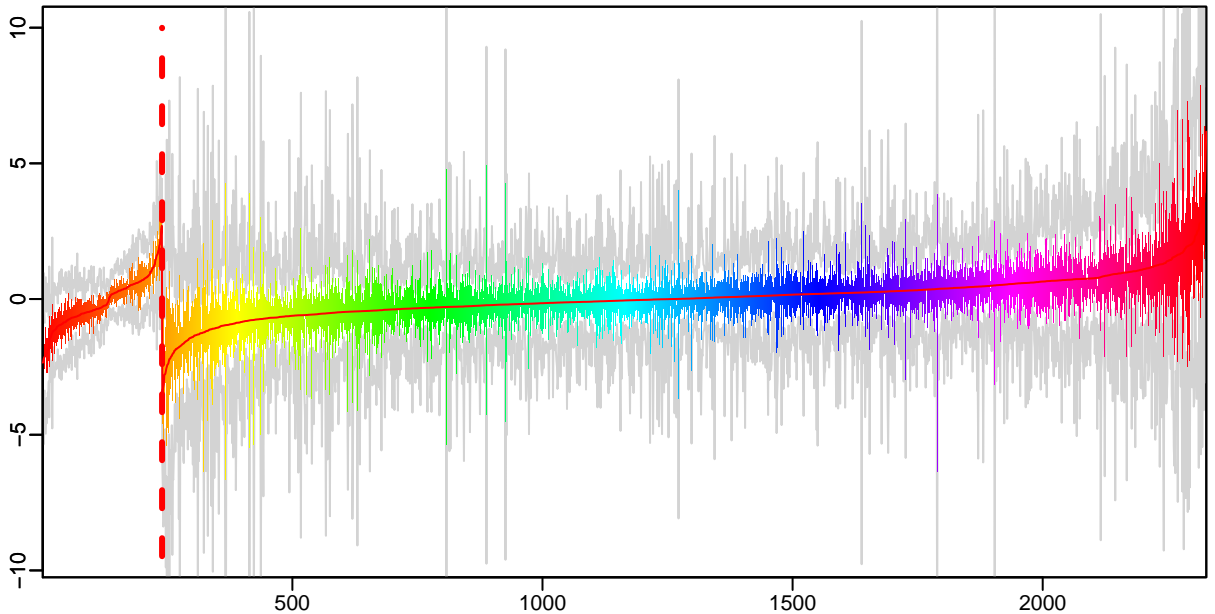
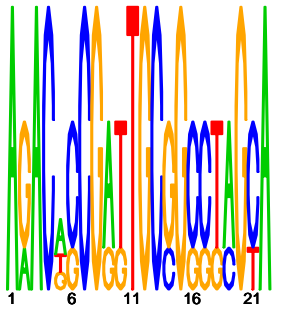


Cluster: 0489 mtu; resid: 0.46; r/c: 6/239

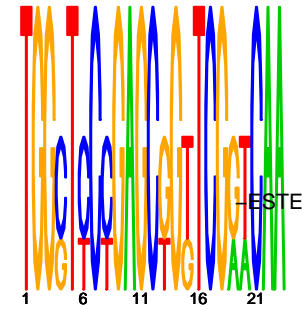
13 Oct 25 14:25:18 iter=2000
cMonkey Version 4.9.8 mtu



upstream meme PSSM #1; e=0.1



upstream meme PSSM #2; e=6.5



log10(P) upstream meme

log10(P.clust)=-12.99; 6 seqs; 5 uniq

-19.39 PROBABLE ACYL-CoA DEHYDROGENASE FADE4: RV0231

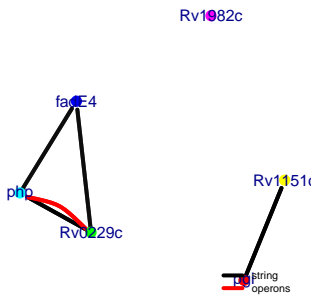
-19.39 ESTERASE (ARYLTRIPHOSPHATASE) (PARAOXON HYDROLASE): RV0230C

-19.39 POSSIBLE CONSERVED MEMBRANE PROTEIN: RV0229C

-8.30 NAD-dependent deacetylase: RV1151C

-5.86 hypothetical protein: RV1982C

-5.60 glucose-6-phosphate isomerase: RV0946C



-200 -100 -1