





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