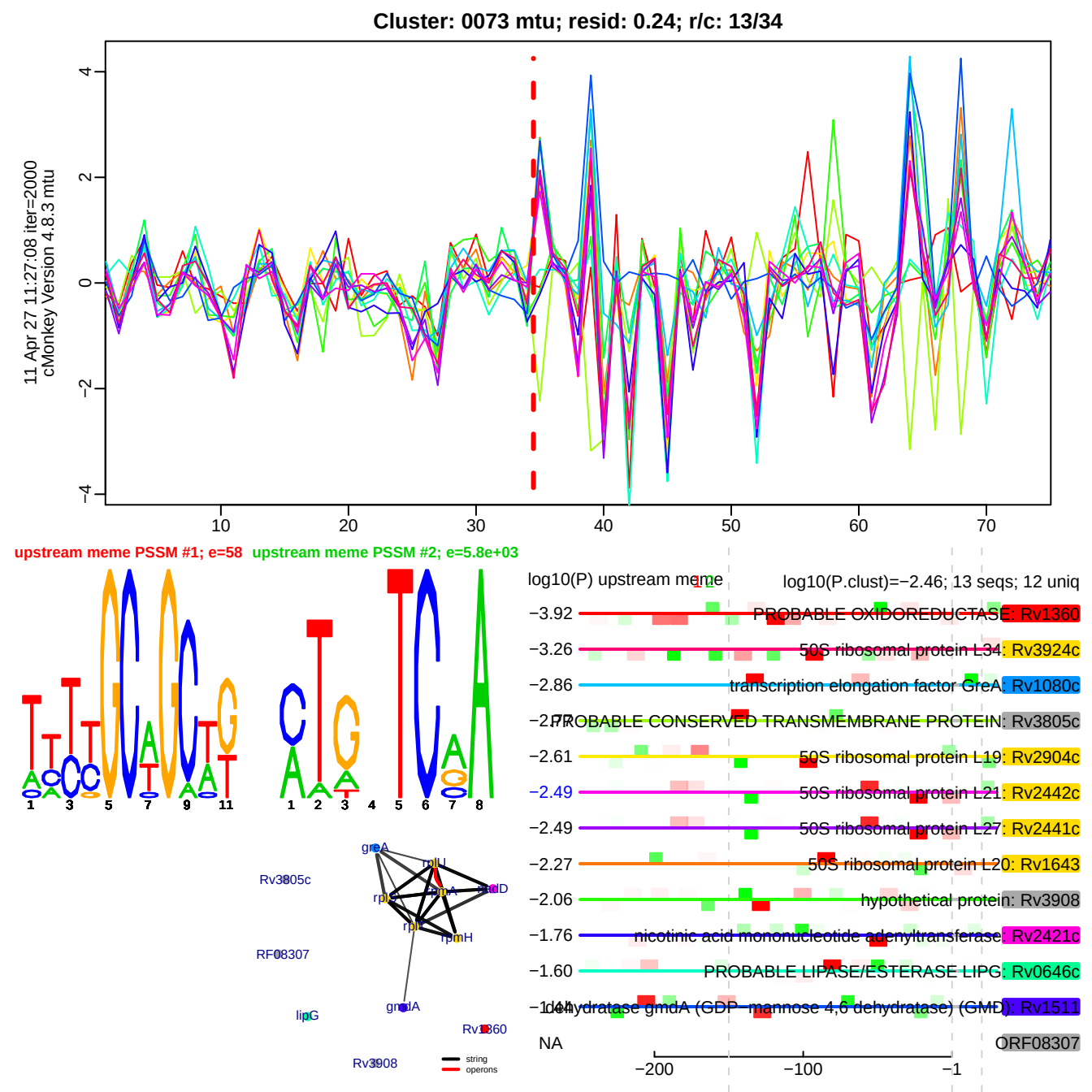

Cluster: 0073 mtu; resid: 0.24; r/c: 13/34
11 Apr 27 11:27:08 iter=2000
cMonkey Version 4.8.3 mtu
upstream meme PSSM #1; e=58
upstream meme PSSM #2; e=5.8e+03
log10(P) upstream meme
log10(P.clust)=−2.46; 13 seqs; 12 uniq
−3.92 PROBABLE OXIDOREDUCTASE: Rv1360
−3.26 50S ribosomal protein L34: Rv3924c
−2.86 transcription elongation factor GreA: Rv1080c
−2.77 PROBABLE CONSERVED TRANSMEMBRANE PROTEIN: Rv3805c
−2.61 50S ribosomal protein L19: Rv2904c
−2.49 50S ribosomal protein L21: Rv2442c
−2.49 50S ribosomal protein L27: Rv2441c
−2.27 50S ribosomal protein L20: Rv1643
−2.06 hypothetical protein: Rv3908
−1.76 nicotinic acid mononucleotide adenyltransferase: Rv2421c
−1.60 PROBABLE LIPASE/ESTERASE LIPG: Rv0646c
−1.44 dehydratase gmdA (GDP−mannose 4,6 dehydratase) (GMD): Rv1511
NA ORF08307
string
operons