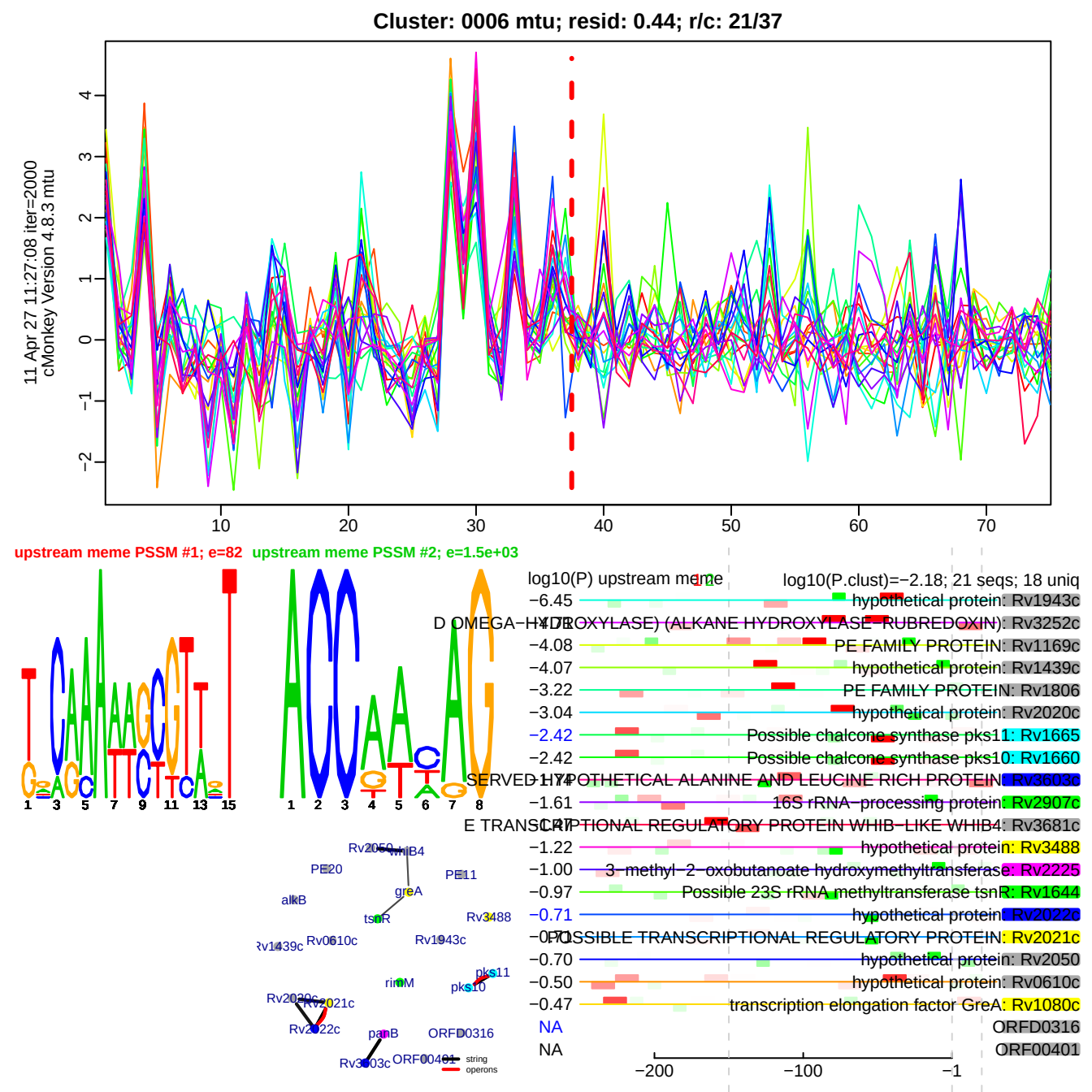
Cluster: 0006 mtu; resid: 0.44; r/c: 21/37
11 Apr 27 11:27:08 iter=2000
cMonkey Version 4.8.3 mtu
upstream meme PSSM #1; e=82
upstream meme PSSM #2; e=1.5e+03
log10(P) upstream meme
log10(P.clust)=−2.18; 21 seqs; 18 uniq
−6.45 hypothetical protein: Rv1943c
D OMEGA−HYDROXYLASE) (ALKANE HYDROXYLASE−RUBREDOXIN): Rv3252c
−4.08 PE FAMILY PROTEIN: Rv1169c
−4.07 hypothetical protein: Rv1439c
−3.22 PE FAMILY PROTEIN: Rv1806
−3.04 hypothetical protein: Rv2020c
−2.42 Possible chalcone synthase pks11: Rv1665
−2.42 Possible chalcone synthase pks10: Rv1660
SERVED HYPOTHETICAL ALANINE AND LEUCINE RICH PROTEIN: Rv3603c
−1.61 16S rRNA−processing protein: Rv2907c
E TRANSCRIPTIONAL REGULATORY PROTEIN WHIB−LIKE WHIB4: Rv3681c
−1.22 hypothetical protein: Rv3488
−1.00 3−methyl−2−oxobutanoate hydroxymethyltransferase: Rv2225
−0.97 Possible 23S rRNA methyltransferase tsnR: Rv1644
−0.71 hypothetical protein: Rv2022c
−0.70 POSSIBLE TRANSCRIPTIONAL REGULATORY PROTEIN: Rv2021c
−0.70 hypothetical protein: Rv2050
−0.50 hypothetical protein: Rv0610c
−0.47 transcription elongation factor GreA: Rv1080c
NA ORFD0316
NA ORF00401
Rv2050
whiB4
PE20
PE11
greA
alkB
tsnR
Rv3488
Rv1439c
Rv0610c
Rv1943c
pks11
rimM
pks10
Rv2020c
Rv2021c
Rv2022c
panB
ORFD0316
Rv3603c
ORF00401
string
operons