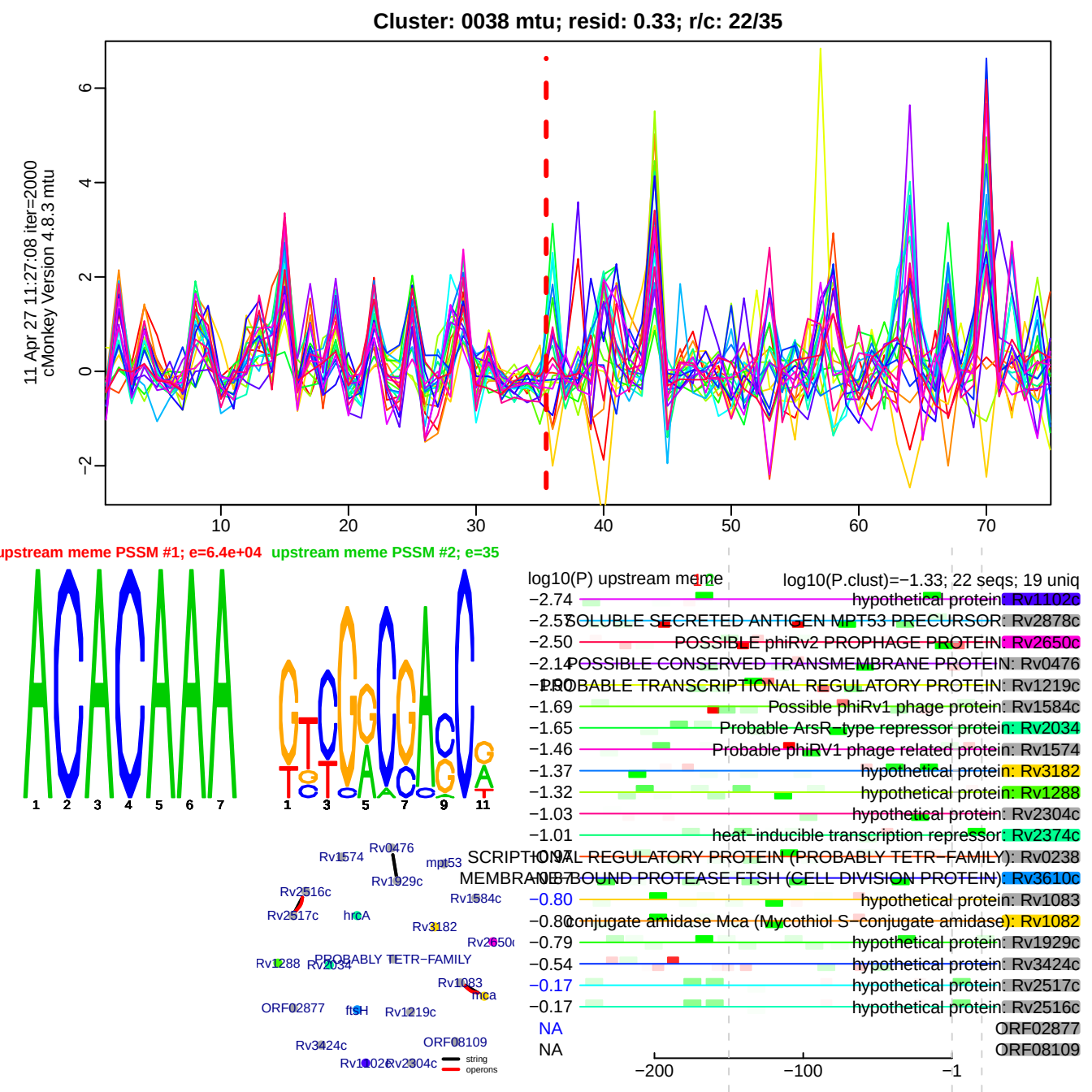
Cluster: 0038 mtu; resid: 0.33; r/c: 22/35
11 Apr 27 11:27:08 iter=2000
cMonkey Version 4.8.3 mtu
upstream meme PSSM #1; e=6.4e+04
upstream meme PSSM #2; e=35
log10(P) upstream meme
log10(P.clust)=−1.33; 22 seqs; 19 uniq
−2.74 hypothetical protein: Rv1102c
−2.57 SOLUBLE SECRETED ANTIGEN MPT53 PRECURSOR: Rv2878c
−2.50 POSSIBLE phiRv2 PROPHAGE PROTEIN: Rv2650c
−2.14 POSSIBLE CONSERVED TRANSMEMBRANE PROTEIN: Rv0476
−1.80 PROBABLE TRANSCRIPTIONAL REGULATORY PROTEIN: Rv1219c
−1.69 Possible phiRv1 phage protein: Rv1584c
−1.65 Probable ArsR−type repressor protein: Rv2034
−1.46 Probable phiRV1 phage related protein: Rv1574
−1.37 hypothetical protein: Rv3182
−1.32 hypothetical protein: Rv1288
−1.03 hypothetical protein: Rv2304c
−1.01 heat−inducible transcription repressor: Rv2374c
−0.97 TRANSCRIPTIONAL REGULATORY PROTEIN (PROBABLY TETR−FAMILY): Rv0238
−0.87 MEMBRANE−BOUND PROTEASE FTSH (CELL DIVISION PROTEIN): Rv3610c
−0.80 hypothetical protein: Rv1083
−0.80 conjugate amidase Mca (Mycothiol S−conjugate amidase): Rv1082
−0.79 hypothetical protein: Rv1929c
−0.54 hypothetical protein: Rv3424c
−0.17 hypothetical protein: Rv2517c
−0.17 hypothetical protein: Rv2516c
NA ORF02877
NA ORF08109
Rv1574
Rv0476
mpt53
Rv1929c
Rv2516c
Rv1584c
Rv2517c
hrcA
Rv3182
Rv2650c
Rv1288
PROBABLY TETR−FAMILY
Rv2034
Rv1083
mca
ORF02877
ftsH
Rv1219c
Rv3424c
ORF08109
Rv1102c
Rv2304c
string
operons
−200
−100
−1