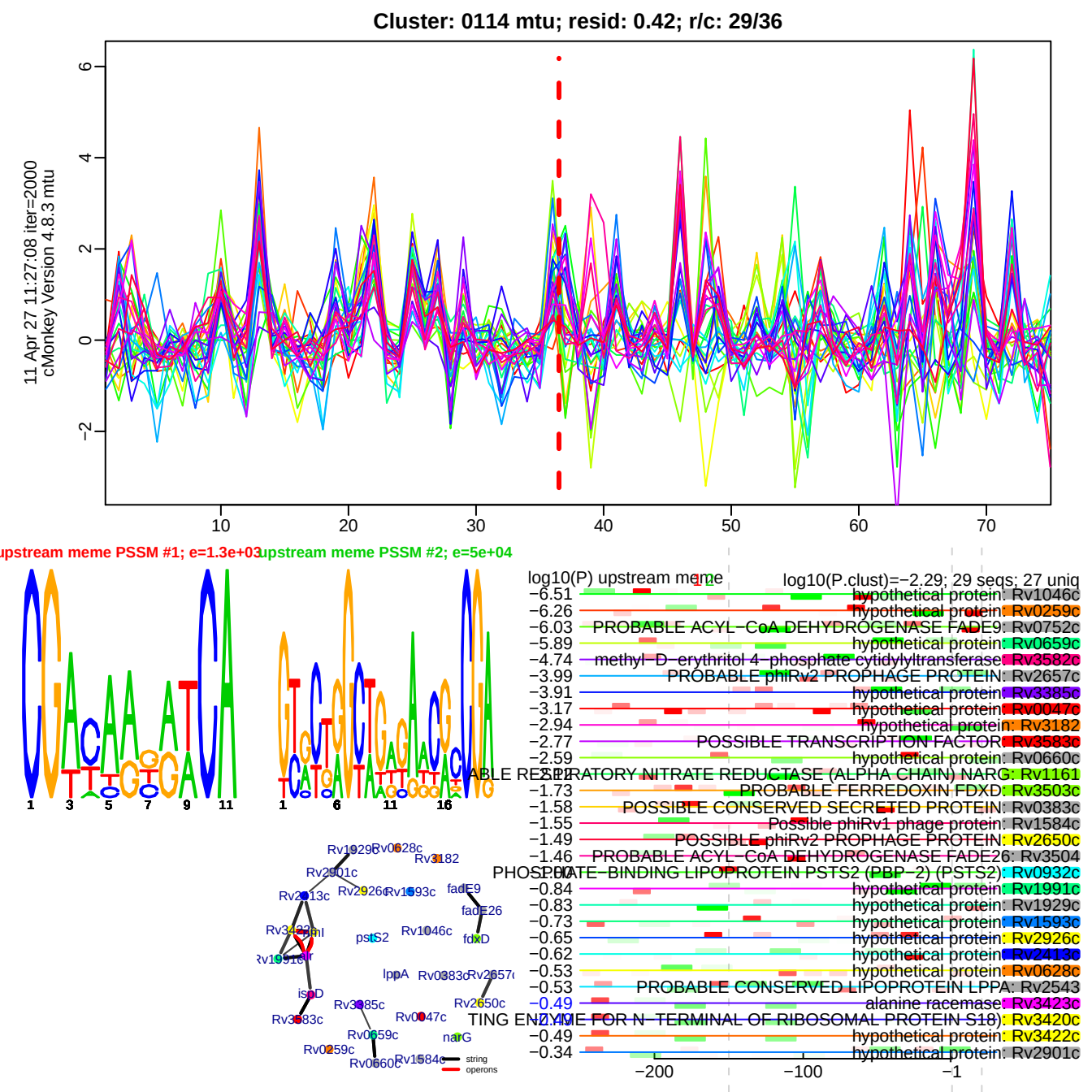
Cluster: 0114 mtu; resid: 0.42; r/c: 29/36
11 Apr 27 11:27:08 iter=2000
cMonkey Version 4.8.3 mtu
upstream meme PSSM #1; e=1.3e+03
upstream meme PSSM #2; e=5e+04
log10(P) upstream meme
log10(P.clust)=−2.29; 29 seqs; 27 uniq
−6.51 hypothetical protein: Rv1046c
−6.26 hypothetical protein: Rv0259c
−6.03 PROBABLE ACYL−CoA DEHYDROGENASE FADE9: Rv0752c
−5.89 hypothetical protein: Rv0659c
−4.74 methyl−D−erythritol 4−phosphate cytidylyltransferase: Rv3582c
−3.99 PROBABLE phiRv2 PROPHAGE PROTEIN: Rv2657c
−3.91 hypothetical protein: Rv3385c
−3.17 hypothetical protein: Rv0047c
−2.94 hypothetical protein: Rv3182
−2.77 POSSIBLE TRANSCRIPTION FACTOR: Rv3583c
−2.59 hypothetical protein: Rv0660c
−2.29 PROBABLE RESPIRATORY NITRATE REDUCTASE (ALPHA CHAIN) NARG: Rv1161
−1.73 PROBABLE FERREDOXIN FDXD: Rv3503c
−1.58 POSSIBLE CONSERVED SECRETED PROTEIN: Rv0383c
−1.55 Possible phiRv1 phage protein: Rv1584c
−1.49 POSSIBLE phiRv2 PROPHAGE PROTEIN: Rv2650c
−1.46 PROBABLE ACYL−CoA DEHYDROGENASE FADE26: Rv3504
−1.14 PHOSPHATE−BINDING LIPOPROTEIN PSTS2 (PBP−2) (PSTS2): Rv0932c
−0.84 hypothetical protein: Rv1991c
−0.83 hypothetical protein: Rv1929c
−0.73 hypothetical protein: Rv1593c
−0.65 hypothetical protein: Rv2926c
−0.62 hypothetical protein: Rv2413c
−0.53 hypothetical protein: Rv0628c
−0.53 PROBABLE CONSERVED LIPOPROTEIN LPPA: Rv2543
−0.49 alanine racemase: Rv3423c
−0.49 TING ENZYME FOR N−TERMINAL OF RIBOSOMAL PROTEIN S18): Rv3420c
−0.49 hypothetical protein: Rv3422c
−0.34 hypothetical protein: Rv2901c
−200
−100
−1
Rv1929c
Rv0628c
Rv3182
Rv2901c
Rv2926c
Rv1993c
fadE9
Rv2413c
fadE26
Rv3422c
phiI
pstS2
Rv1046c
fdxD
Rv1991c
lppA
Rv0383c
Rv2657c
isoD
Rv3385c
Rv2650c
Rv3383c
Rv0047c
Rv0659c
narG
Rv0259c
Rv0660c
Rv1584c
string
operons