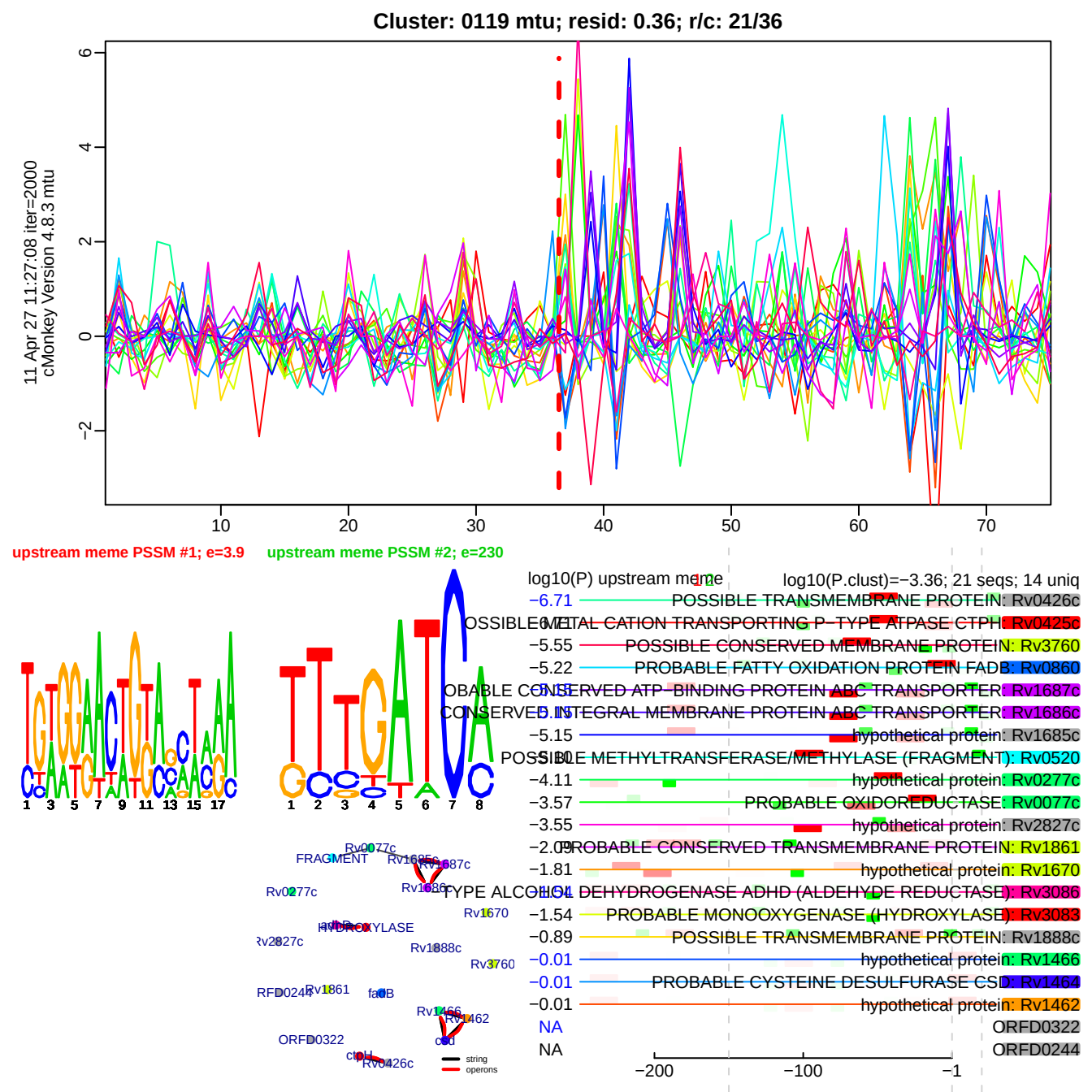
Cluster: 0119 mtu; resid: 0.36; r/c: 21/36
11 Apr 27 11:27:08 iter=2000
cMonkey Version 4.8.3 mtu
upstream meme PSSM #1; e=3.9
upstream meme PSSM #2; e=230
log10(P) upstream meme
log10(P.clust)=−3.36; 21 seqs; 14 uniq
−6.71 POSSIBLE TRANSMEMBRANE PROTEIN: Rv0426c
POSSIBLE METAL CATION TRANSPORTING P−TYPE ATPASE CTPH: Rv0425c
−5.55 POSSIBLE CONSERVED MEMBRANE PROTEIN: Rv3760
−5.22 PROBABLE FATTY OXIDATION PROTEIN FADB: Rv0860
PROBABLE CONSERVED ATP−BINDING PROTEIN ABC TRANSPORTER: Rv1687c
CONSERVED INTEGRAL MEMBRANE PROTEIN ABC TRANSPORTER: Rv1686c
−5.15 hypothetical protein: Rv1685c
POSSIBLE METHYLTRANSFERASE/METHYLASE (FRAGMENT): Rv0520
−4.11 hypothetical protein: Rv0277c
−3.57 PROBABLE OXIDOREDUCTASE: Rv0077c
−3.55 hypothetical protein: Rv2827c
−2.09 PROBABLE CONSERVED TRANSMEMBRANE PROTEIN: Rv1861
−1.81 hypothetical protein: Rv1670
PROBABLE ALCOHOL DEHYDROGENASE ADHD (ALDEHYDE REDUCTASE): Rv3086
−1.54 PROBABLE MONOOXYGENASE (HYDROXYLASE): Rv3083
−0.89 POSSIBLE TRANSMEMBRANE PROTEIN: Rv1888c
−0.01 hypothetical protein: Rv1466
−0.01 PROBABLE CYSTEINE DESULFURASE CSD: Rv1464
−0.01 hypothetical protein: Rv1462
NA ORFD0322
NA ORFD0244
−200
−100
−1
string
operons