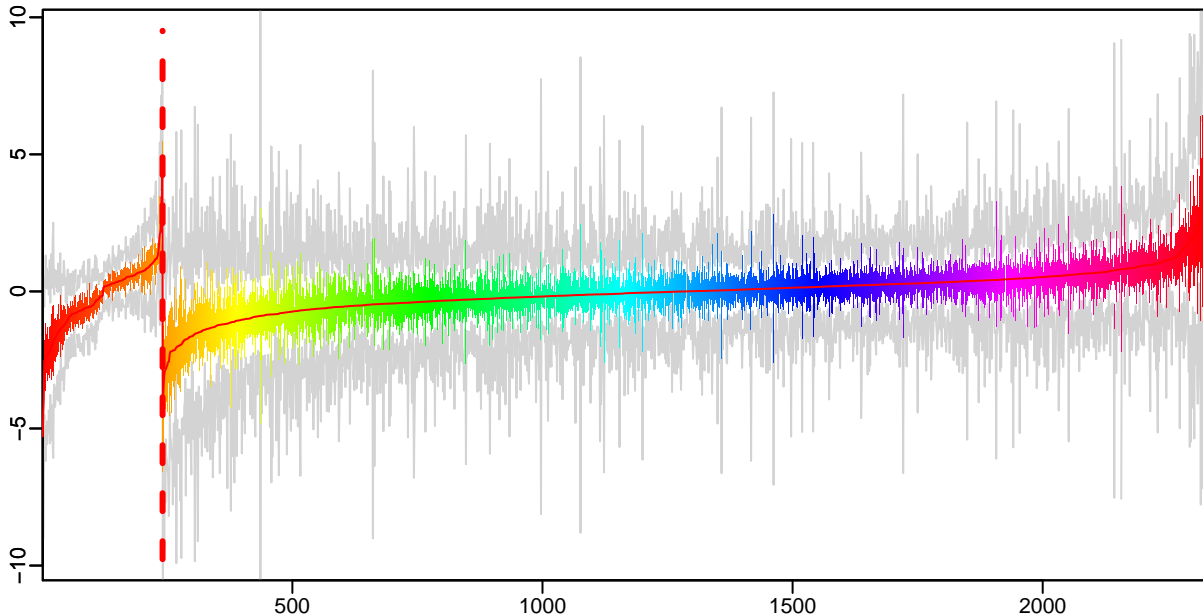
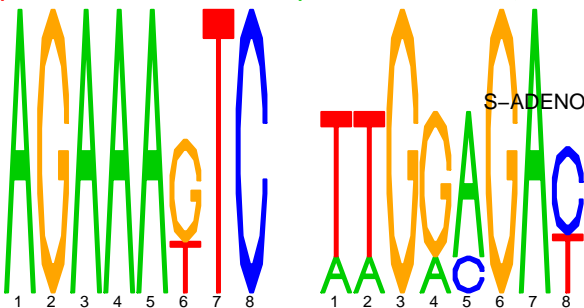


Cluster: 0579 mtu; resid: 0.51; r/c: 7/240

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



upstream meme PSSM #1; e=940 upstream meme PSSM #2; e=3.5e+03



Rv1607c

Rv2297

Rv0695 Rv0696

Rv0697

mRA

— string
— operons

log10(P) upstream meme log10(P.clust)=-2.35; 7 seqs; 5 uniq

S-ADENOSYL-L-HOMOCYSTEINE HOMOCYSTEINYL-RIBOHYDROLASE: RV0091

-3.39 hypothetical protein: RV1507C

-2.50 PROBABLE DEHYDROGENASE: RV0697

-2.50 PROBABLE MEMBRANE SUGAR TRANSFERASE: RV0696

-2.50 hypothetical protein: RV0695

-1.04 hypothetical protein: RV2297

-0.76 UDP-N-acetylglucosamine 1-carboxyvinyltransferase: RV1315

-200 -100 -1