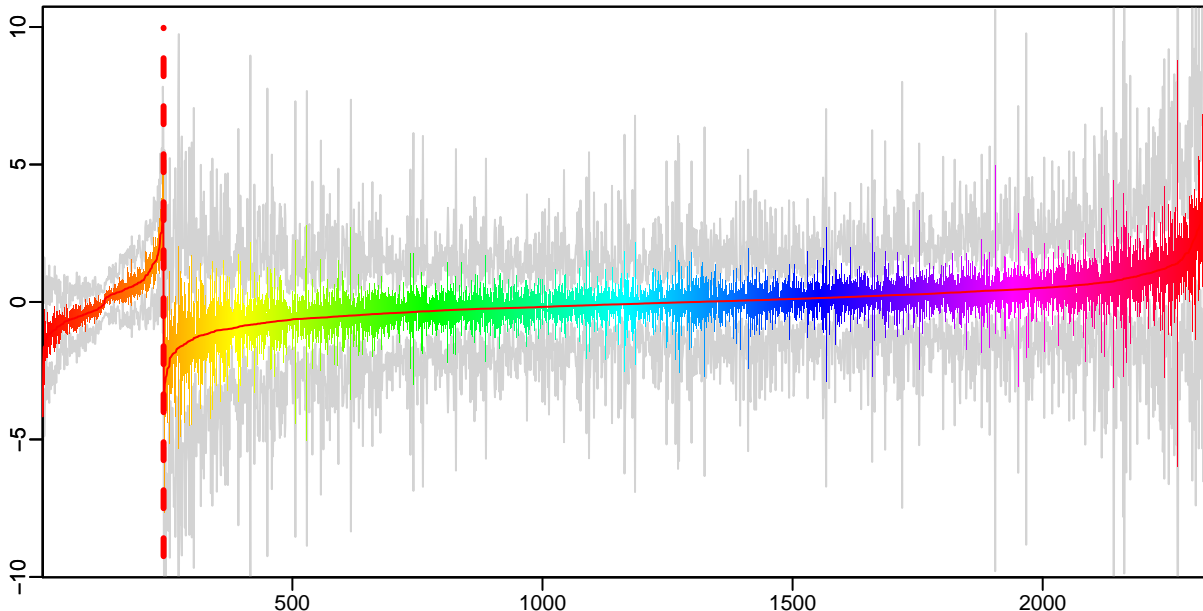
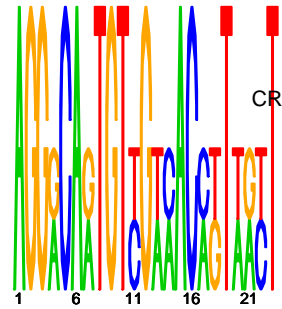
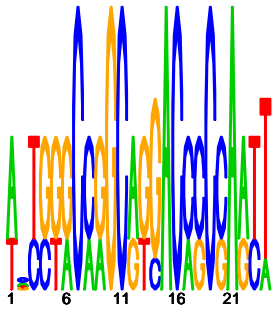


Cluster: 0249 mtu; resid: 0.44; r/c: 7/242

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



upstream meme PSSM #1; e=1e-05 upstream meme PSSM #2; e=0.00019



log10(P) upstream meme log10(P.clust)=-12.83; 7 seqs; 6 uniq

CRIP10A5 REGULATORY PROTEIN (POSSIBLY TETR-FAMILY): RV0472C

-17.95 POSSIBLE CONSERVED TRANSMEMBRANE PROTEIN: RV0473

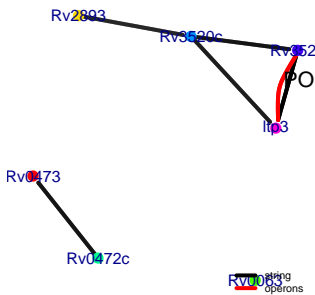
-14.07 acetyl-CoA acetyltransferase: RV3523

-14.07 hypothetical protein: RV3521

POSSIBLE COENZYME F420-DEPENDENT OXIDOREDUCTASE: RV3520C

-6.00 POSSIBLE OXIDOREDUCTASE: RV2893

-5.69 POSSIBLE OXIDOREDUCTASE: RV0063



-200 -100 -1