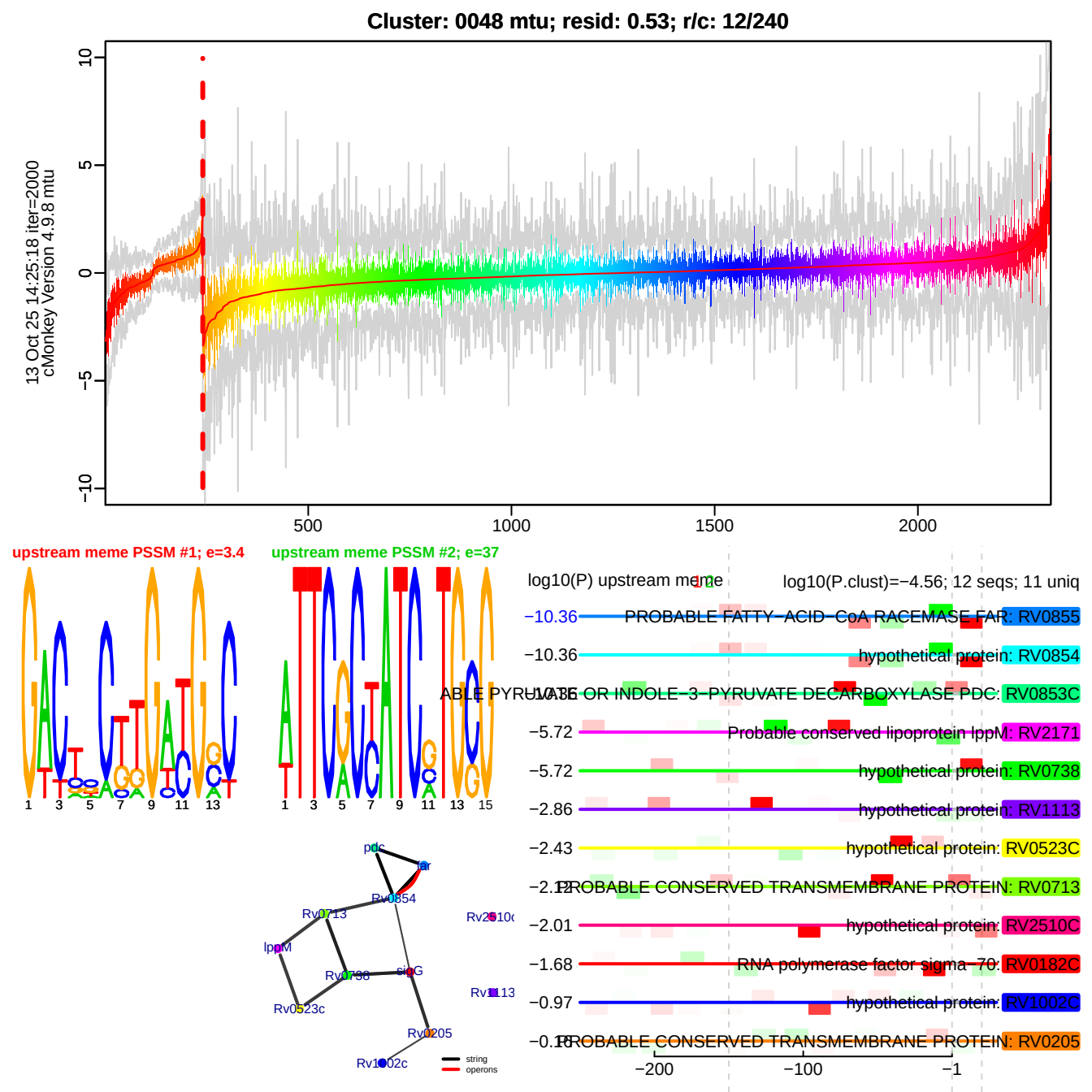
Cluster: 0048 mtu; resid: 0.53; r/c: 12/240
13 Oct 25 14:25:18 iter=2000
cMonkey Version 4.9.8 mtu
upstream meme PSSM #1; e=3.4
upstream meme PSSM #2; e=37
log10(P) upstream meme
log10(P.clust)=−4.56; 12 seqs; 11 uniq
−10.36 PROBABLE FATTY−ACID−CoA RACEMASE FAR: RV0855
−10.36 hypothetical protein: RV0854
ABLE PYRUVATE OR INDOLE−3−PYRUVATE DECARBOXYLASE PDC: RV0853C
−5.72 Probable conserved lipoprotein lppM: RV2171
−5.72 hypothetical protein: RV0738
−2.86 hypothetical protein: RV1113
−2.43 hypothetical protein: RV0523C
PROBABLE CONSERVED TRANSMEMBRANE PROTEIN: RV0713
−2.01 hypothetical protein: RV2510C
−1.68 RNA polymerase factor sigma−70: RV0182C
−0.97 hypothetical protein: RV1002C
PROBABLE CONSERVED TRANSMEMBRANE PROTEIN: RV0205
string
operons