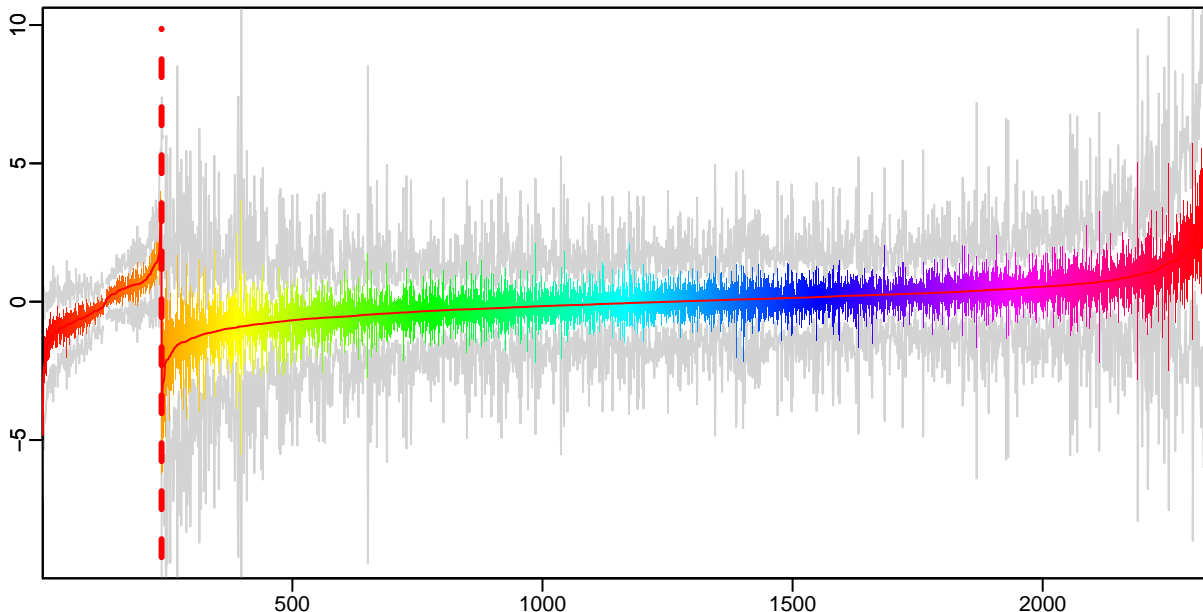
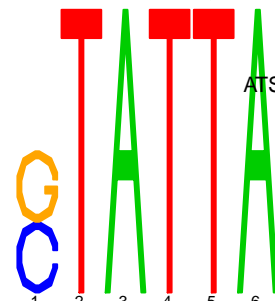
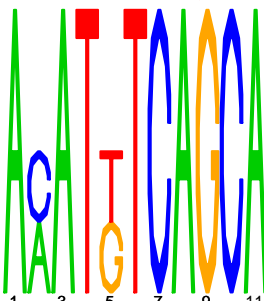


Cluster: 0189 mtu; resid: 0.44; r/c: 9/238

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



upstream meme PSSM #1; e=1.3e+04 upstream meme PSSM #2; e=1.3e+04



log10(P) upstream meme log10(P.clust)=-2.01; 9 seqs; 6 uniq

ATSD (ARYL SULFATE SULPHOHYDROLASE) (ARYLSULPHATASE): RV0863

-2.87 hypothetical protein: RV2975C

-2.00 CONSERVED HYPOTHETICAL ALANINE RICH PROTEIN: RV2974C

-2.87 PROBABLE ATP-DEPENDENT DNA HELICASE RECG: RV2973C

POSSIBLE CONSERVED MEMBRANE OR EXPORTED PROTEIN: RV2972C

-0.87 methionyl-tRNA synthetase: RV1007C

-0.75 hypothetical protein: RV1002C

-0.35 hypothetical protein: RV1896C

-0.01 PROBABLE DNA HELICASE ERCC3: RV0861C

