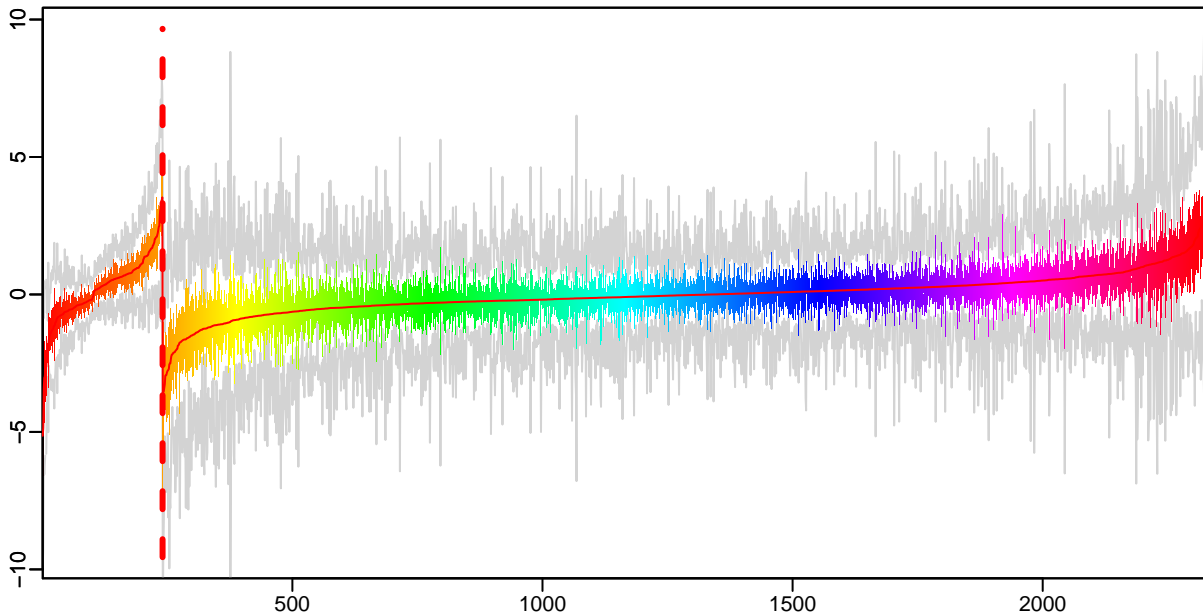
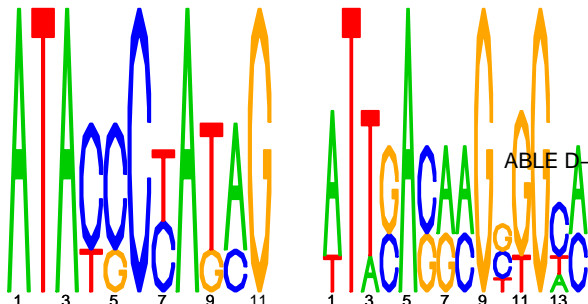


Cluster: 0512 mtu; resid: 0.54; r/c: 8/240

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



upstream meme PSSM #1; e=1.3e+00 upstream meme PSSM #2; e=6.6e+03



log10(P) upstream meme log10(P.clust)=-3.68; 8 seqs; 8 uniq

-6.52 PROBABLE BACTERIOFERRITIN BFRA: RV1876

ABLE D-3-PHOSPHOGLYCERATE DEHYDROGENASE SERA1 (PGDH): RV2996C

-4.69 PROBABLE DEHYDROGENASE: RV1812C

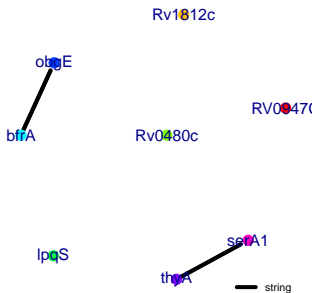
-3.57 PROBABLE LIPOPROTEIN LPQS: RV0847

-3.32 GTPase ObgE: RV2440C

-2.49 thymidylate synthase: RV2764C

-0.00 POSSIBLE AMIDOHYDROLASE: RV0480C

NA RV0947C



-200 -100 -1