

jaspar_MA0035.3 (Gata1) (RC)



cisbp_M1867 (grn)



cisbp_M6253 (grn)



jaspar_MA0036.2 (GATA2) (RC)



bergman_pnr (pnr)



homer_ACTGATAAGA_PQM-1 (PQM-1)



cisbp_M4600 (grn)



cisbp_M4622 (grn)



cisbp_M4595 (grn)



cisbp_M1868 (GATA2)



homer_NBWGATAAGR_Gata4 (Gata4)



cisbp_M3320 (GATA2)



cisbp_M1336 (M1336)



cisbp_M0795 (GLN3)



homer_AWTGATAAGA_elt-3 (ELT-3)



cisbp_M4714 (grn)



homer_NCCTTATCTC_Gata2 (Gata2) (RC)



homer_CAGATAAGGN_Gata1 (Gata1)



cisbp_M3315 (grn)



cisbp_M3316 (grn)



jaspar_MA0037.2 (GATA3)



cisbp_M1869 (grn)



flyfactorsurvey_pnr_SANGER_5_FBgn0003117 (pnr)



cisbp_M5156 (grn)



jaspar_MA0482.1 (Gata4) (RC)



cisbp_M2284 (grn)



flyfactorsurvey_GATAe_SANGER_5_FBgn0038391 (GATAe)



cisbp_M4983 (GATAe)



flyfactorsurvey_srp_SANGER_5_FBgn0003507 (srp) (RC)



cisbp_M5214 (srp)



stark_ANHDBHGATAASSDNNB (pnr)



cisbp_M4320 (GZF3)



stark_YGATAAGC (pnr)



jaspar_MA0542.1 (ELT-3) (RC)



cisbp_M2335 (M2335)



cisbp_M4715 (M4715)



cisbp_M0001 (ABF1) (RC)



bergman_TFAM (TFAM) (RC)



stark_TTATS (mitochondrial transcription factor A) (RC)



flyfactorsurvey_GATAAd_SANGER_5_FBgn0032223 (GATAAd) (RC)



cisbp_M4982 (GATAAd)



homer_GCTGATAASV_Unknown5 (Unknown5)



cisbp_M2575 (TFAM) (RC)



cisbp_M0772 (M0772)



position