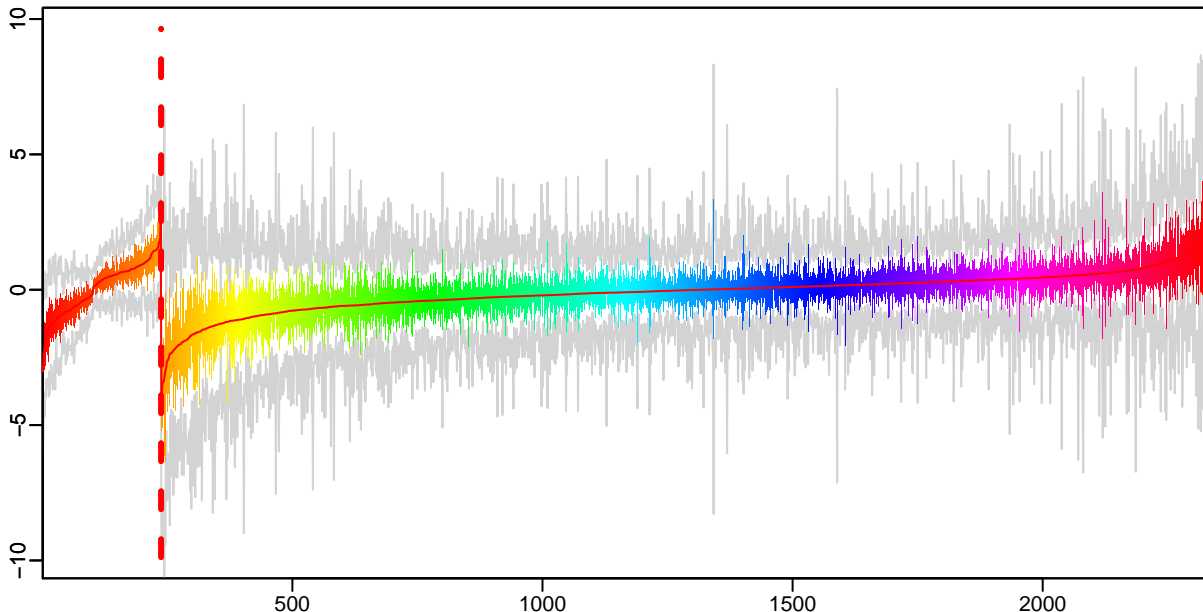
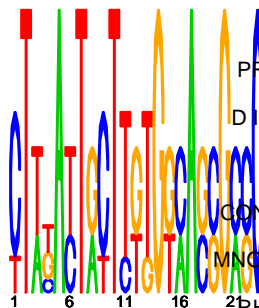
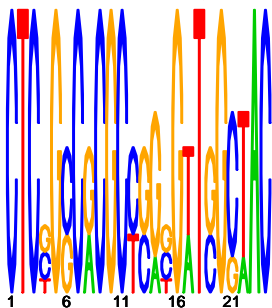


Cluster: 0455 mtu; resid: 0.53; r/c: 12/237

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



upstream meme PSSM #1; e=0.00016 upstream meme PSSM #2; e=0.00028



log10(P) upstream meme log10(P.clust)=-10.90; 12 seqs; 7 uniq

PROBABLE SHORT-CHAIN TYPE DEHYDROGENASE/REDUCTASE: RV2509

D INTEGRAL MEMBRANE LEUCINE AND ALANINE RICH PROTEIN: RV2508C

-15.36 hypothetical protein: RV3268

CONSERVED HYPOTHETICAL PROTEIN (CPSA-RELATED PROTEIN): RV3267

MNOSE BIOSYNTHESIS PROTEIN) (TDP-RHAMNOSE SYNTHASE): RV3266C

2 RHAMNOSE 3(4->3)-ALPHA-D-GlcNAc(1->6)-P-DECAPRENYL: RV3265C

(D-ALPHA-D-5-PYRROSE-1-PHOSPHATE GUANYLYLTRANSFERASE): RV3264C

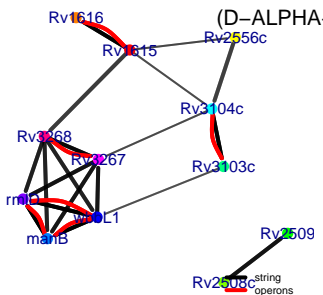
-6.90 hypothetical protein: RV2556C

-6.22 hypothetical protein: RV1616

-6.22 Probable hypothetical membrane protein: RV1615

-0.00 POSSIBLE CONSERVED TRANSMEMBRANE PROTEIN: RV3104C

-0.03 HYPOTHETICAL PROLINE-RICH PROTEIN: RV3103C



-200 -100 -1