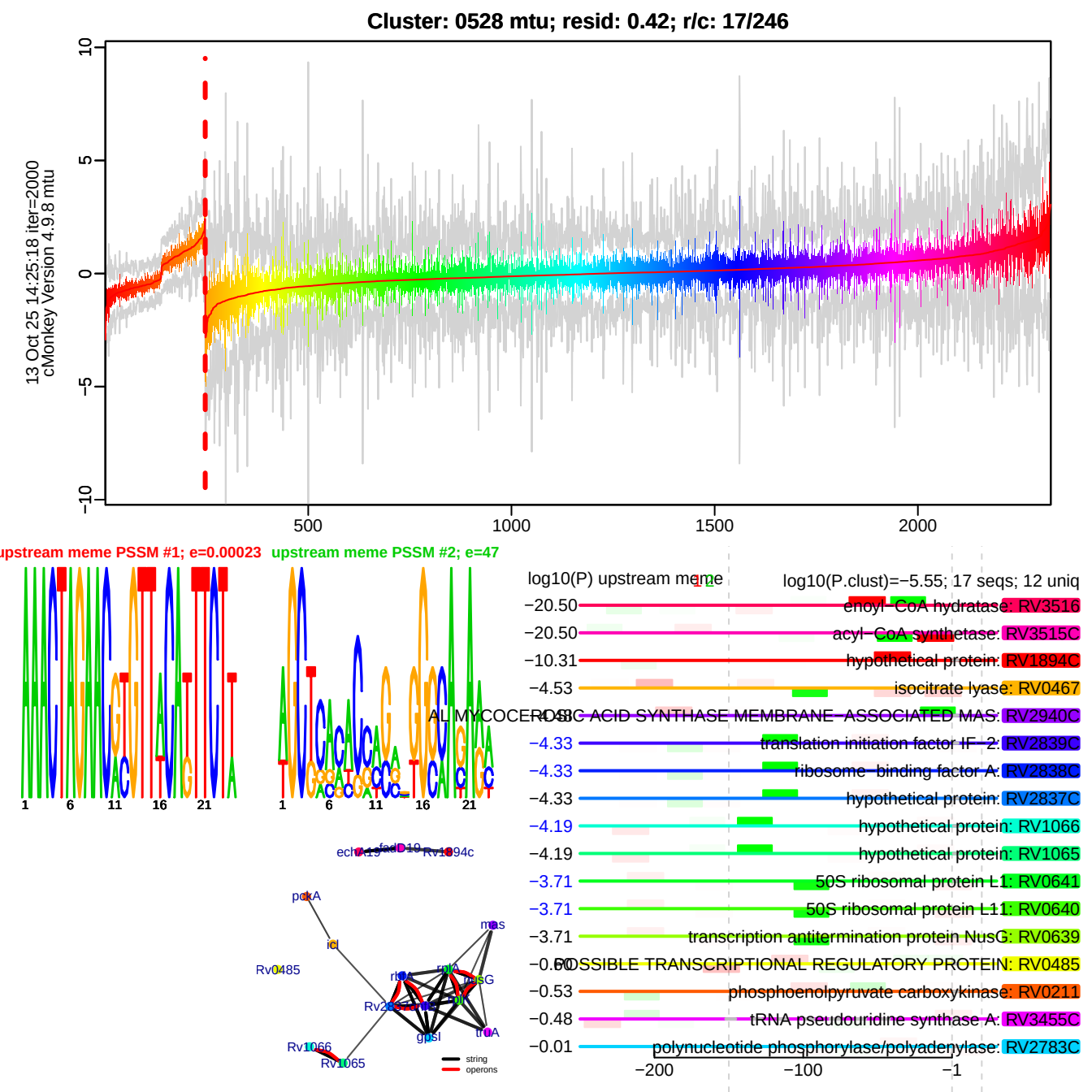
Cluster: 0528 mtu; resid: 0.42; r/c: 17/246
13 Oct 25 14:25:18 iter=2000
cMonkey Version 4.9.8 mtu
upstream meme PSSM #1; e=0.00023
upstream meme PSSM #2; e=47
log10(P) upstream meme
log10(P.clust)=−5.55; 17 seqs; 12 uniq
−20.50 enoyl−CoA hydratase: RV3516
−20.50 acyl−CoA synthetase: RV3515C
−10.31 hypothetical protein: RV1894C
−4.53 isocitrate lyase: RV0467
−4.33 MYCOCEROSIC ACID SYNTHASE MEMBRANE−ASSOCIATED MAS: RV2940C
−4.33 translation initiation factor IF−2: RV2839C
−4.33 ribosome−binding factor A: RV2838C
−4.33 hypothetical protein: RV2837C
−4.19 hypothetical protein: RV1066
−4.19 hypothetical protein: RV1065
−3.71 50S ribosomal protein L1: RV0641
−3.71 50S ribosomal protein L11: RV0640
−3.71 transcription antitermination protein NusG: RV0639
−0.60 POSSIBLE TRANSCRIPTIONAL REGULATORY PROTEIN: RV0485
−0.53 phosphoenolpyruvate carboxykinase: RV0211
−0.48 tRNA pseudouridine synthase A: RV3455C
−0.01 polynucleotide phosphorylase/polyadenylase: RV2783C
string
operons