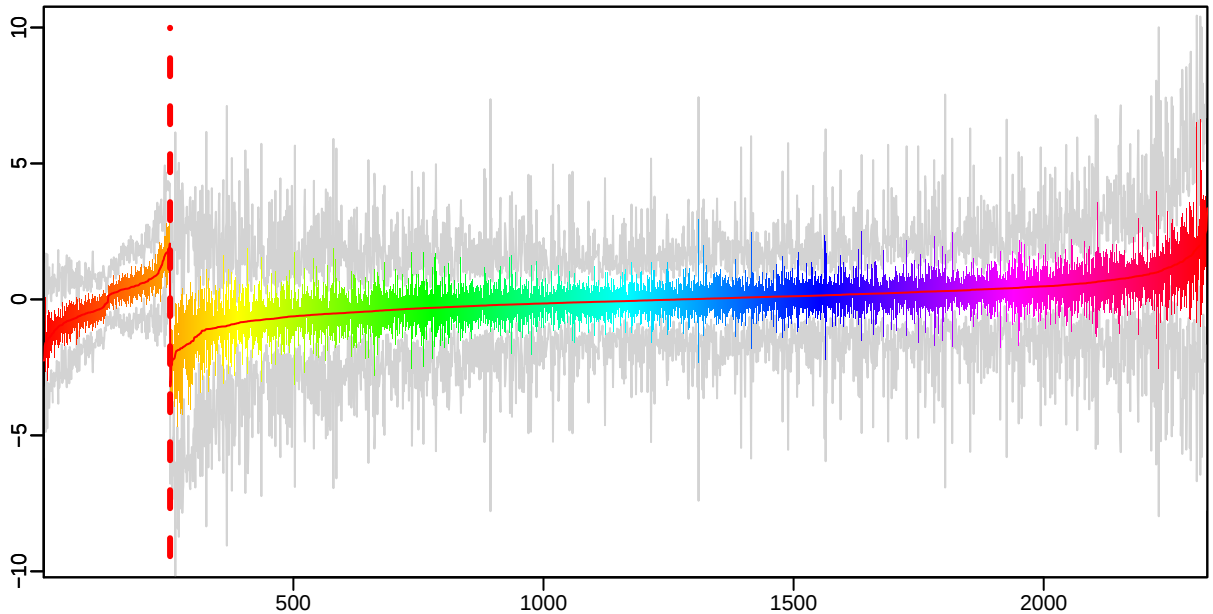
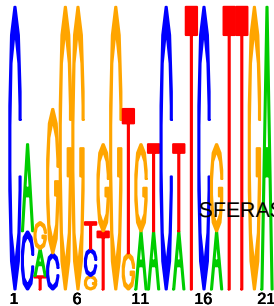
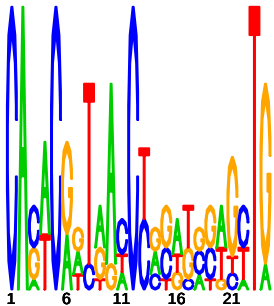


Cluster: 0033 mtu; resid: 0.56; r/c: 12/253

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



upstream meme PSSM #1; e=0.00027 upstream meme PSSM #2; e=0.00094



log10(P) upstream meme log10(P.clust)=-8.12; 12 seqs; 10 uniq

HYPOTHETICAL TRANSCRIPTIONAL REGULATORY PROTEIN: RV3095

-13.78 hypothetical protein: RV3094C

-13.78 HYPOTHETICAL OXIDOREDUCTASE: RV3093C

SFERASE) (1.107) ADENOSINE DIPHOSPHOGLUCOSE PYROPHOSPHORYLASE: RV0993

-13.07 hypothetical protein: RV0992C

-10.41 hypothetical protein: RV3849

PROBABLE CONSERVED INTEGRAL MEMBRANE PROTEIN: RV3092C

-4.46 hypothetical protein: RV1906C

-4.46 PROBABLE D-AMINO ACID OXIDASE AAO: RV1905C

-3.79 POSSIBLE MEMBRANE PROTEIN: RV3851

-0.53 PROBABLE OXIDOREDUCTASE: RV1257C

-0.52 hypothetical protein: RV1919C

