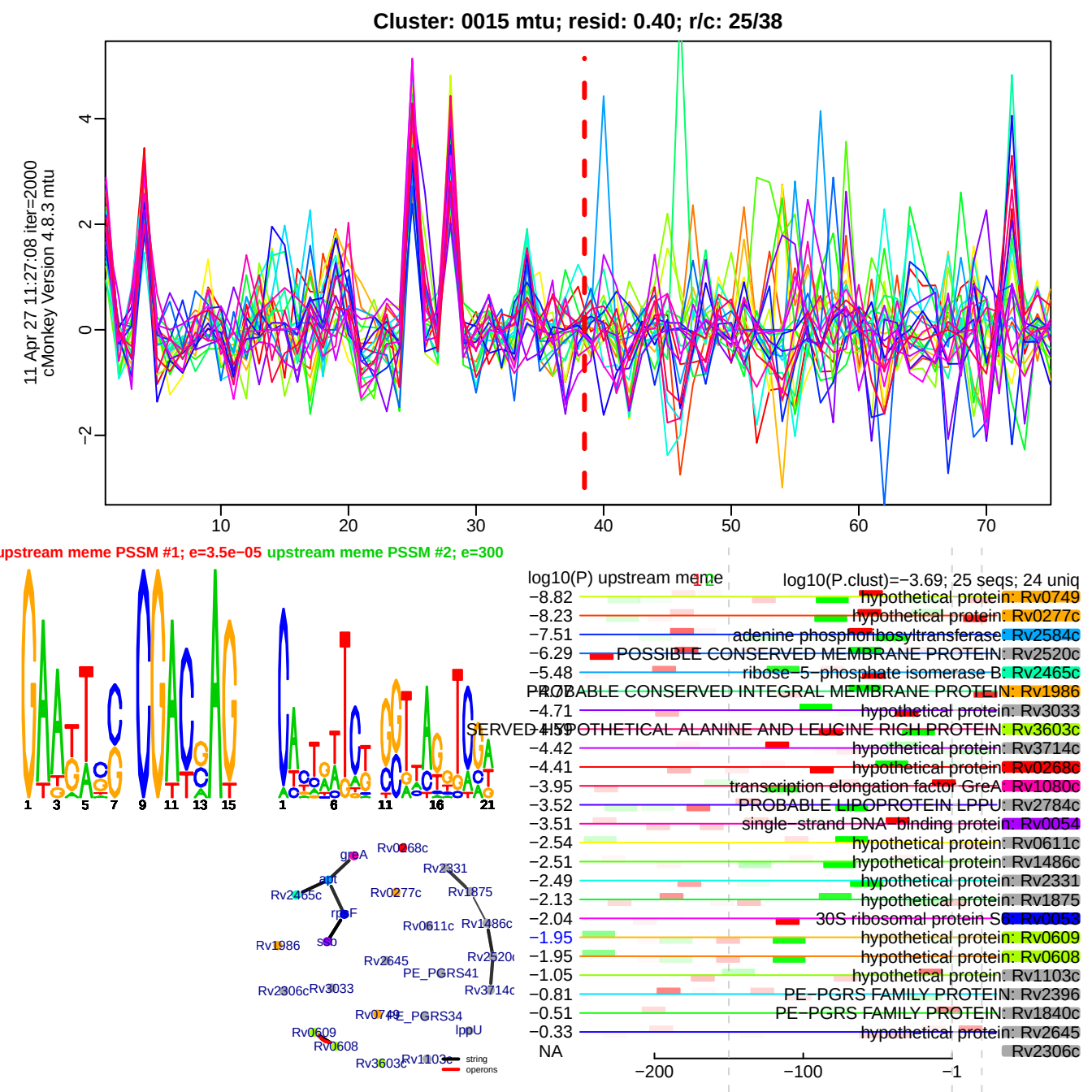
Cluster: 0015 mtu; resid: 0.40; r/c: 25/38
11 Apr 27 11:27:08 iter=2000
cMonkey Version 4.8.3 mtu
upstream meme PSSM #1; e=3.5e−05
upstream meme PSSM #2; e=300
log10(P) upstream meme
log10(P.clust)=−3.69; 25 seqs; 24 uniq
−8.82 hypothetical protein: Rv0749
−8.23 hypothetical protein: Rv0277c
−7.51 adenine phosphoribosyltransferase: Rv2584c
−6.29 POSSIBLE CONSERVED MEMBRANE PROTEIN: Rv2520c
−5.48 ribose−5−phosphate isomerase B: Rv2465c
−4.73 PROBABLE CONSERVED INTEGRAL MEMBRANE PROTEIN: Rv1986
−4.71 hypothetical protein: Rv3033
−4.59 HYPOTHETICAL ALANINE AND LEUCINE RICH PROTEIN: Rv3603c
−4.42 hypothetical protein: Rv3714c
−4.41 hypothetical protein: Rv0268c
−3.95 transcription elongation factor GreA: Rv1080c
−3.52 PROBABLE LIPOPROTEIN LPPU: Rv2784c
−3.51 single−strand DNA−binding protein: Rv0054
−2.54 hypothetical protein: Rv0611c
−2.51 hypothetical protein: Rv1486c
−2.49 hypothetical protein: Rv2331
−2.13 hypothetical protein: Rv1875
−2.04 30S ribosomal protein S6: Rv0053
−1.95 hypothetical protein: Rv0609
−1.95 hypothetical protein: Rv0608
−1.05 hypothetical protein: Rv1103c
−0.81 PE−PGRS FAMILY PROTEIN: Rv2396
−0.51 PE−PGRS FAMILY PROTEIN: Rv1840c
−0.33 hypothetical protein: Rv2645
NA Rv2306c
−200 −100 −1
greA
Rv0268c
Rv2331
apt
Rv2465c
Rv0277c
Rv1875
rpsF
Rv0611c
Rv1486c
Rv1986
ssb
Rv2645
Rv2520c
PE_PGRS41
Rv2306c
Rv3033
Rv3714c
Rv0749
PE_PGRS34
Rv0609
lppU
Rv0608
Rv3603c
Rv1103c
string
operons