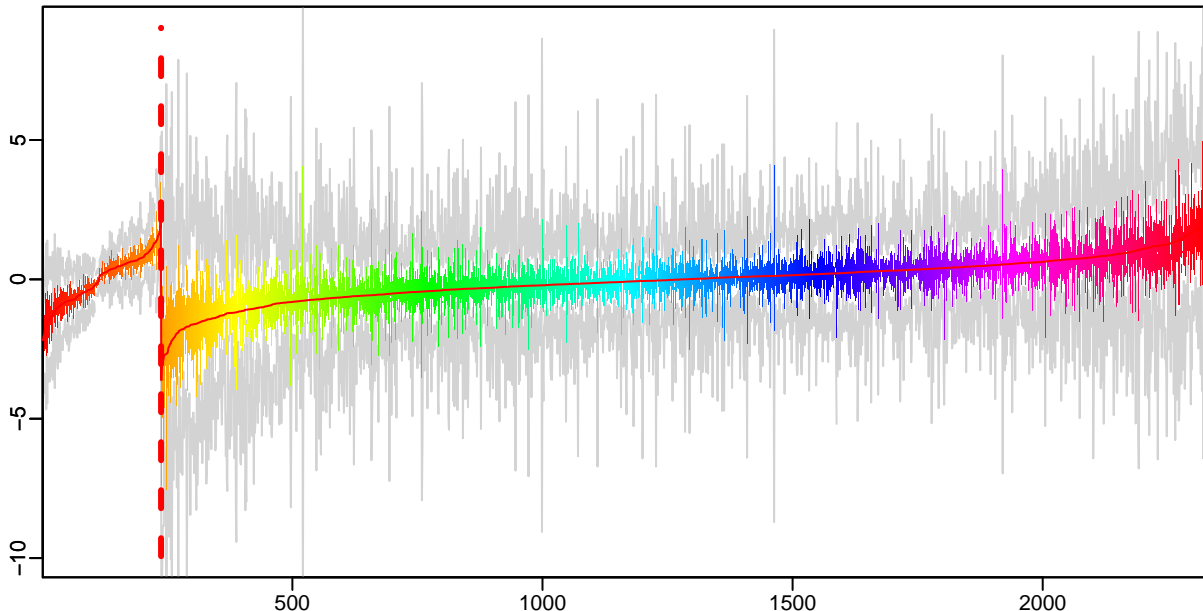
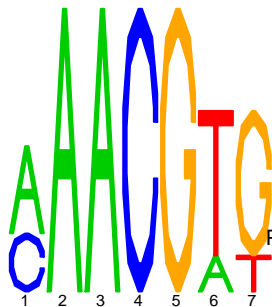
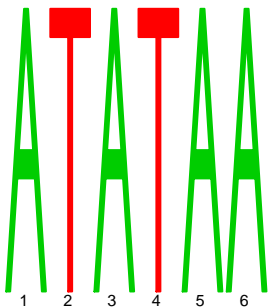


Cluster: 0309 mtu; resid: 0.40; r/c: 5/237

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



upstream meme PSSM #1; e=1.4e+03 upstream meme PSSM #2; e=2e+03



log10(P) upstream meme

log10(P.clust)=-1.39; 5 seqs; 5 uniq

-2.44 PROBABLE BACTERIOFERRITIN BFRA: RV1876

-1.01 POSSIBLE ALANINE AND PROLINE RICH MEMBRANE PROTEIN: RV2473

-0.86 PROBABLE HELICASE HELZ: RV2101

-0.63 PE FAMILY PROTEIN: RV1430

-0.63 ribonucleotide diphosphate reductase subunit beta: RV1981C

bfrA

ndrF

PS16

helZ

RV2473

-200

-100

-1