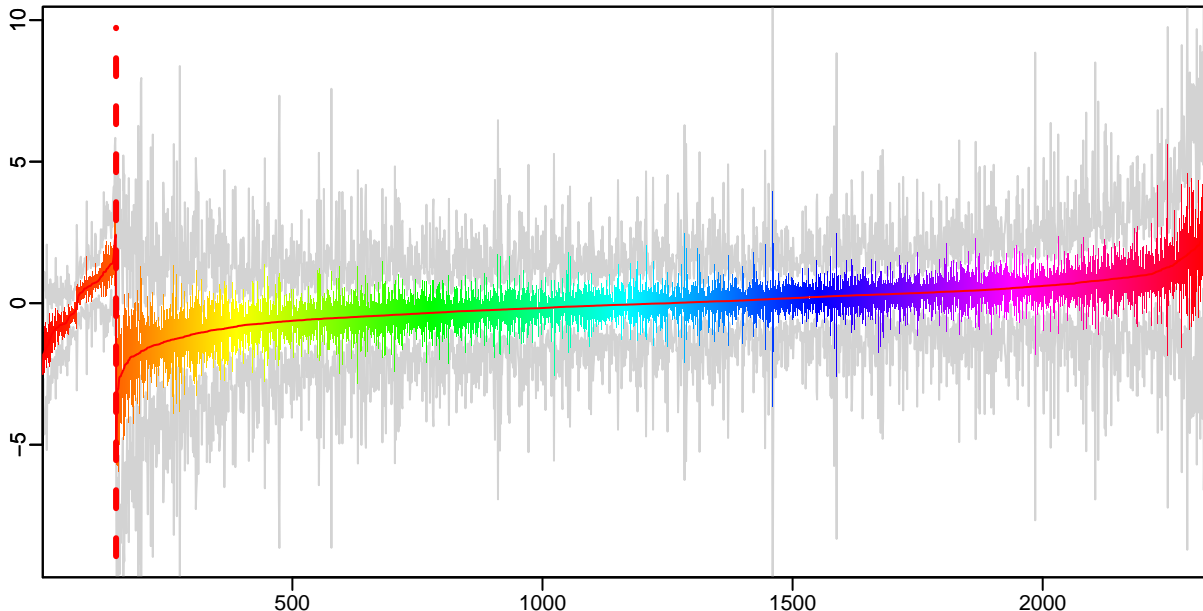
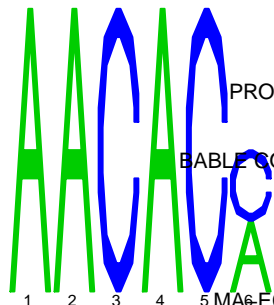
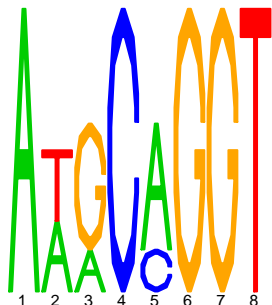


Cluster: 0050 mtu; resid: 0.49; r/c: 8/147

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



upstream meme PSSM #1; e=2.3e+00 upstream meme PSSM #2; e=2.8e+03



log10(P) upstream meme

log10(P.clust)=-1.78; 8 seqs; 4 uniq

PROBABLE CYTOCHROME P450 MONOOXYGENASE 142 CYP142: RV3518C

BABLE CONSERVED TRANSMEMBRANE TRANSPORT PROTEIN MMPL9: RV2339

-1.85 hypothetical protein: RV0052

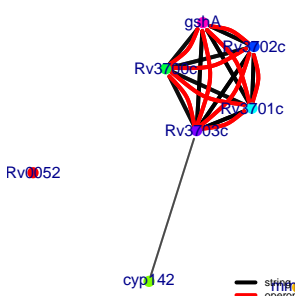
-1.67 (GCS) (GAMMA-GLUTAMYL-L-CYSTEINE SYNTHETASE): RV3704C

-1.67 hypothetical protein: RV3703C

-1.67 hypothetical protein: RV3702C

-1.67 hypothetical protein: RV3701C

-1.67 hypothetical protein: RV3700C



syn operons

-200 -100 -1