



- DR3\_Q4
- STTTCRNNTT\_IRF\_Q6
- GATA\_Q6
- FREAC4\_Q1
- SYNCRIP\_TARGET\_GENES
- EVI1\_Q2
- GCCNNNWTAAAR\_UNKNOWN
- PR\_Q1
- HFH3\_Q1
- TAXCREB\_Q1
- AAANWWTGC\_UNKNOWN
- RORA2\_Q1
- STAT3\_Q2
- YTAATTAA\_LHX3\_Q1
- HEY2\_TARGET\_GENES
- WWTAAAGGC\_UNKNOWN
- YTAAYNGCT\_UNKNOWN
- TSHZ1\_TARGET\_GENES
- PR\_Q2
- TAAWWATAG\_RSRFC4\_Q2
- GGGNNTTCC\_NFKB\_Q6\_Q1
- CDP\_Q2
- OCT1\_Q7
- HMEF2\_Q6
- CGGAARNGGCNG\_UNKNOWN
- ZBTB1\_TARGET\_GENES
- CCGNMNTNACG\_UNKNOWN
- NR1D1\_TARGET\_GENES
- TFCP2\_TARGET\_GENES
- ZNF584\_TARGET\_GENES
- PPARA\_TARGET\_GENES
- SIPA1\_TARGET\_GENES
- KDM1B\_TARGET\_GENES
- ATGGYGA\_UNKNOWN
- ZNF133\_TARGET\_GENES
- TAANNYSGCG\_UNKNOWN
- MCAATNNNNNGCG\_UNKNOWN
- ZNF585B\_TARGET\_GENES
- ZNF37A\_TARGET\_GENES
- CGTSACG\_PAX3\_B
- CREBP1\_Q1
- CCCNNNNNNAAGWT\_UNKNOWN
- CDH4\_TARGET\_GENES
- EGR2\_Q1
- EGR3\_Q1
- TERF2\_TARGET\_GENES
- ZNF165\_TARGET\_GENES
- GTF3A\_TARGET\_GENES
- THRAP3\_TARGET\_GENES
- TMTCGCGANR\_UNKNOWN
- YBX3\_TARGET\_GENES
- RAX2\_TARGET\_GENES
- ZNF667\_TARGET\_GENES
- GKCGCNNNNNNTGAYG\_UNKNOWN
- GCGNNANTTCC\_UNKNOWN
- ZNF623\_TARGET\_GENES
- GGCNKCCATNK\_UNKNOWN
- ZNF677\_TARGET\_GENES
- MCM3\_TARGET\_GENES
- HOXA7\_TARGET\_GENES
- RUVBL1\_TARGET\_GENES
- ARHGAP35\_TARGET\_GENES
- AAGWWRNYGGC\_UNKNOWN
- IRX2\_TARGET\_GENES
- ZBTB5\_TARGET\_GENES
- HOXC13\_TARGET\_GENES
- ZMYM2\_TARGET\_GENES
- CHAF1A\_TARGET\_GENES
- ZSCAN5C\_TARGET\_GENES
- ZNF622\_TARGET\_GENES
- PCGF6\_TARGET\_GENES
- PPARGC1A\_TARGET\_GENES
- ZNF704\_TARGET\_GENES
- MCM2\_TARGET\_GENES
- E2F\_Q2
- MAPK3\_TARGET\_GENES
- ETF\_Q6
- PAX8\_TARGET\_GENES
- WCTCNATGGY\_UNKNOWN
- THRA\_TARGET\_GENES
- TOX4\_TARGET\_GENES
- ACTWSNACTNY\_UNKNOWN
- SREBP1\_Q1
- NCX\_Q1
- KRCTCNNNNMANAGC\_UNKNOWN
- YYCATTCWW\_UNKNOWN
- RBM15\_TARGET\_GENES
- TTTNNANAGCYR\_UNKNOWN
- PAX5\_Q1
- PER1\_TARGET\_GENES
- ZNF777\_TARGET\_GENES
- RBL1\_TARGET\_GENES
- HDAC8\_TARGET\_GENES
- STAT1\_Q1
- MCRS1\_TARGET\_GENES
- SQSTM1\_TARGET\_GENES
- SETBP1\_TARGET\_GENES
- ZNF532\_TARGET\_GENES
- GGAMTNNNNNTCCY\_UNKNOWN
- METHYL CYTOSINE\_DIOXYGENASE\_TET\_UNIPROT\_A0A023HHK9
- SGCGSSAAA\_E2F1DP2\_Q1
- ZBED4\_TARGET\_GENES
- MXD1\_TARGET\_GENES
- RRCCGTTA\_UNKNOWN
- GGAANCGGAANY\_UNKNOWN
- GCGSCMNTTT\_UNKNOWN
- GGCNNMSMYNTTG\_UNKNOWN
- CREB3L2\_TARGET\_GENES
- HOXD11\_TARGET\_GENES
- ZNF669\_TARGET\_GENES
- MYC\_Q2
- TNCATNTCCYR\_UNKNOWN
- AACWWCAANK\_UNKNOWN
- TTCNRGNNTTC\_HSF\_Q6
- GGCNRNWCTTYS\_UNKNOWN
- SNACANNNYSYAGA\_UNKNOWN
- GATGKMRGCG\_UNKNOWN
- ZNF595\_TARGET\_GENES
- DNMT1\_TARGET\_GENES
- AACYNNTTCCS\_UNKNOWN
- SOX11\_TARGET\_GENES
- ZZZ3\_TARGET\_GENES
- KCCGNSWTTT\_UNKNOWN
- COBLL1\_TARGET\_GENES
- ZNF547\_TARGET\_GENES
- RACTNNRTTNC\_UNKNOWN
- TBX1\_TARGET\_GENES
- MYBL1\_TARGET\_GENES
- MIF1\_Q1
- FOXQ1\_TARGET\_GENES
- CCAWYNNGAAR\_UNKNOWN
- E2F\_Q1
- NUFIP1\_TARGET\_GENES
- DICER1\_TARGET\_GENES