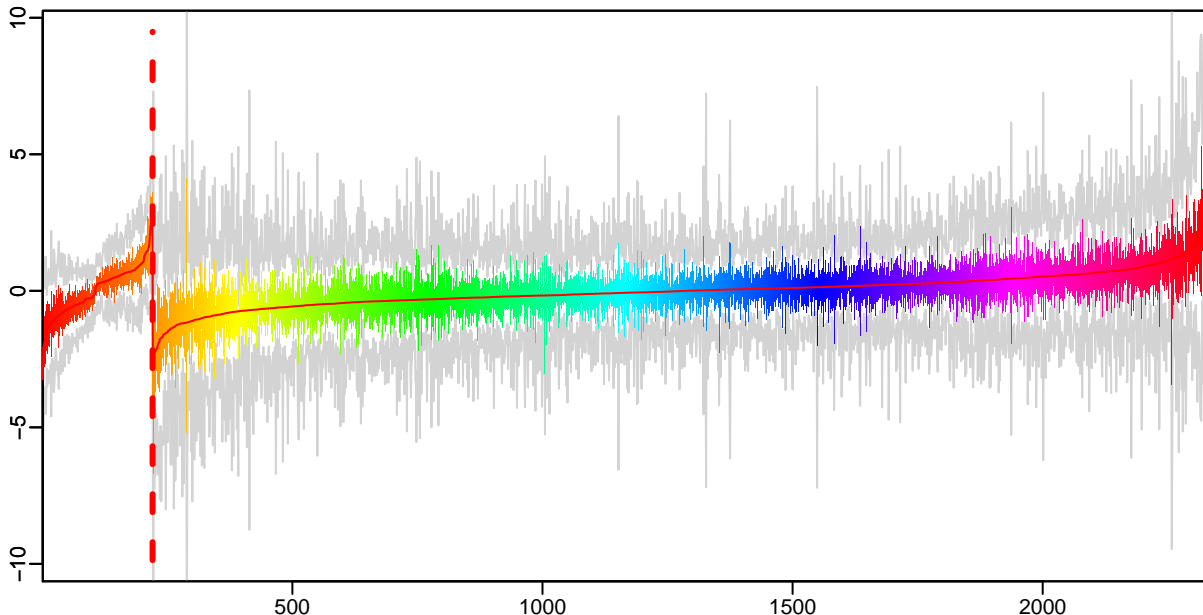
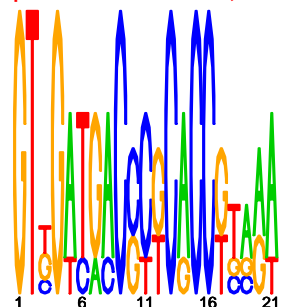


Cluster: 0024 mtu; resid: 0.54; r/c: 10/220

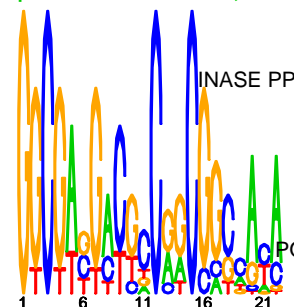
13 Oct 25 14:25:18 iter=2000  
cMonkey Version 4.9.8 mtu



upstream meme PSSM #1; e=1.3



upstream meme PSSM #2; e=1.5



log10(P) upstream meme

log10(P.clust)=-7.41; 10 seqs; 8 uniq

PHOSPHATE-GLUCOSE PHOSPHOTRANSFERASE: RV2702

POSSIBLE EXTRAGENIC SUPPRESSOR PROTEIN SUHB: RV2701C

hypothetical protein: RV2311

POSSIBLE CELL DIVISION TRANSMEMBRANE PROTEIN FTSK: RV2748C

TIGR ALANINE RICH PROTEIN (ESAT-6 LIKE PROTEIN 12): RV3904C

PROBABLE MEMBRANE PROTEIN: RV1744C

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

POSSIBLE SUGAR TRANSPORT INTEGRAL MEMBRANE PROTEIN SUGI: RV3331

hypothetical protein: RV3440C

SERVED-2YB HYPOTHETICAL ALANINE AND PROLINE RICH PROTEIN: RV3439C

