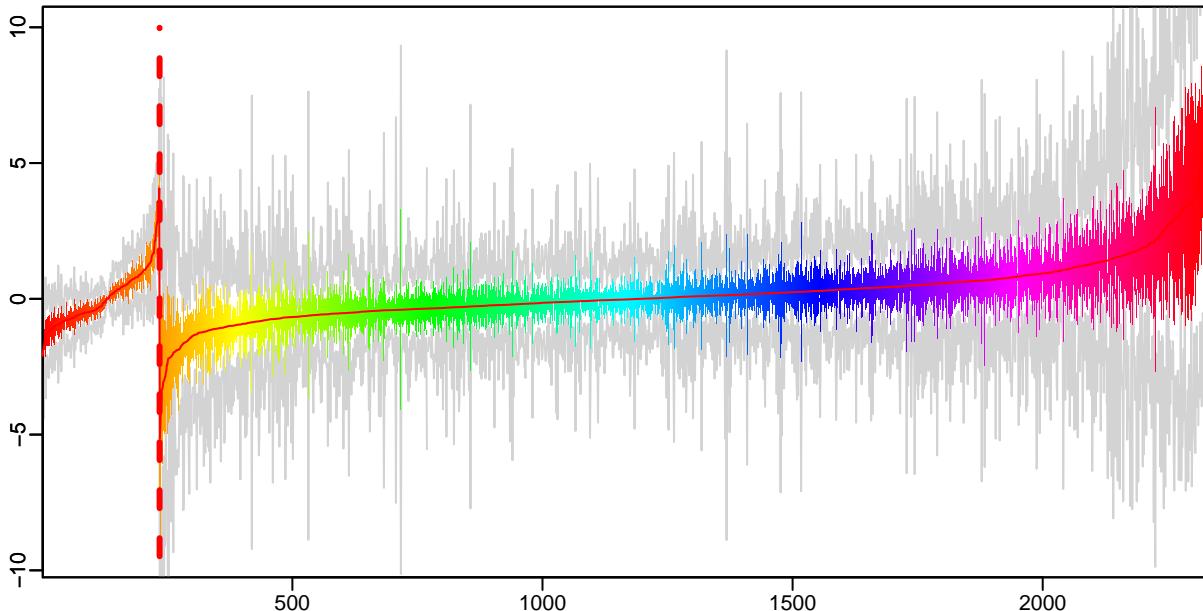
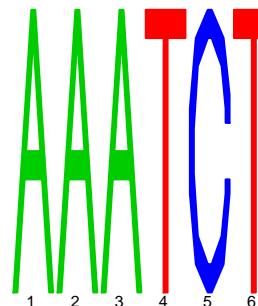
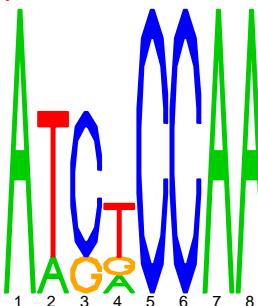


Cluster: 0328 mtu; resid: 0.36; r/c: 5/234

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



upstream meme PSSM #1; e=94 upstream meme PSSM #2; e=3.7e+03



log10(P) upstream meme

log10(P.clust)=-1.82; 5 seqs; 5 uniq

-2.88 PUTATIVE TRANSFERASE: RV1377C

-2.61 hypothetical protein: RV1982C

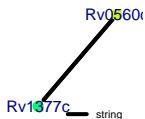
POSSIBLE BENZOQUINONE METHYLTRANSFERASE (METHYLASE): RV0560C

REGULATORY PROTEIN PKNK (PROTEIN KINASE K) (STPK K): RV3080C

-1.15 6-phosphofructokinase: RV3010C

Rv1082c

p16A



-200 -100 -1