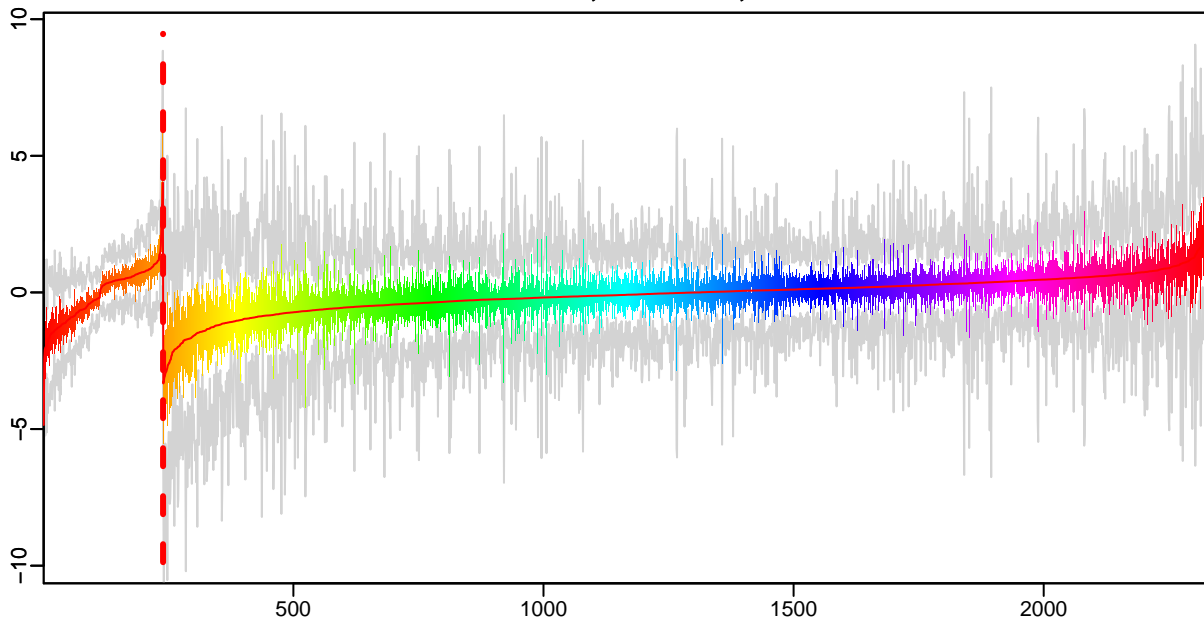


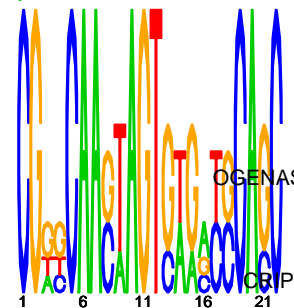
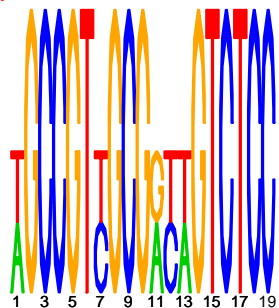
Cluster: 0561 mtu; resid: 0.52; r/c: 11/239

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



upstream meme PSSM #1; e=0.027

upstream meme PSSM #2; e=9.5



log10(P) upstream meme

log10(P.clust)=-6.80; 11 seqs; 7 uniq

-14.68 oligoribonuclease: RV2511

-14.68 hypothetical protein: RV2510C

OGENASE PCD4 PIPERIDEINE-6-CARBOXYLATE DEHYDROGENASE: RV3293

-7.24 hypothetical protein: RV3292

ORPTIONAL REGULATORY PROTEIN (PROBABLY ASNC-FAMILY): RV3291C

PROBABLE NADP-DEPENDENT ALCOHOL DEHYDROGENASE ADHC: RV3045

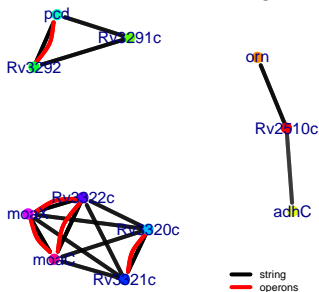
-5.77 molybdenum cofactor biosynthesis protein C: RV3324C

-5.77 PROBABLE MOAD-MOAE FUSION PROTEIN MOAX: RV3323C

-5.77 POSSIBLE METHYLTRANSFERASE: RV3322C

-0.22 hypothetical protein: RV3321C

-0.22 hypothetical protein: RV3320C



— string  
— operons

-200 -100 -1