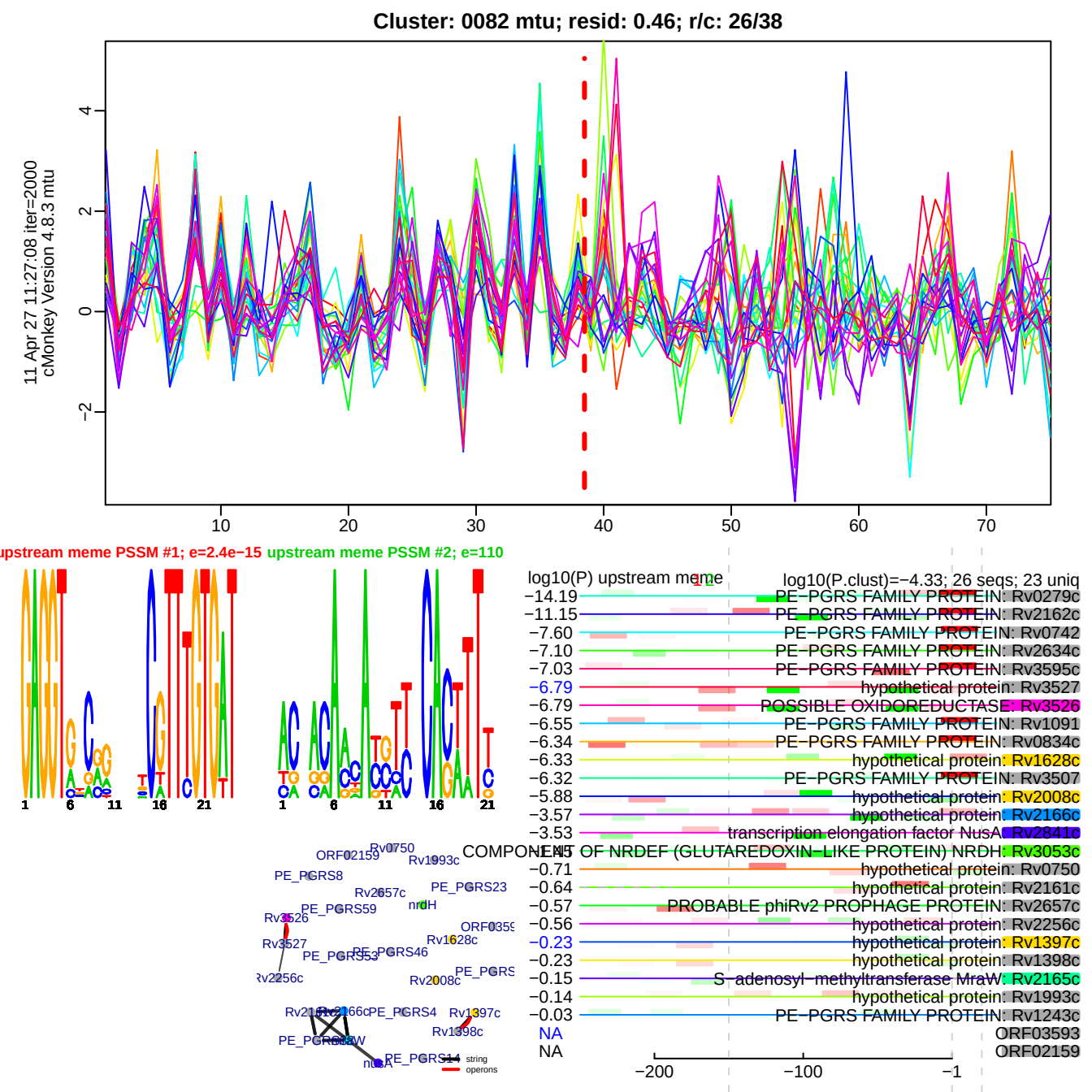
Cluster: 0082 mtu; resid: 0.46; r/c: 26/38
11 Apr 27 11:27:08 iter=2000
cMonkey Version 4.8.3 mtu
upstream meme PSSM #1; e=2.4e−15
upstream meme PSSM #2; e=110
log10(P) upstream meme
log10(P.clust)=−4.33; 26 seqs; 23 uniq
−14.19 PE−PGRS FAMILY PROTEIN: Rv0279c
−11.15 PE−PGRS FAMILY PROTEIN: Rv2162c
−7.60 PE−PGRS FAMILY PROTEIN: Rv0742
−7.10 PE−PGRS FAMILY PROTEIN: Rv2634c
−7.03 PE−PGRS FAMILY PROTEIN: Rv3595c
−6.79 hypothetical protein: Rv3527
−6.79 POSSIBLE OXIDOREDUCTASE: Rv3526
−6.55 PE−PGRS FAMILY PROTEIN: Rv1091
−6.34 PE−PGRS FAMILY PROTEIN: Rv0834c
−6.33 hypothetical protein: Rv1628c
−6.32 PE−PGRS FAMILY PROTEIN: Rv3507
−5.88 hypothetical protein: Rv2008c
−3.57 hypothetical protein: Rv2166c
−3.53 transcription elongation factor NusA: Rv2841c
−1.45 COMPONENT OF NRDEF (GLUTAREDOXIN−LIKE PROTEIN) NRDH: Rv3053c
−0.71 hypothetical protein: Rv0750
−0.64 hypothetical protein: Rv2161c
−0.57 PROBABLE phiRv2 PROPHAGE PROTEIN: Rv2657c
−0.56 hypothetical protein: Rv2256c
−0.23 hypothetical protein: Rv1397c
−0.23 hypothetical protein: Rv1398c
−0.15 S−adenosyl−methyltransferase MraW: Rv2165c
−0.14 hypothetical protein: Rv1993c
−0.03 PE−PGRS FAMILY PROTEIN: Rv1243c
NA ORF03593
NA ORF02159
−200
−100
−1
string
operons