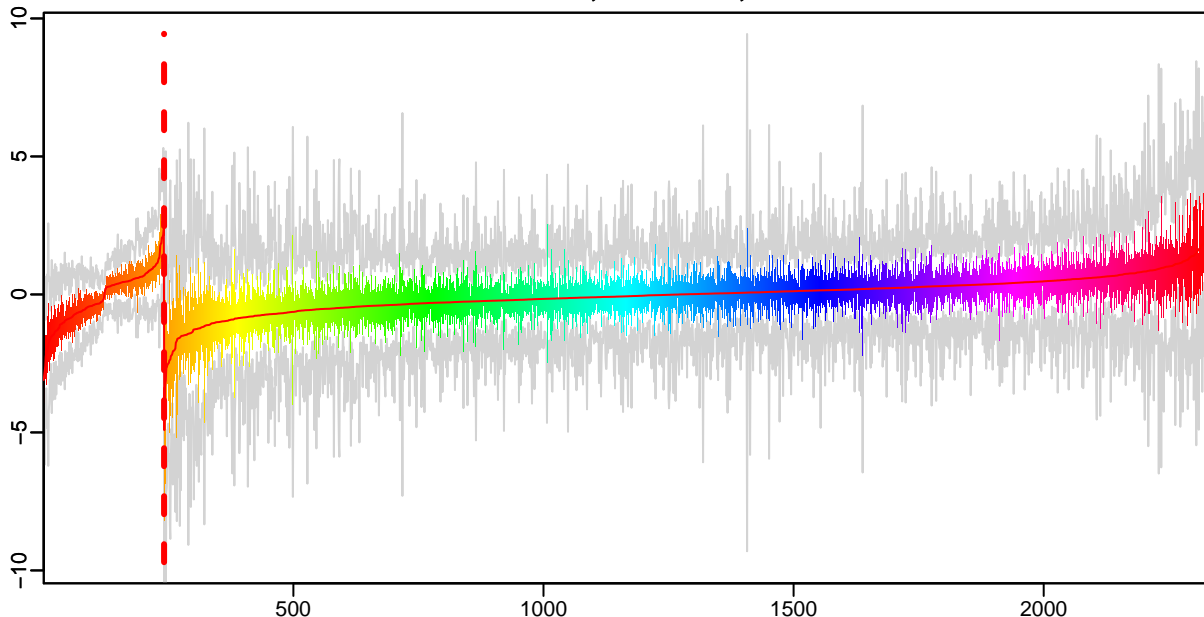
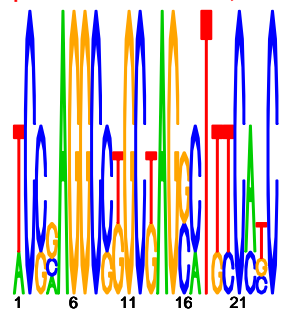


Cluster: 0379 mtu; resid: 0.53; r/c: 10/241

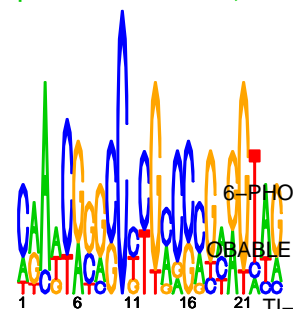
13 Oct 25 14:25:18 iter=2000  
cMonkey Version 4.9.8 mtu



upstream meme PSSM #1; e=8.6



upstream meme PSSM #2; e=9.1



log10(P) upstream meme log10(P.clust)=-8.51; 10 seqs; 9 uniq

-17.20 PROBABLE CONSERVED TRANSMEMBRANE PROTEIN: RV2206

-17.20 hypothetical protein: RV2205C

6-PHOSPHATE DEACETYLASE NAGA (GLCNAC 6-P DEACETYLASE): RV3332

PROBABLE SUGAR TRANSPORT INTEGRAL MEMBRANE PROTEIN SUGI: RV3331

TI-SIGNAL FACTOR ANTAGONIST (REGULATOR OF SIGMA F B): RV3687C

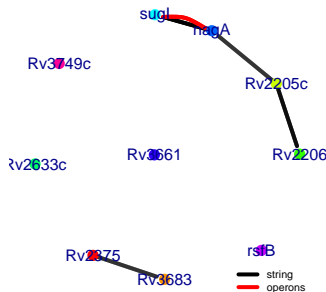
-4.89 hypothetical protein: RV3683

-3.62 hypothetical protein: RV3749C

-3.22 hypothetical protein: RV2375

-2.76 hypothetical protein: RV2633C

-2.50 hypothetical protein: RV3661



-200 -100 -1