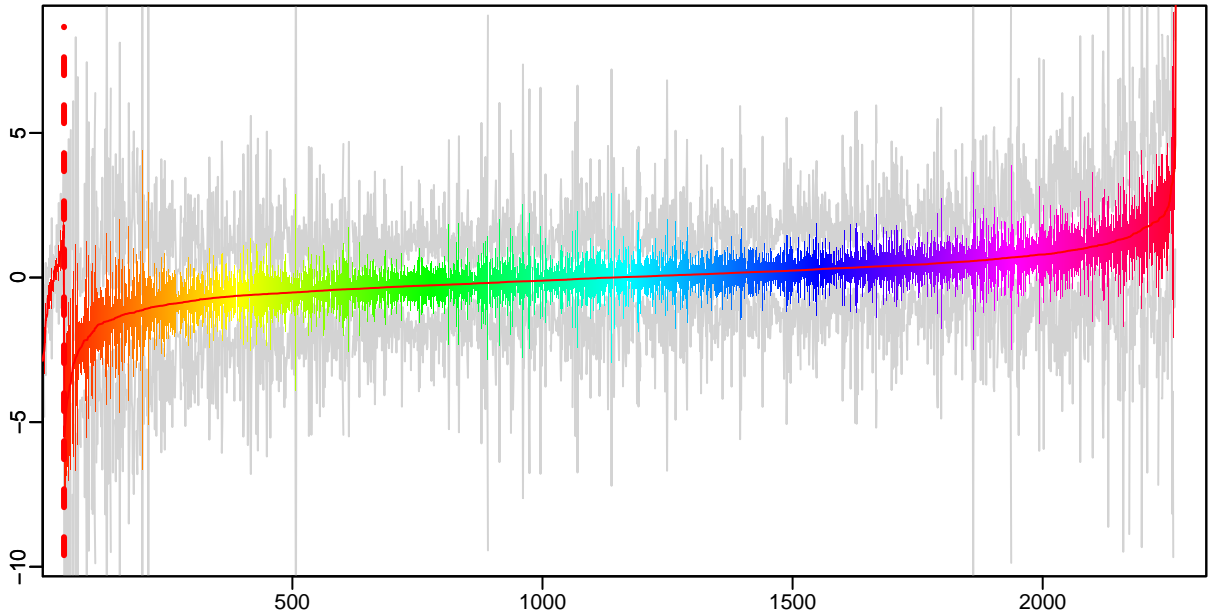
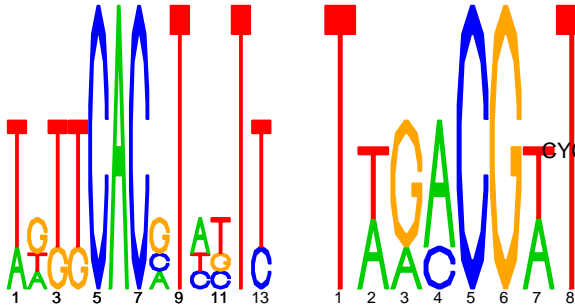


Cluster: 0095 mtu; resid: 0.35; r/c: 4/43

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



upstream meme PSSM #1; e=1.5e+00 upstream meme PSSM #2; e=3.4e+03



log10(P) upstream meme

log10(P.clust)=-4.26; 4 seqs; 4 uniq

CYCLASE 3 (ADP-PYROPHOSPHATE-LYASE) (ADENYLATE CYCLASE): RV1264

-4.41 PROBABLE TRANSPOSASE: RV3348

Rv1076 -3.93 hypothetical protein: RV1176C

Rv1264 -3.29 PROBABLE ATP-DEPENDENT HELICASE DING: RV1329C

Rv3348

-200

-100

-1