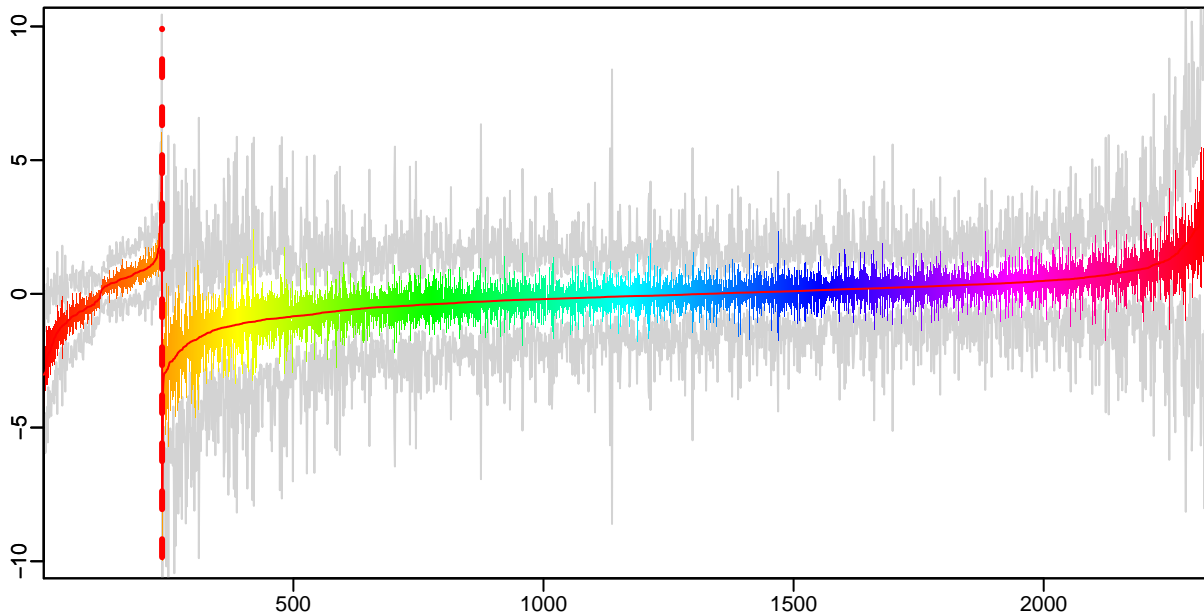
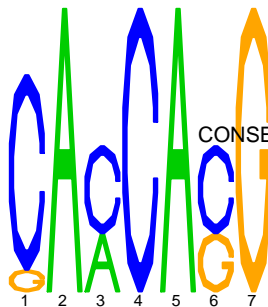
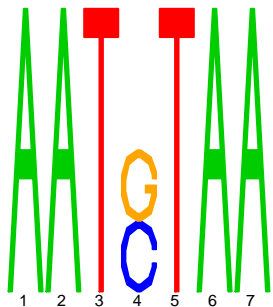


Cluster: 0303 mtu; resid: 0.47; r/c: 10/237

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



upstream meme PSSM #1; e=1.8e+00 upstream meme PSSM #2; e=1.8e+03



log10(P) upstream meme log10(P.clust)=-1.02; 10 seqs; 10 uniq

-2.41 hypothetical protein: RV2481C

CONSERVED INTEGRAL MEMBRANE TRANSPORT PROTEIN Mmpl8: RV3823C

-2.04 leucyl-tRNA synthetase: RV0041

-0.84 DNA polymerase I: RV1629

-0.78 acyl-CoA synthetase: RV1193

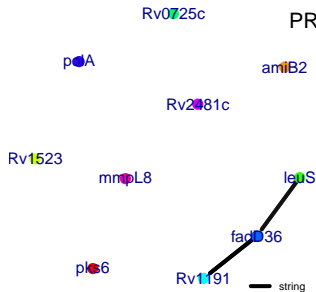
PROBABLE MEMBRANE BOUND POLYKETIDE SYNTHASE PKS6: RV0405

-0.52 Probable methyltransferase: RV1523

-0.33 hypothetical protein: RV1191

-0.33 hypothetical protein: RV0725C

-0.29 amidase: RV1263



-200 -100 -1