001 |
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| AC007255.1 | RPIA | 0.575 | < 0.001 |
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| AC007255.1 | IL9RP3 | 0.575 | < 0.001 |
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| AC007255.1 | CA8 | 0.575 | < 0.001 |
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| AC007255.1 | ARHGAP44 | 0.575 | < 0.001 |
| AC007255.1 | KRT8P20 | 0.575 | < 0.001 |
| AC007255.1 | GATA4 | 0.575 | < 0.001 |
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| AC007255.1 | PLEKHA7 | 0.574 | < 0.001 |
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| AC007255.1 | KRT18P4 | 0.573 | < 0.001 |
| AC007255.1 | LHFPL3 | 0.573 | < 0.001 |
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| AC007255.1 | ARHGAP42 | 0.572 | < 0.001 |
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| AC007255.1 | BSPRY | 0.569 | < 0.001 |
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| AC007255.1 | RNA5SP311 | 0.569 | < 0.001 |
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| AC007255.1 | ERBB2 | 0.569 | < 0.001 |
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| AC007255.1 | PDE11A | 0.562 | < 0.001 |
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| AC007255.1 | RNU6-702P | 0.562 | < 0.001 |
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| AC007255.1 | SELENOS | 0.462 | < 0.001 |
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| AC007255.1 | MT-CO1 | 0.462 | < 0.001 |
| AC007255.1 | STEAP2 | 0.462 | < 0.001 |
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| AC007255.1 | MSLNL | 0.461 | < 0.001 |
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| AC007255.1 | REG3G | 0.460 | < 0.001 |
| AC007255.1 | KRT8P39 | 0.460 | < 0.001 |
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| AC007255.1 | HOXB8 | 0.443 | < 0.001 |
| AC007255.1 | TNFRSF10D | 0.443 | < 0.001 |
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| AC007255.1 | SLC7A9 | 0.428 | < 0.001 |
| AC007255.1 | TPM3P6 | 0.428 | < 0.001 |
| AC007255.1 | LINC02649 | 0.428 | < 0.001 |
| AC007255.1 | CYP4F2 | 0.428 | < 0.001 |
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| AC007255.1 | CWC25 | 0.427 | < 0.001 |
| AC007255.1 | NFKBIE | 0.427 | < 0.001 |
| AC007255.1 | AP001893.1 | 0.427 | < 0.001 |
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| AC007255.1 | AL365205.1 | 0.427 | < 0.001 |
| AC007255.1 | SNORA15B-1 | 0.427 | < 0.001 |
| AC007255.1 | IL13RA1 | 0.427 | < 0.001 |
| AC007255.1 | AL590822.2 | 0.427 | < 0.001 |
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| AC007255.1 | ATXN7 | 0.427 | < 0.001 |
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| AC007255.1 | MTTP | 0.427 | < 0.001 |
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| AC007255.1 | OR2I1P | 0.426 | < 0.001 |
| AC007255.1 | CFL1P5 | 0.426 | < 0.001 |
| AC007255.1 | MGLL | 0.426 | < 0.001 |
| AC007255.1 | LSMEM2 | 0.426 | < 0.001 |
| AC007255.1 | LINC01273 | 0.426 | < 0.001 |
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| AC007255.1 | LINC02418 | 0.426 | < 0.001 |
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| AC007255.1 | TIMP4 | 0.426 | < 0.001 |
| AC007255.1 | MAGI1-IT1 | 0.426 | < 0.001 |
| AC007255.1 | DOP1A | 0.426 | < 0.001 |
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| AC007255.1 | AC139887.2 | 0.423 | < 0.001 |
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| AC007255.1 | FAM199X | 0.419 | < 0.001 |
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| AC007255.1 | PLEKHA8 | 0.418 | < 0.001 |
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