





SCGGAAGY_V\$ELK1_02
V\$MYOGENIN_Q6
RGAANNTTC_V\$HSF1_01
V\$IK2_01
WGGAATGY_V\$TEF1_Q6
V\$E12_Q6
CTGCAGY_UNKNOWN
RYTTCCTG_V\$ETS2_B
V\$E2F_Q6
RNGTG GGC_UNKNOWN
V\$E2F1_Q4_01
CAGGTA_V\$AREB6_01
CAGGTG_V\$E12_Q6
V\$E2F_Q4
V\$E2F_Q3_01
TGACAGNY_V\$MEIS1_01
GGGTG GRR_V\$PAX4_03
TTANTCA_UNKNOWN
CAGCTG_V\$AP4_Q5
WTGAAAT_UNKNOWN
AACTTT_UNKNOWN
RTAAACA_V\$FREAC2_01
TGANTCA_V\$AP1_C
GGGAGGRR_V\$MAZ_Q6
YTATTTTNR_V\$MEF2_02
TTGTTT_V\$FOXO4_01
TGTTTGY_V\$HNF3_Q6
CTTTGA_V\$LEF1_Q2
GCANCTGNY_V\$MYOD_Q6
TGGAAA_V\$NFAT_Q4_01