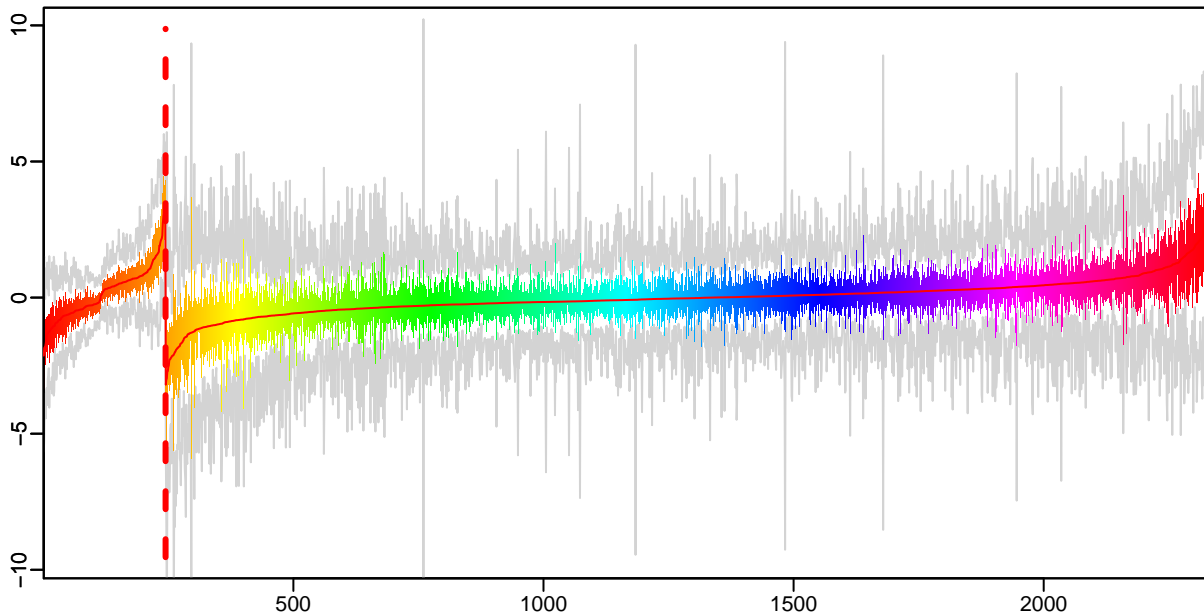
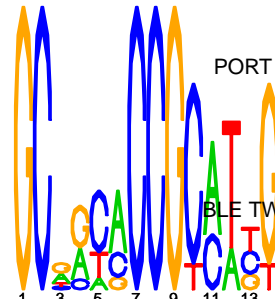
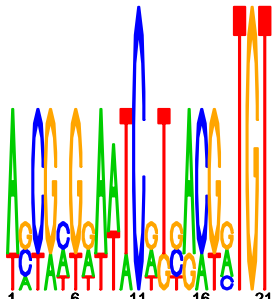


Cluster: 0526 mtu; resid: 0.56; r/c: 12/244

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



upstream meme PSSM #1; e=450 upstream meme PSSM #2; e=8.1e+03



log10(P) upstream meme log10(P.clust)=-4.50; 12 seqs; 10 uniq

PORT SYSTEM TRANSCRIPTIONAL REGULATORY PROTEIN PHOY2: RV0821C

-7.27 hypothetical protein: RV0857

-7.27 hypothetical protein: RV3272

ABLE TWO COMPONENT TRANSCRIPTIONAL REGULATORY PROTEIN: RV3765C

-6.94 hypothetical protein: RV1425

-3.53 POSSIBLE CONSERVED MEMBRANE PROTEIN: RV1417

-3.53 riboflavin synthase subunit beta: RV1416

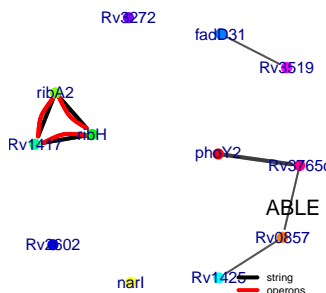
-3.53 ne 4-phosphate synthase/GTP cyclohydrolase II protein: RV1415

-2.44 acyl-CoA synthetase: RV1925

-2.24 hypothetical protein: RV2602

ABLE RESPIRATORY NITRATE REDUCTASE (GAMMA CHAIN) NAR1: RV1164

-0.13 hypothetical protein: RV3519



-200 -100 -1