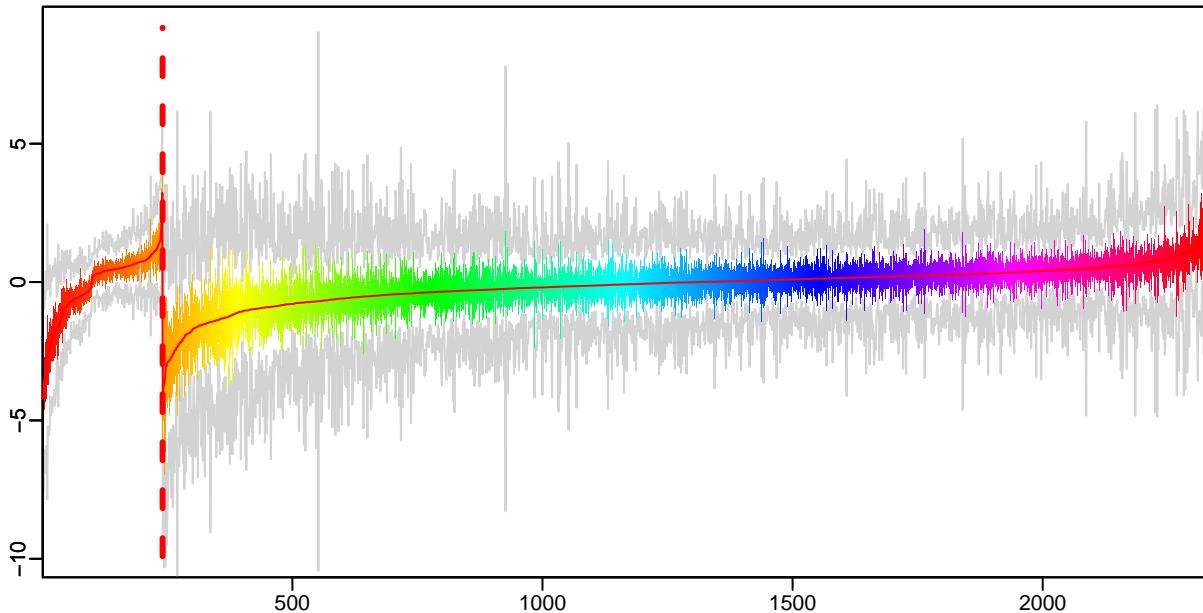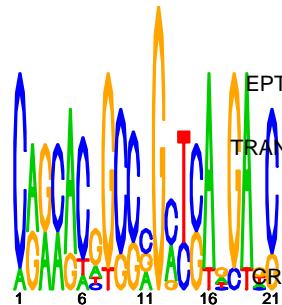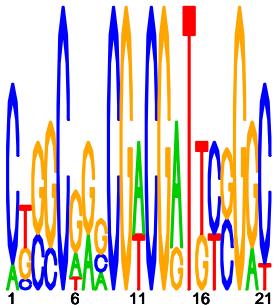# Cluster: 0212 mtu; resid: 0.55; r/c: 18/240

13 Oct 25 14:25:18 iter=2000
cMonkey Version 4.9.8 mtu

upstream meme PSSM #1; e=2.5e−06   upstream meme PSSM #2; e=0.00031

log10(P) upstream meme   log10(P.clust)=−6.37; 18 seqs; 13 uniq

PROBABLE CONSERVED INTEGRAL MEMBRANE PROTEIN: RV2994
EPTA−2,4−DIENE−1,7−DIOATE ISOMERASE (HHDD ISOMERASE): RV2993C
−11.75   glutamyl−tRNA synthetase: RV2992C
TRANSMEMBRANE CYTOCHROME C OXIDASE (SUBUNIT II) CTAC: RV2200C
−10.78   Possible conserved integral membrane protein: RV2199C
−10.78   Probable asparagine synthetase AsnB: RV2201
−9.71   POSSIBLE CONSERVED LIPOPROTEIN LPPH: RV3576
−8.09   TRANSCRIPTIONAL REGULATORY PROTEIN (PROBABLY LACI−FAMILY): RV3575C
−8.09   lysyl−tRNA synthetase: RV3598C
−6.95   POSSIBLE CONSERVED TRANSMEMBRANE PROTEIN: RV0879C
AMINE TRANSPORT TRANSMEMBRANE PROTEIN ABC TRANSPORTER: RV2563
NSERVED HYPOTHETICAL ALANINE AND LEUCINE RICH PROTEIN: RV2567
−2.46
−2.21   hypothetical protein: RV2231C
−2.21   hypothetical protein: RV2230C
−2.21   hypothetical protein: RV2229C
−0.22   hypothetical protein: RV3234C
−0.22   hypothetical protein: RV3233C
−0.06   cystathionine gamma−synthase: RV1079