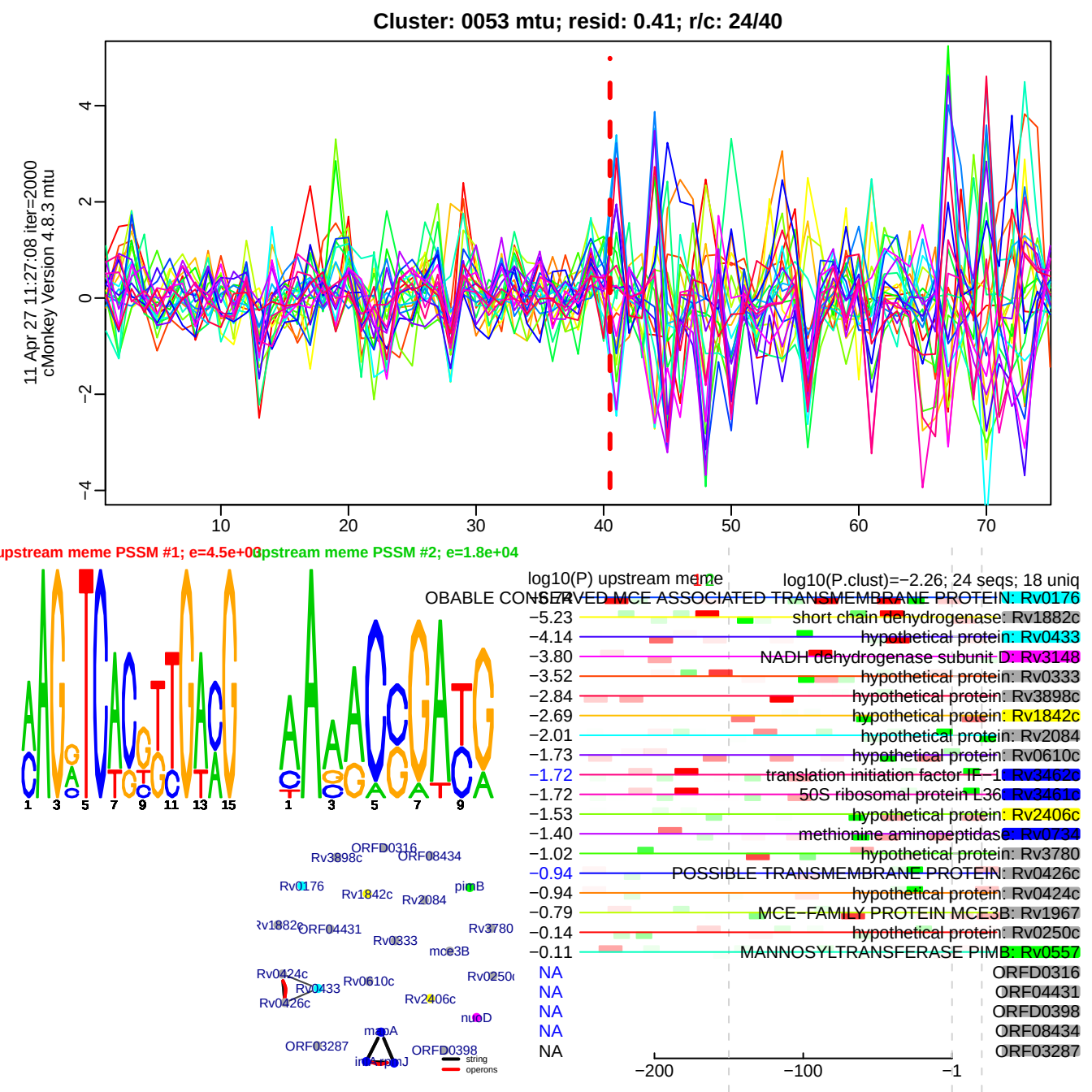
Cluster: 0053 mtu; resid: 0.41; r/c: 24/40
11 Apr 27 11:27:08 iter=2000
cMonkey Version 4.8.3 mtu
upstream meme PSSM #1; e=4.5e+03
upstream meme PSSM #2; e=1.8e+04
log10(P) upstream meme
log10(P.clust)=−2.26; 24 seqs; 18 uniq
−5.72 OBABLE CONSERVED MCE ASSOCIATED TRANSMEMBRANE PROTEIN: Rv0176
−5.23 short chain dehydrogenase: Rv1882c
−4.14 hypothetical protein: Rv0433
−3.80 NADH dehydrogenase subunit D: Rv3148
−3.52 hypothetical protein: Rv0333
−2.84 hypothetical protein: Rv3898c
−2.69 hypothetical protein: Rv1842c
−2.01 hypothetical protein: Rv2084
−1.73 hypothetical protein: Rv0610c
−1.72 translation initiation factor IF−1: Rv3462c
−1.72 50S ribosomal protein L36: Rv3461c
−1.53 hypothetical protein: Rv2406c
−1.40 methionine aminopeptidase: Rv0734
−1.02 hypothetical protein: Rv3780
−0.94 POSSIBLE TRANSMEMBRANE PROTEIN: Rv0426c
−0.94 hypothetical protein: Rv0424c
−0.79 MCE−FAMILY PROTEIN MCE3B: Rv1967
−0.14 hypothetical protein: Rv0250c
−0.11 MANNOSYLTRANSFERASE PIMB: Rv0557
NA ORFD0316
NA ORF04431
NA ORFD0398
NA ORF08434
NA ORF03287
−200
−100
−1
string
operons