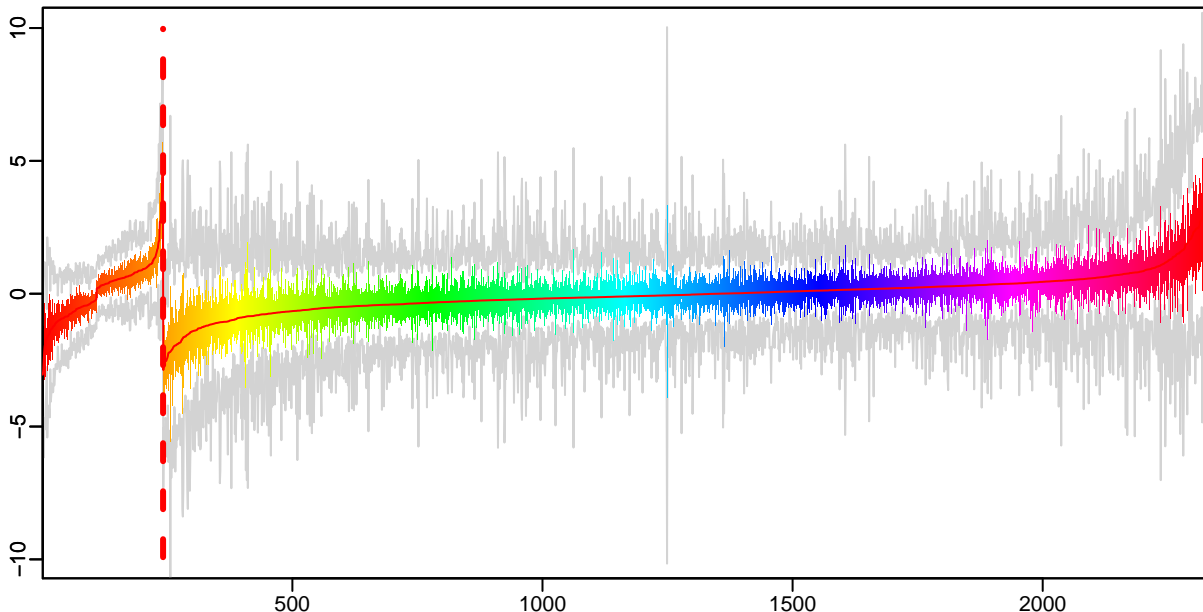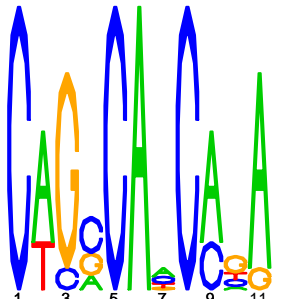# Cluster: 0405 mtu; resid: 0.59; r/c: 20/241
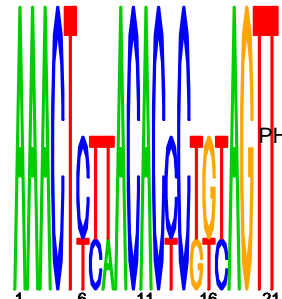

13 Oct 25 14:25:18 iter=2000
cMonkey Version 4.9.8 mtu

upstream meme PSSM #1; e=0.58

upstream meme PSSM #2; e=2.9

log10(P) upstream meme

log10(P.clust)=−5.03; 20 seqs; 10 uniq

log10(P)	Annotation
−12.49	hypothetical protein: RV0525
−12.49	glutamate−1−semialdehyde aminotransferase: RV0524
−12.49	hypothetical protein: RV0523C
−11.01	PHOH1 (PHOSPHATE STARVATION−INDUCIBLE PROTEIN PSIH): RV2368C
−10.02	hypothetical protein: RV2367C
−10.02	PROBABLE CONSERVED TRANSMEMBRANE PROTEIN: RV2366C
−10.02	hypothetical protein: RV2365C
−2.79	hypothetical protein: RV2325C
−2.42	phosphomannomutase/phosphoglucomutase: RV3257C
−2.42	hypothetical protein: RV3256C
−2.42	SE) (PMI) (PHOSPHOHEXOISOMERASE) (PHOSPHOHEXOMUTASE): RV3255C
−2.15	malate:quinone oxidoreductase: RV2852C
−2.15	hypothetical protein: RV2851C
−2.15	cobyrinic acid a,c−diamide synthase: RV2848C
−1.48	hypothetical protein: RV1026
−1.48	hypothetical protein: RV1025
−1.04	PHOH2 (PHOSPHATE STARVATION−INDUCIBLE PROTEIN PSIH): RV1095
−1.04	amidophosphoribosyltransferase: RV0808
−0.74	1−deoxy−D−xylulose 5−phosphate reductoisomerase: RV2870C
−0.77	PROBABLE CONSERVED TRANSMEMBRANE PROTEIN: RV2869C

string
operons