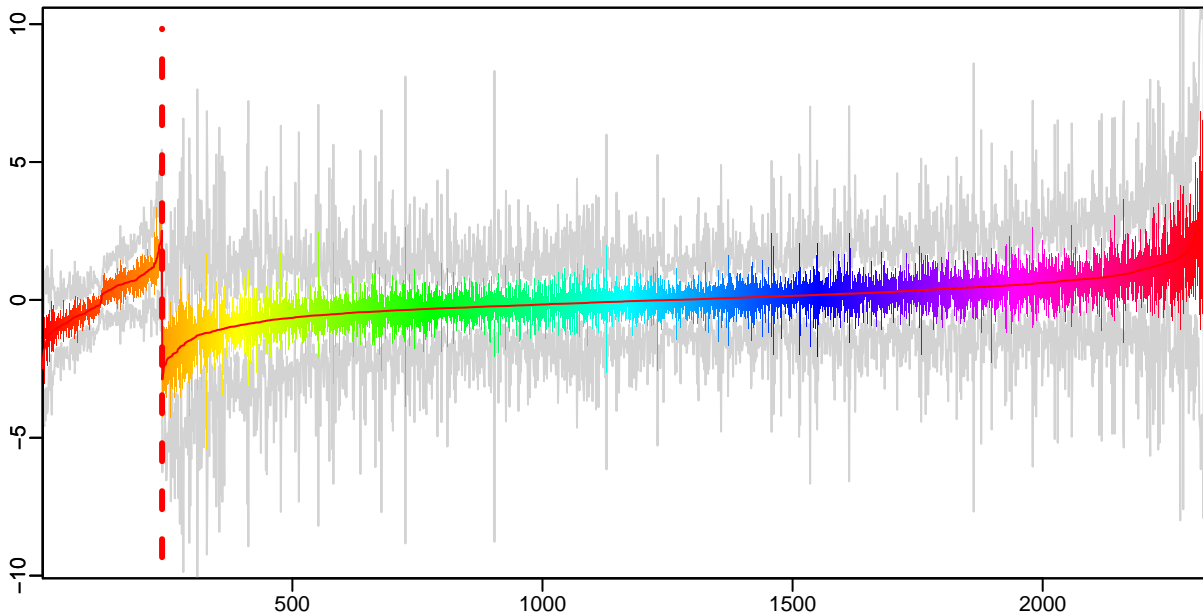
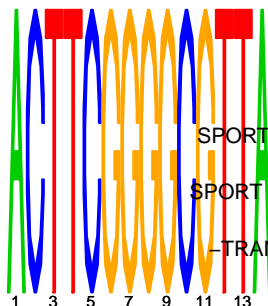
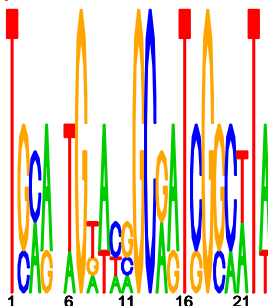


Cluster: 0518 mtu; resid: 0.50; r/c: 10/239

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



upstream meme PSSM #1; e=410 upstream meme PSSM #2; e=4.1e+03



log10(P) upstream meme log10(P.clust)=-6.50; 10 seqs; 6 uniq

-7.38 PROBABLE SULFATE-BINDING LIPOPROTEIN SUB: RV2400C

SPORT INTEGRAL-MEMBRANE PROTEIN ABC TRANSPORTER-CYST: RV2399C

SPORT INTEGRAL-MEMBRANE PROTEIN ABC TRANSPORTER-CYSW: RV2398C

-TRANSPORT ATP-BINDING PROTEIN ABC TRANSPORTER-CYSA1: RV2397C

-6.82 hypothetical protein: RV3054C

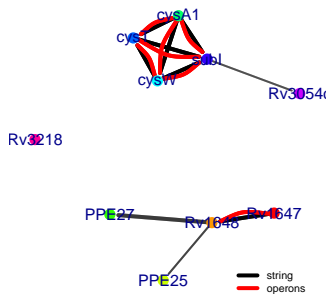
-6.51 Probable transmembrane protein: RV1648

-6.51 hypothetical protein: RV1647

-5.42 hypothetical protein: RV3218

-5.32 PPE FAMILY PROTEIN: RV1790

-4.88 PPE FAMILY PROTEIN: RV1787



-200 -100 -1