



GATA_Q6

AR_Q2

TGASTMAGC_NFE2_01

CCGNMNNTNACG_UNKNOWN

SREBP1_01

AR_03

GGCNRNWCTTYS_UNKNOWN

CRGAARNNNNCGA_UNKNOWN

MEF2_01

HMEF2_Q6

NRSF_01

CAGNWMCNNGAC_UNKNOWN

TMTCGCGANR_UNKNOWN

GGAANCGGAANY_UNKNOWN

GGCNKCCATNK_UNKNOWN

AAGWWRNYGGC_UNKNOWN

TAANNYSGCG_UNKNOWN

GCGNNANTTCC_UNKNOWN

KCCGNSWTTT_UNKNOWN

GKCGCNNNNNNNTGAYG_UNKNOWN

GCGSCMNTTT_UNKNOWN

RRCCGTTA_UNKNOWN

ACAWNRRNSRCGG_UNKNOWN

AACYNNNNNTTCCS_UNKNOWN

SGCGSSAAA_E2F1DP2_01

KRCTCNNNNMANAGC_UNKNOWN

TTTNANAGCYR_UNKNOWN

AGCYRW TTC_UNKNOWN

YNTTTNNNANGCARM_UNKNOWN