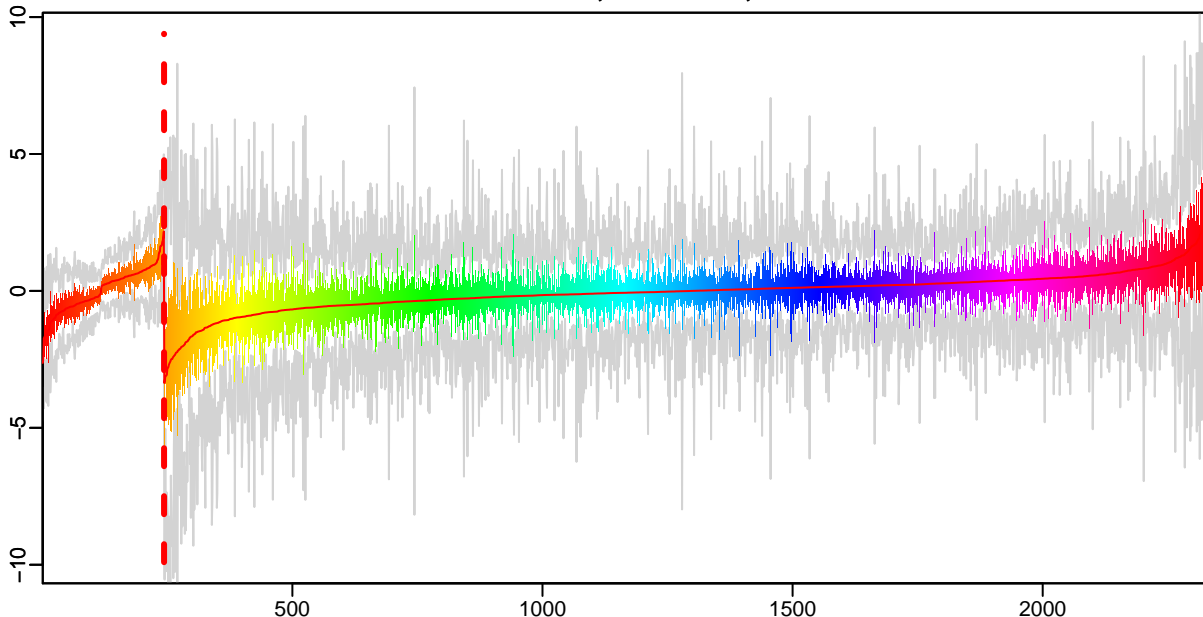
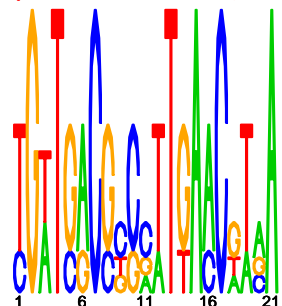


Cluster: 0415 mtu; resid: 0.48; r/c: 9/243

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



upstream meme PSSM #1; e=37 upstream meme PSSM #2; e=3.5e+03



log10(P) upstream meme log10(P.clust)=-6.84; 9 seqs; 5 uniq

IDE (FORMATE DEHYDROGENASE H ALPHA SUBUNIT) (FDH-H): RV2900C

-9.12 formate dehydrogenase accessory protein: RV2899C

-7.19 Probable polyketide synthase pks5: RV1527C

-7.04 spermidine synthase: RV2601

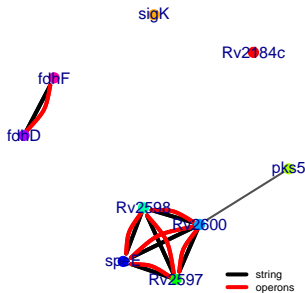
PROBABLE CONSERVED INTEGRAL MEMBRANE PROTEIN: RV2600

-7.04 hypothetical protein: RV2598

-7.04 PROBABLE MEMBRANE PROTEIN: RV2597

-6.54 RNA polymerase sigma factor SigK: RV0445C

-1.39 hypothetical protein: RV2184C



-200 -100 -1