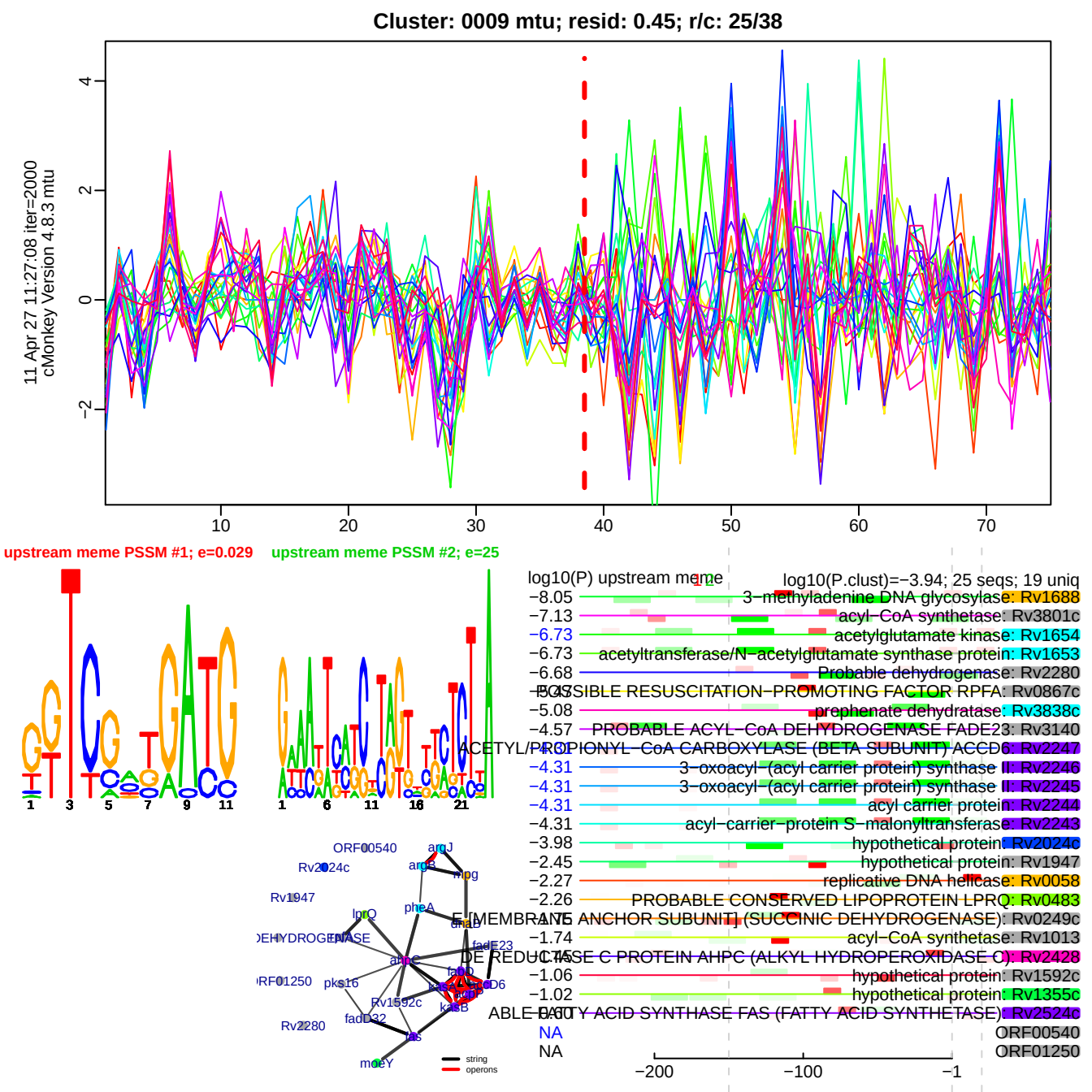
Cluster: 0009 mtu; resid: 0.45; r/c: 25/38
11 Apr 27 11:27:08 iter=2000
cMonkey Version 4.8.3 mtu
upstream meme PSSM #1; e=0.029
upstream meme PSSM #2; e=25
log10(P) upstream meme
log10(P.clust)=−3.94; 25 seqs; 19 uniq
−8.05 3−methyladenine DNA glycosylase: Rv1688
−7.13 acyl−CoA synthetase: Rv3801c
−6.73 acetylglutamate kinase: Rv1654
−6.73 acetyltransferase/N−acetylglutamate synthase protein: Rv1653
−6.68 Probable dehydrogenase: Rv2280
−5.47 POSSIBLE RESUSCITATION−PROMOTING FACTOR RPFA: Rv0867c
−5.08 prephenate dehydratase: Rv3838c
−4.57 PROBABLE ACYL−CoA DEHYDROGENASE FADE23: Rv3140
−4.31 ACETYL/PROPIONYL−CoA CARBOXYLASE (BETA SUBUNIT) ACCD6: Rv2247
−4.31 3−oxoacyl−(acyl carrier protein) synthase II: Rv2246
−4.31 3−oxoacyl−(acyl carrier protein) synthase II: Rv2245
−4.31 acyl carrier protein: Rv2244
−4.31 acyl−carrier−protein S−malonyltransferase: Rv2243
−3.98 hypothetical protein: Rv2024c
−2.45 hypothetical protein: Rv1947
−2.27 replicative DNA helicase: Rv0058
−2.26 PROBABLE CONSERVED LIPOPROTEIN LPRQ: Rv0483
−1.75 F [MEMBRANE ANCHOR SUBUNIT] (SUCCINIC DEHYDROGENASE): Rv0249c
−1.74 acyl−CoA synthetase: Rv1013
−1.45 DE REDUCTASE C PROTEIN AHPC (ALKYL HYDROPEROXIDASE C): Rv2428
−1.06 hypothetical protein: Rv1592c
−1.02 hypothetical protein: Rv1355c
ABLE FATTY ACID SYNTHASE FAS (FATTY ACID SYNTHETASE): Rv2524c
NA
NA
ORF00540
ORF01250
string
operons