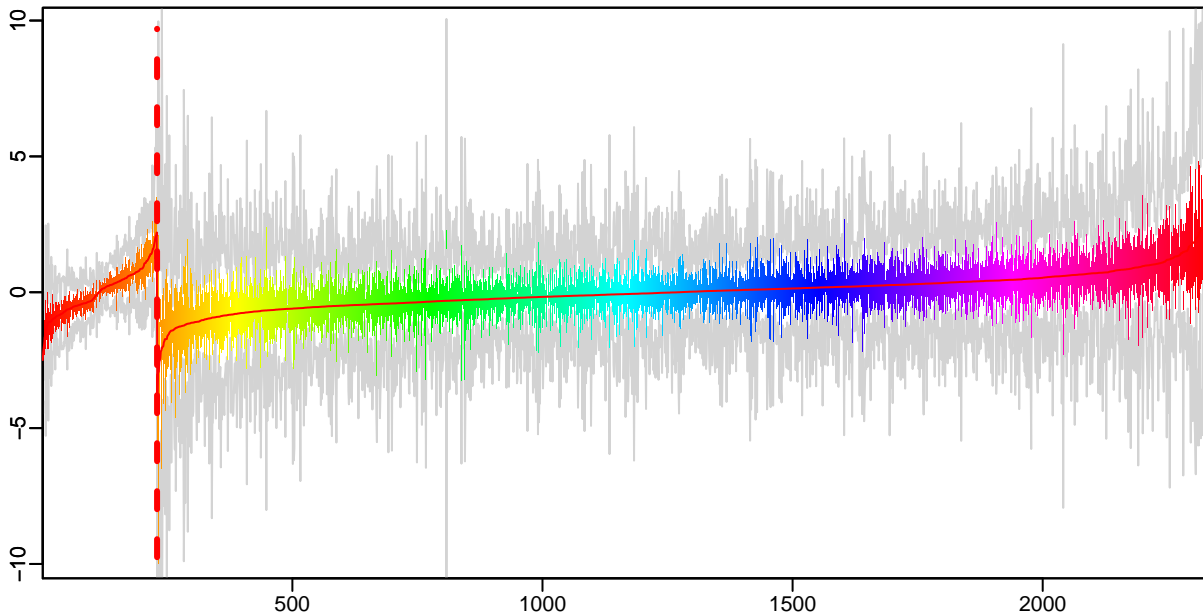
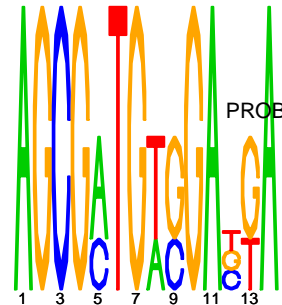
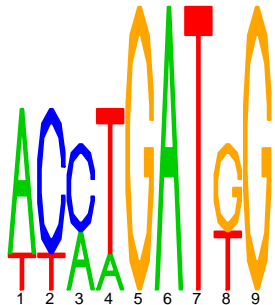


Cluster: 0147 mtu; resid: 0.43; r/c: 6/229

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



upstream meme PSSM #1; e=160 upstream meme PSSM #2; e=4.8e+03



log10(P) upstream meme

log10(P.clust)=-3.85; 6 seqs; 5 uniq

PROBABLE SHORT-CHAIN TYPE DEHYDROGENASE/REDUCTASE: RV1245C

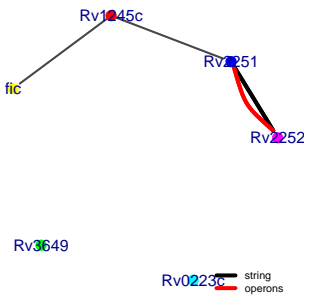
-5.42 POSSIBLE CELL FILAMENTATION PROTEIN FIC: RV3641C

-5.31 PROBABLE ALDEHYDE DEHYDROGENASE: RV0223C

-3.30 PROBABLE HELICASE: RV3649

-1.09 diacylglycerol kinase: RV2252

-1.09 POSSIBLE FLAVOPROTEIN: RV2251



-200 -100 -1