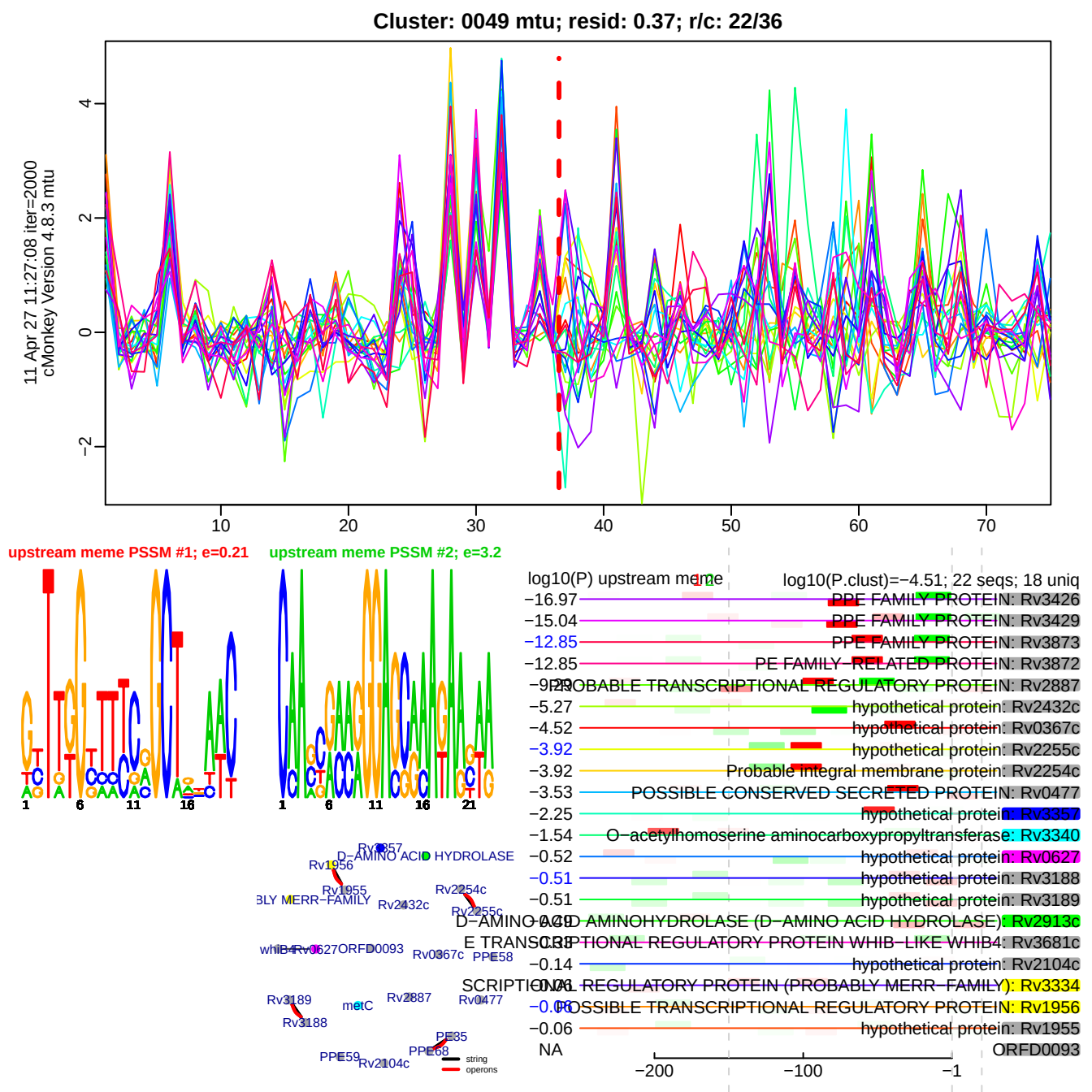
Cluster: 0049 mtu; resid: 0.37; r/c: 22/36
11 Apr 27 11:27:08 iter=2000
cMonkey Version 4.8.3 mtu
upstream meme PSSM #1; e=0.21
upstream meme PSSM #2; e=3.2
log10(P) upstream meme
log10(P.clust)=−4.51; 22 seqs; 18 uniq
−16.97 PPE FAMILY PROTEIN: Rv3426
−15.04 PPE FAMILY PROTEIN: Rv3429
−12.85 PPE FAMILY PROTEIN: Rv3873
−12.85 PE FAMILY−RELATED PROTEIN: Rv3872
−9.29 PROBABLE TRANSCRIPTIONAL REGULATORY PROTEIN: Rv2887
−5.27 hypothetical protein: Rv2432c
−4.52 hypothetical protein: Rv0367c
−3.92 hypothetical protein: Rv2255c
−3.92 Probable integral membrane protein: Rv2254c
−3.53 POSSIBLE CONSERVED SECRETED PROTEIN: Rv0477
−2.25 hypothetical protein: Rv3357
−1.54 O−acetylhomoserine aminocarboxypropyltransferase: Rv3340
−0.52 hypothetical protein: Rv0627
−0.51 hypothetical protein: Rv3188
−0.51 hypothetical protein: Rv3189
−0.38 D−AMINO ACID AMINOHYDROLASE (D−AMINO ACID HYDROLASE): Rv2913c
−0.38 TRANSCRIPTIONAL REGULATORY PROTEIN WHIB−LIKE WHIB4: Rv3681c
−0.14 hypothetical protein: Rv2104c
−0.06 TRANSCRIPTIONAL REGULATORY PROTEIN (PROBABLY MERR−FAMILY): Rv3334
−0.06 POSSIBLE TRANSCRIPTIONAL REGULATORY PROTEIN: Rv1956
−0.06 hypothetical protein: Rv1955
NA ORFD0093
−200 −100 −1
Rv0357
D−AMINO ACID HYDROLASE
Rv1956
Rv1955
Rv2254c
Rv2432c
Rv2255c
Rv0367c
PPE58
Rv3189
metC
Rv2887
Rv0477
Rv3188
PE35
PPE68
PPE59
Rv2104c
string
operons