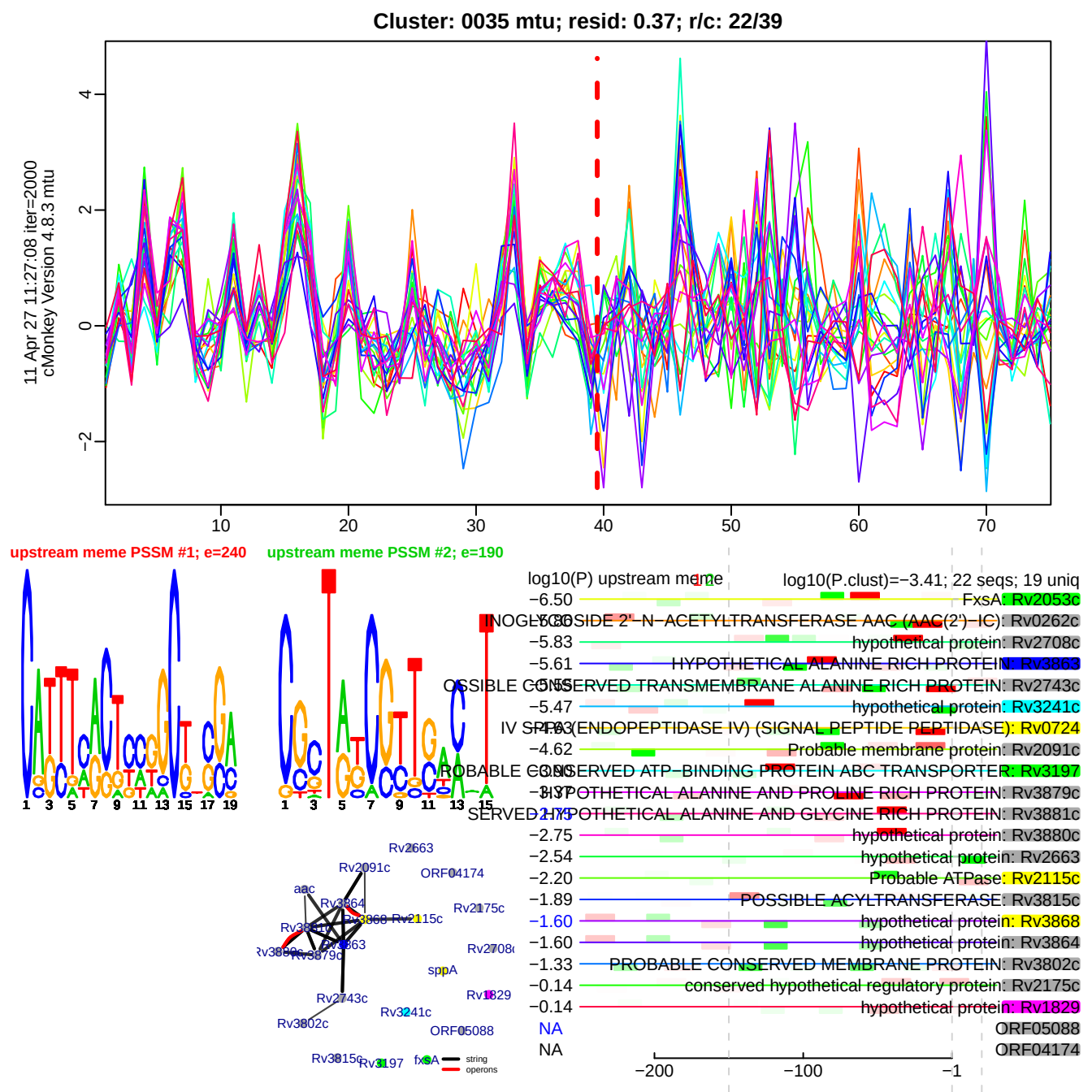
Cluster: 0035 mtu; resid: 0.37; r/c: 22/39
11 Apr 27 11:27:08 iter=2000
cMonkey Version 4.8.3 mtu
upstream meme PSSM #1; e=240
upstream meme PSSM #2; e=190
log10(P) upstream meme
log10(P.clust)=−3.41; 22 seqs; 19 uniq
−6.50 FxsA: Rv2053c
INOGLYCOSIDE 2''−N−ACETYLTRANSFERASE AAC (AAC(2')−IC): Rv0262c
−5.83 hypothetical protein: Rv2708c
−5.61 HYPOTHETICAL ALANINE RICH PROTEIN: Rv3863
OSSIBLE CONSERVED TRANSMEMBRANE ALANINE RICH PROTEIN: Rv2743c
−5.47 hypothetical protein: Rv3241c
IV SPASE (ENDOPEPTIDASE IV) (SIGNAL PEPTIDE PEPTIDASE): Rv0724
−4.62 Probable membrane protein: Rv2091c
ROBABLE CONSERVED ATP−BINDING PROTEIN ABC TRANSPORTER: Rv3197
HYPOTHETICAL ALANINE AND PROLINE RICH PROTEIN: Rv3879c
SERVED HYPOTHETICAL ALANINE AND GLYCINE RICH PROTEIN: Rv3881c
−2.75 hypothetical protein: Rv3880c
−2.54 hypothetical protein: Rv2663
−2.20 Probable ATPase: Rv2115c
−1.89 POSSIBLE ACYLTRANSFERASE: Rv3815c
−1.60 hypothetical protein: Rv3868
−1.60 hypothetical protein: Rv3864
−1.33 PROBABLE CONSERVED MEMBRANE PROTEIN: Rv3802c
−0.14 conserved hypothetical regulatory protein: Rv2175c
−0.14 hypothetical protein: Rv1829
NA ORF05088
NA ORF04174
−200 −100 −1
Rv2663
Rv2091c
ORF04174
aac
Rv3864
Rv2175c
Rv3868
Rv2115c
Rv3863
Rv2708c
Rv3879c
sppA
Rv1829
Rv2743c
Rv3241c
Rv3802c
ORF05088
Rv3815c
Rv3197
fxsA
string
operons