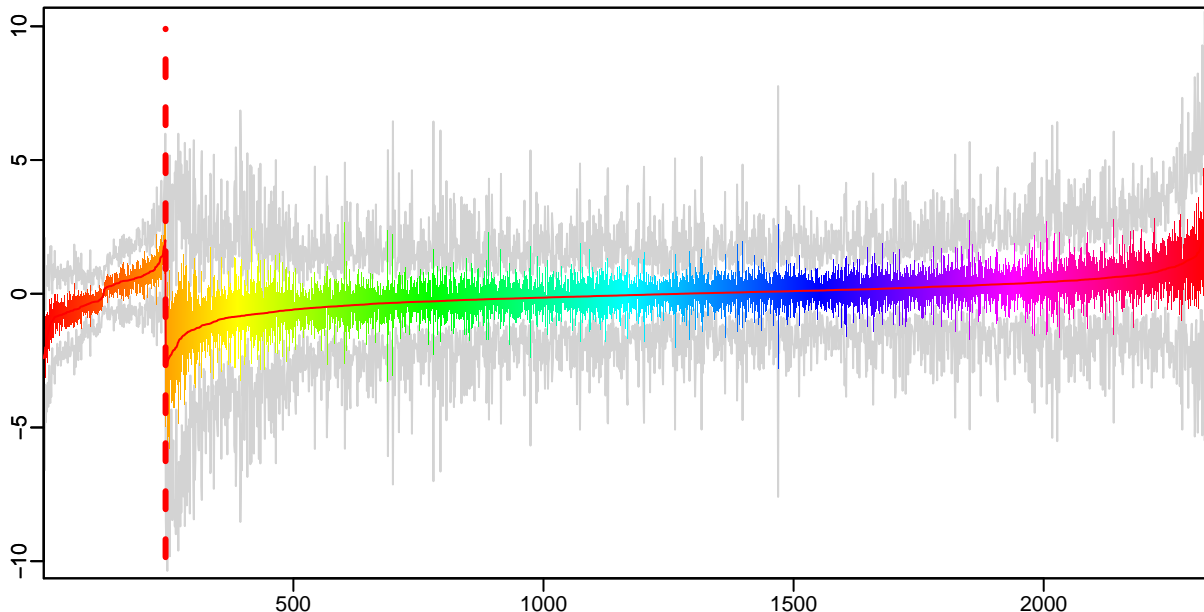
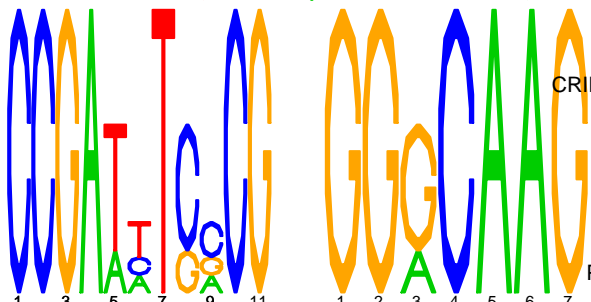


Cluster: 0457 mtu; resid: 0.53; r/c: 9/244

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



upstream meme PSSM #1; e=2.2e+03 upstream meme PSSM #2; e=4e+03



log10(P) upstream meme log10(P.clust)=-2.96; 9 seqs; 4 uniq

CRIP1ONAL REGULATORY PROTEIN (PROBABLY TETR-FAMILY): RV3557C

-3.14 acetyl-CoA acetyltransferase: RV3556C

-2.99 POSSIBLE MEMBRANE PROTEIN: RV0900

POSSIBLE CONSERVED EXPORTED OR MEMBRANE PROTEIN: RV0901

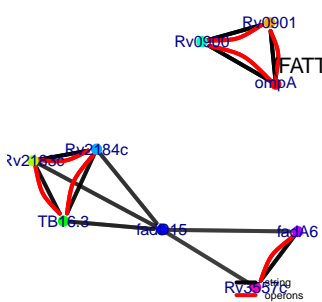
-2.99 OUTER MEMBRANE PROTEIN A OMPA: RV0899

FATTY-ACID-CoA SYNTHETASE (FATTY ACID-CoA SYNTHASE): RV2187

-2.82 hypothetical protein: RV2184C

-2.82 hypothetical protein: RV2185C

-2.82 hypothetical protein: RV2183C



-200 -100 -1