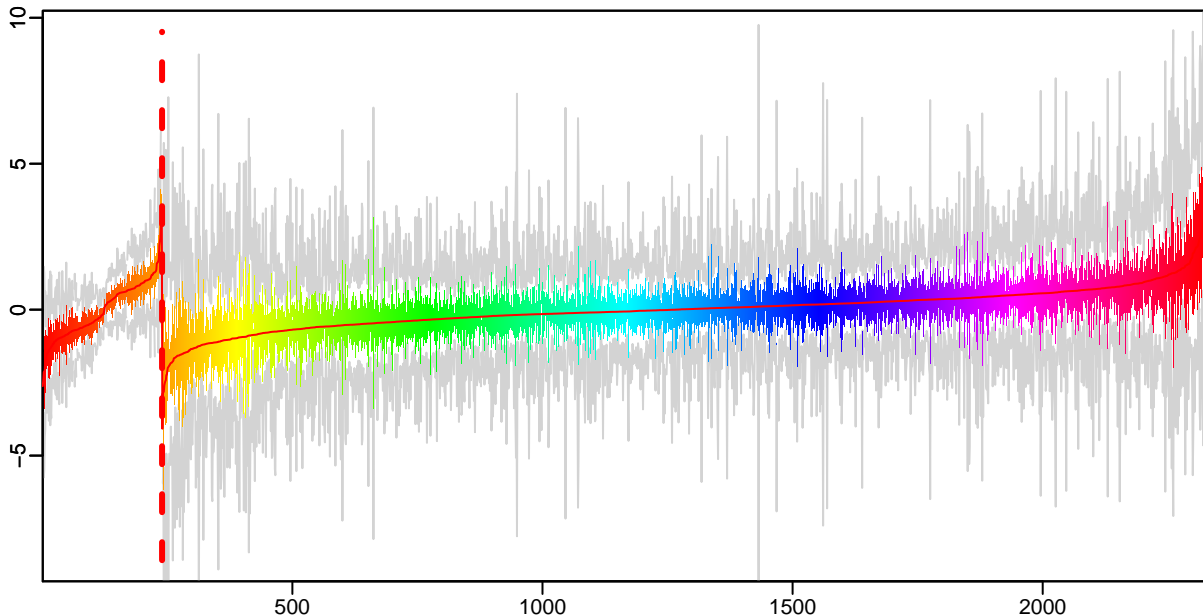
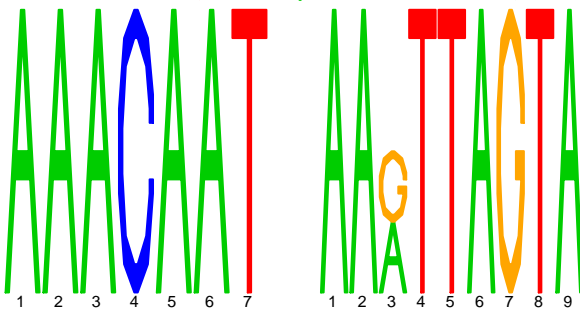


Cluster: 0338 mtu; resid: 0.49; r/c: 8/239

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



upstream meme PSSM #1; e=3.5e+00 upstream meme PSSM #2; e=3.8e+03



log10(P) upstream meme log10(P.clust)=-1.69; 8 seqs; 7 uniq

-4.66 PROBABLE TRANSPOSASE: RV3191C

PROBABLE POLYKETIDE BETA-KETOACYL SYNTHASE PKS4: RV1181

PROBABLE POLYKETIDE BETA-KETOACYL SYNTHASE PKS3: RV1180

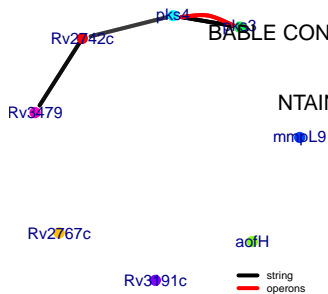
CONSERVED HYPOTHETICAL ARGININE RICH PROTEIN: RV2742C

PROBABLE CONSERVED TRANSMEMBRANE TRANSPORT PROTEIN Mmpl9: RV2339

CONTAINING MONOAMINE OXIDASE AOFH (AMINE OXIDASE) (MAO): RV3170

-0.00 POSSIBLE TRANSMEMBRANE PROTEIN: RV3479

-0.00 POSSIBLE MEMBRANE PROTEIN: RV2767C



-200 -100 -1