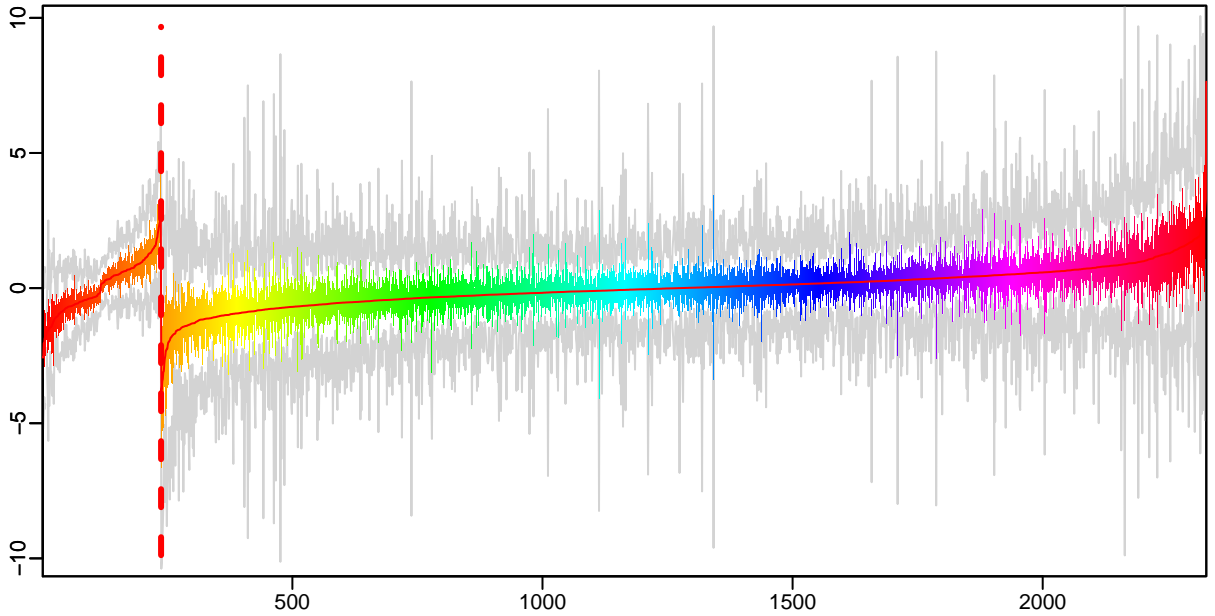
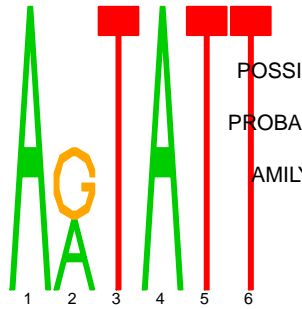
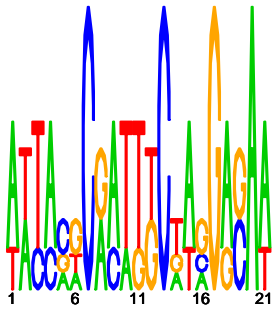


Cluster: 0435 mtu; resid: 0.53; r/c: 11/237

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



upstream meme PSSM #1; e=2e+03 upstream meme PSSM #2; e=7.7e+03



log10(P) upstream meme log10(P.clust)=-5.68; 11 seqs; 5 uniq

POSSIBLE CONSERVED MCE ASSOCIATED MEMBRANE PROTEIN: RV1973

PROBABLE CONSERVED MCE ASSOCIATED MEMBRANE PROTEIN: RV1972

FAMILY LIPOPROTEIN LPRM (MCE-FAMILY LIPOPROTEIN MCE3E): RV1970

-6.87 MCE-FAMILY PROTEIN MCE3C: RV1968

-6.87 MCE-FAMILY PROTEIN MCE3B: RV1967

NSERVED HYPOTHETICAL INTEGRAL MEMBRANE PROTEIN YRBE3B: RV1965

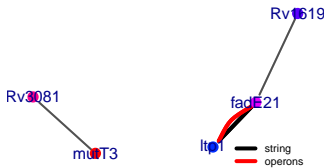
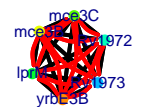
SPHATASE (820XO-DGTPASE) (DGTP PYROPHOSPHOHYDROLASE): RV0413

-5.94 hypothetical protein: RV3081

-4.22 PROBABLE ACYL-CoA DEHYDROGENASE FADE21: RV2789C

-4.22 lipid-transfer protein: RV2790C

-0.71 hypothetical protein: RV1619



-200 -100 -1