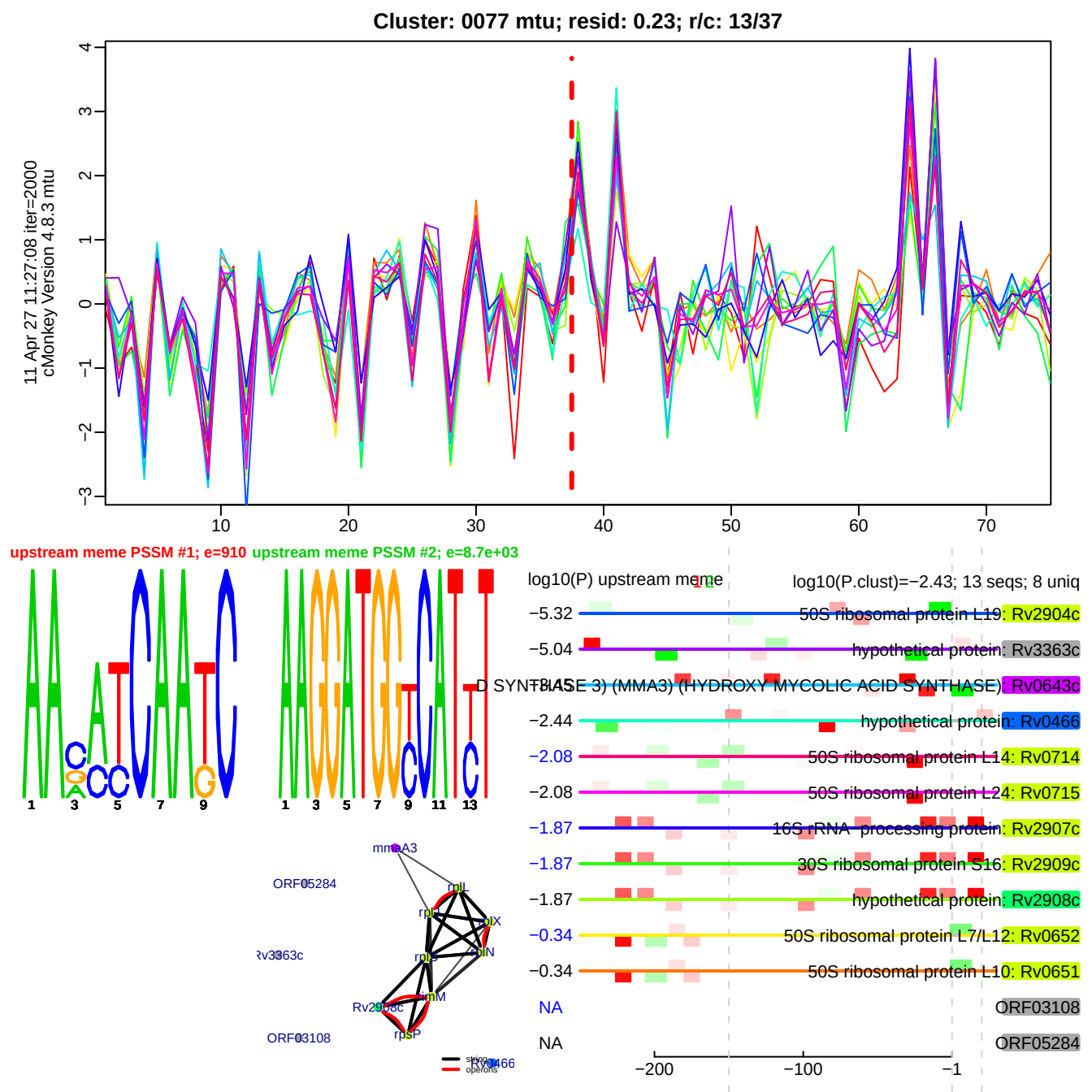

Cluster: 0077 mtu; resid: 0.23; r/c: 13/37
11 Apr 27 11:27:08 iter=2000
cMonkey Version 4.8.3 mtu
upstream meme PSSM #1; e=910
upstream meme PSSM #2; e=8.7e+03
log10(P) upstream meme
log10(P.clust)=−2.43; 13 seqs; 8 uniq
−5.32 50S ribosomal protein L19: Rv2904c
−5.04 hypothetical protein: Rv3363c
D SYNTHASE 3) (MMA3) (HYDROXY MYCOLIC ACID SYNTHASE): Rv0643c
−2.44 hypothetical protein: Rv0466
−2.08 50S ribosomal protein L14: Rv0714
−2.08 50S ribosomal protein L24: Rv0715
−1.87 16S rRNA processing protein: Rv2907c
−1.87 30S ribosomal protein S16: Rv2909c
−1.87 hypothetical protein: Rv2908c
−0.34 50S ribosomal protein L7/L12: Rv0652
−0.34 50S ribosomal protein L10: Rv0651
NA ORF03108
NA ORF05284
mmaA3
ORF05284
rplL
rplI
rplX
rplN
rplS
Rv3363c
rimM
Rv2908c
rpsP
ORF03108
Rv0466
string
operons