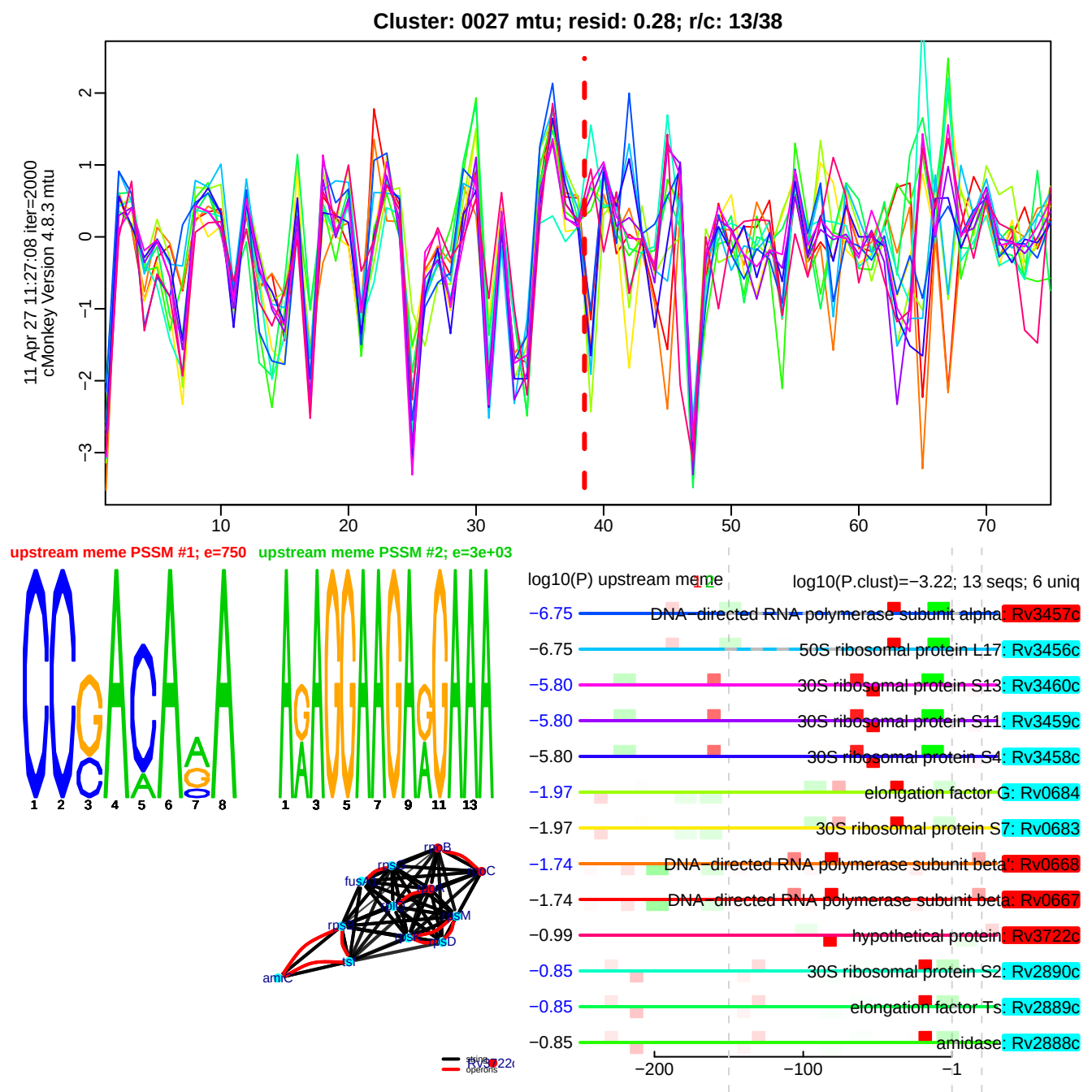
Cluster: 0027 mtu; resid: 0.28; r/c: 13/38
11 Apr 27 11:27:08 iter=2000
cMonkey Version 4.8.3 mtu
upstream meme PSSM #1; e=750
upstream meme PSSM #2; e=3e+03
log10(P) upstream meme
log10(P.clust)=−3.22; 13 seqs; 6 uniq
−6.75 DNA−directed RNA polymerase subunit alpha: Rv3457c
−6.75 50S ribosomal protein L17: Rv3456c
−5.80 30S ribosomal protein S13: Rv3460c
−5.80 30S ribosomal protein S11: Rv3459c
−5.80 30S ribosomal protein S4: Rv3458c
−1.97 elongation factor G: Rv0684
−1.97 30S ribosomal protein S7: Rv0683
−1.74 DNA−directed RNA polymerase subunit beta': Rv0668
−1.74 DNA−directed RNA polymerase subunit beta: Rv0667
−0.99 hypothetical protein: Rv3722c
−0.85 30S ribosomal protein S2: Rv2890c
−0.85 elongation factor Ts: Rv2889c
−0.85 amidase: Rv2888c
rpoB
rpoC
fusA
rplL
rplJ
rpsM
rpsL
rpsG
tsf
amiC
−200
−100
−1