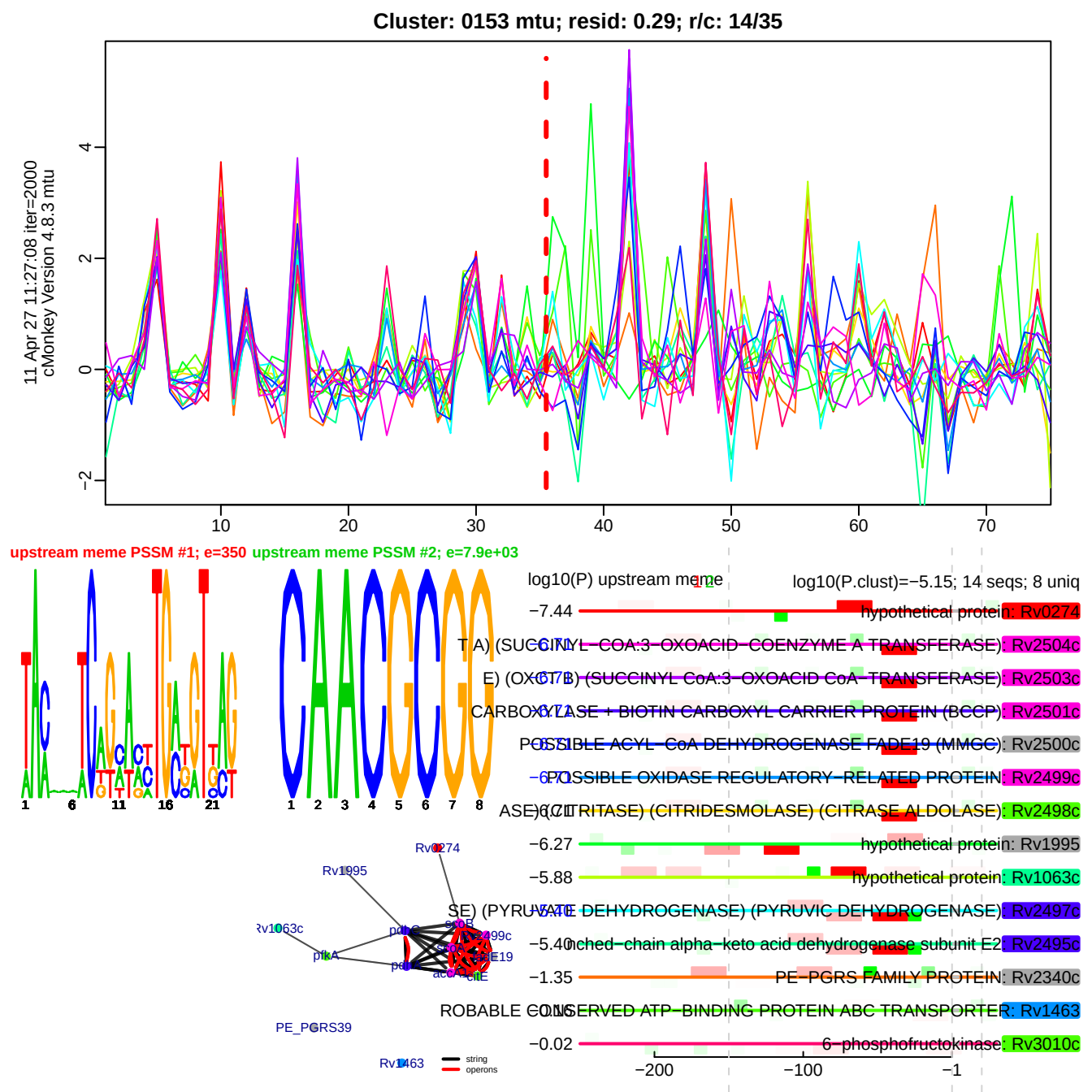
Cluster: 0153 mtu; resid: 0.29; r/c: 14/35
11 Apr 27 11:27:08 iter=2000
cMonkey Version 4.8.3 mtu
upstream meme PSSM #1; e=350
upstream meme PSSM #2; e=7.9e+03
log10(P) upstream meme
log10(P.clust)=−5.15; 14 seqs; 8 uniq
−7.44
hypothetical protein: Rv0274
−6.27
hypothetical protein: Rv1995
−5.88
hypothetical protein: Rv1063c
−5.40
branched−chain alpha−keto acid dehydrogenase subunit E2: Rv2495c
−1.35
PE−PGRS FAMILY PROTEIN: Rv2340c
−0.02
6−phosphofructokinase: Rv3010c
−200
−100
−1
Rv0274
Rv1995
Rv1063c
pfkA
PE_PGRS39
Rv1463
string
operons