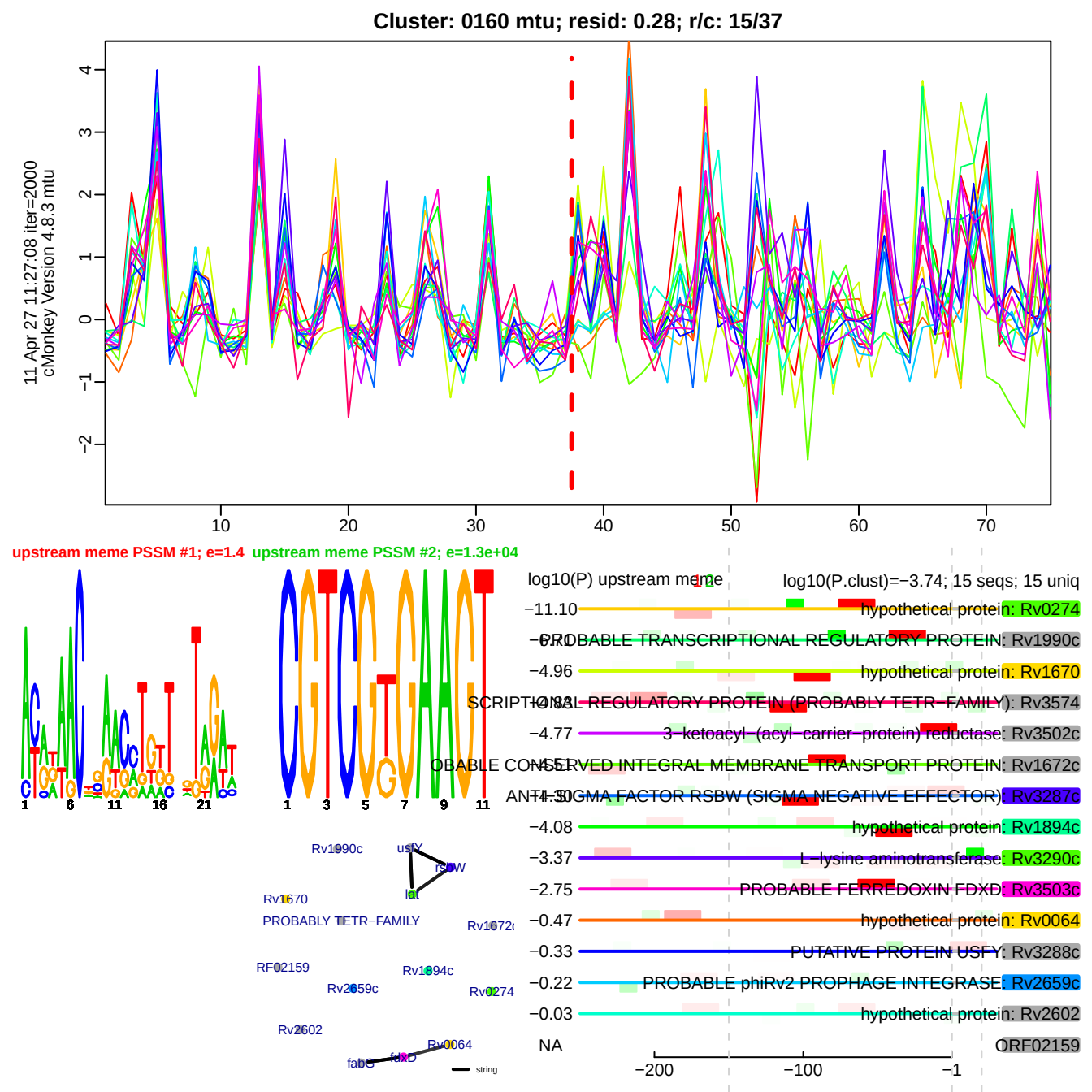
Cluster: 0160 mtu; resid: 0.28; r/c: 15/37
11 Apr 27 11:27:08 iter=2000
cMonkey Version 4.8.3 mtu
upstream meme PSSM #1; e=1.4
upstream meme PSSM #2; e=1.3e+04
log10(P) upstream meme
log10(P.clust)=−3.74; 15 seqs; 15 uniq
−11.10 hypothetical protein: Rv0274
−6.70 PROBABLE TRANSCRIPTIONAL REGULATORY PROTEIN: Rv1990c
−4.96 hypothetical protein: Rv1670
−4.83 TRANSCRIPTIONAL REGULATORY PROTEIN (PROBABLY TETR−FAMILY): Rv3574
−4.77 3−ketoacyl−(acyl−carrier−protein) reductase: Rv3502c
−4.55 PROBABLE CONSERVED INTEGRAL MEMBRANE TRANSPORT PROTEIN: Rv1672c
−4.30 ANTI−SIGMA FACTOR RSBW (SIGMA NEGATIVE EFFECTOR): Rv3287c
−4.08 hypothetical protein: Rv1894c
−3.37 L−lysine aminotransferase: Rv3290c
−2.75 PROBABLE FERREDOXIN FDXD: Rv3503c
−0.47 hypothetical protein: Rv0064
−0.33 PUTATIVE PROTEIN USFY: Rv3288c
−0.22 PROBABLE phiRv2 PROPHAGE INTEGRASE: Rv2659c
−0.03 hypothetical protein: Rv2602
NA ORF02159
−200 −100 −1
Rv1990c
usfY
rsbW
lat
Rv1670
PROBABLY TETR−FAMILY
Rv1672c
RF02159
Rv1894c
Rv2659c
Rv0274
Rv2602
Rv0064
fabG
fdxD
string