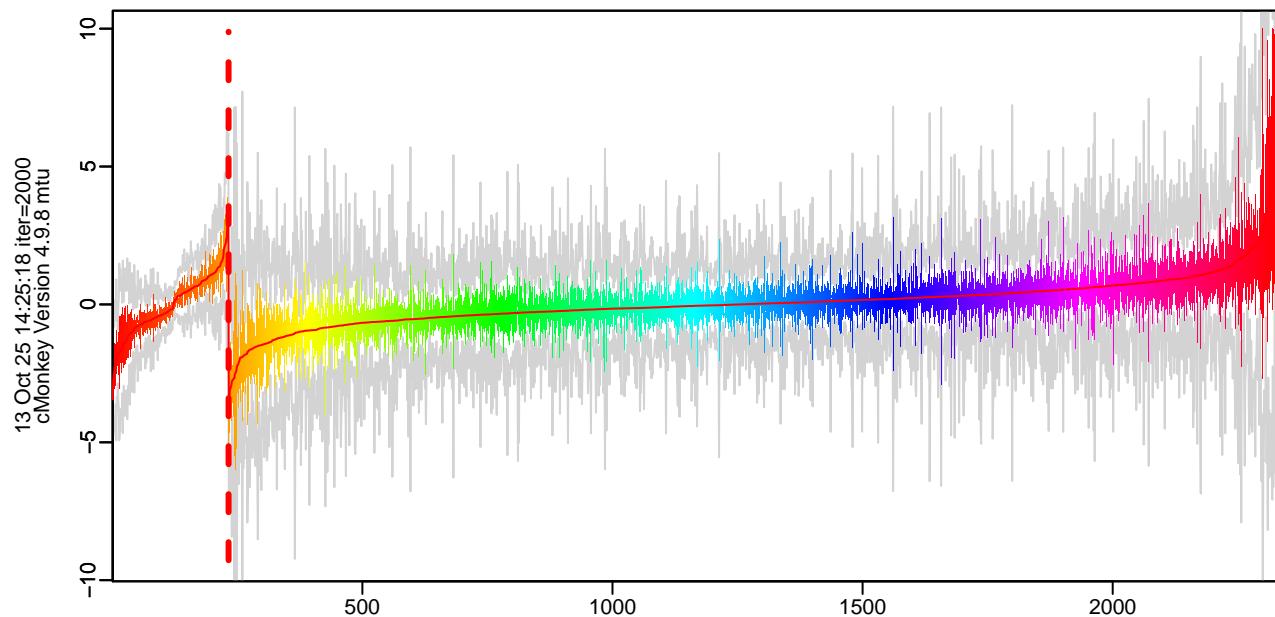
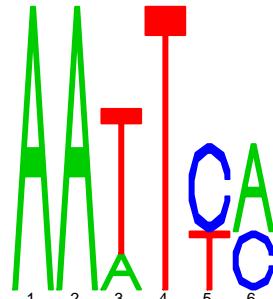


Cluster: 0600 mtu; resid: 0.49; r/c: 6/232



upstream meme PSSM #1; e=2.2e+03 upstream meme PSSM #2; e=750



$\log_{10}(P)$  upstream meme

$\log_{10}(P.\text{clust}) = -1.50$ ; 6 seqs; 5 uniq

-2.20

hypothetical protein: RV1684

-1.83

tRNA (guanine-N(1)-) methyltransferase: RV2906C

-1.66

hypothetical protein: RV3463

-1.66

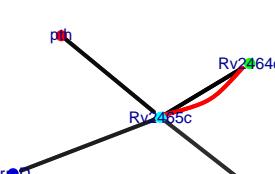
peptidyl tRNA hydrolase: RV1014C

-0.82

ribose-5-phosphate isomerase B: RV2465C

-0.82

POSSIBLE DNA GLYCOSYLASE: RV2464C



Rv3463

string  
operons

-200

-100

-1