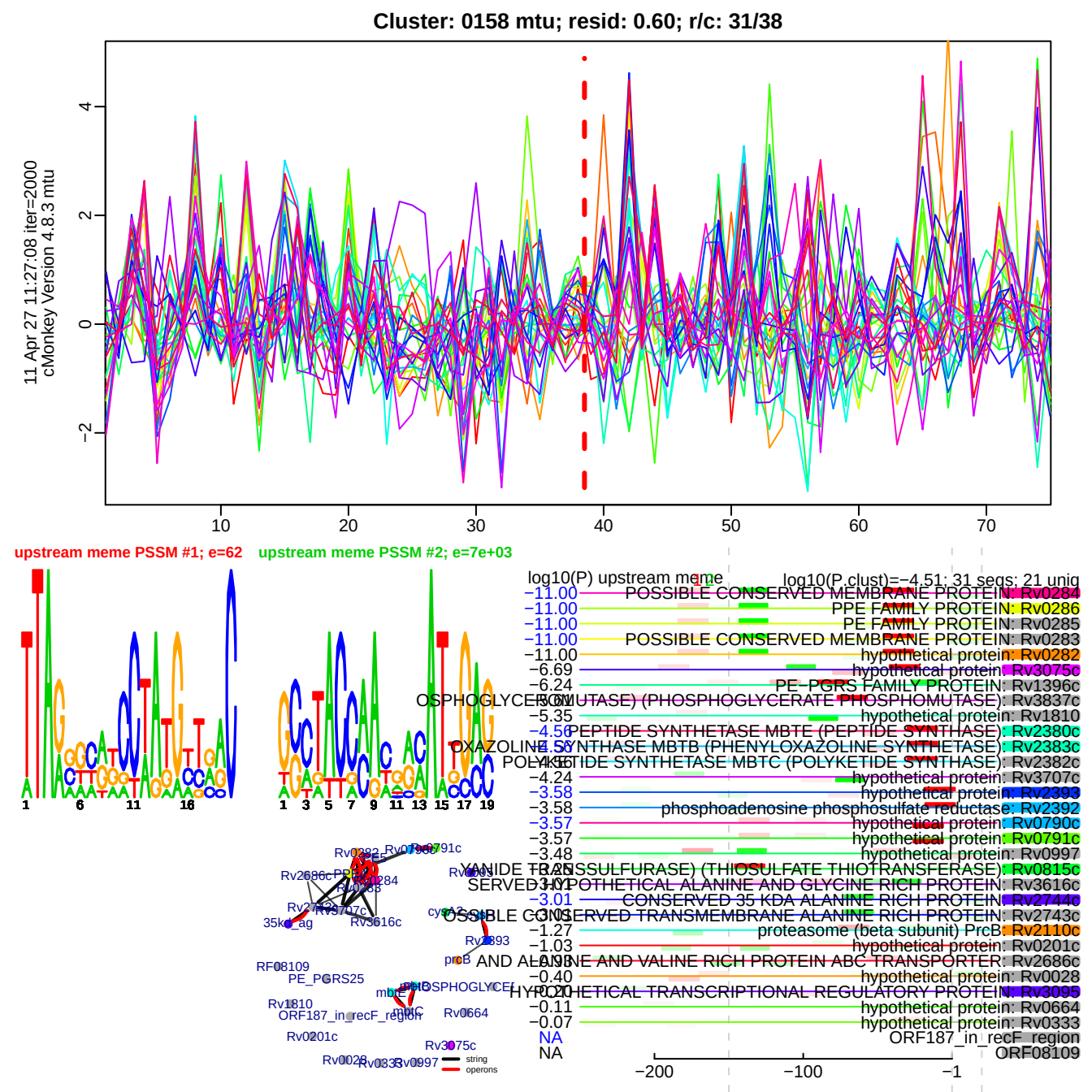
Cluster: 0158 mtu; resid: 0.60; r/c: 31/38
11 Apr 27 11:27:08 iter=2000
cMonkey Version 4.8.3 mtu
upstream meme PSSM #1; e=62
upstream meme PSSM #2; e=7e+03
log10(P) upstream meme
log10(P.clust)=−4.51; 31 seqs; 21 uniq
POSSIBLE CONSERVED MEMBRANE PROTEIN: Rv0284
PPE FAMILY PROTEIN: Rv0286
PE FAMILY PROTEIN: Rv0285
POSSIBLE CONSERVED MEMBRANE PROTEIN: Rv0283
hypothetical protein: Rv0282
hypothetical protein: Rv3075c
PE−PGRS FAMILY PROTEIN: Rv1396c
OSPHOGLYCERATE MUTASE) (PHOSPHOGLYCERATE PHOSPHOMUTASE): Rv3837c
hypothetical protein: Rv1810
PEPTIDE SYNTHETASE MBTE (PEPTIDE SYNTHASE): Rv2380c
OXAZOLINE SYNTHASE MBTB (PHENYLOXAZOLINE SYNTHETASE): Rv2383c
POLYKETIDE SYNTHETASE MBTC (POLYKETIDE SYNTHASE): Rv2382c
hypothetical protein: Rv3707c
hypothetical protein: Rv2393
phosphoadenosine phosphosulfate reductase: Rv2392
hypothetical protein: Rv0790c
hypothetical protein: Rv0791c
hypothetical protein: Rv0997
YANIDE TRANSSULFURASE) (THIOSULFATE THIOTRANSFERASE): Rv0815c
SERVED HYPOTHETICAL ALANINE AND GLYCINE RICH PROTEIN: Rv3616c
CONSERVED 35 KDA ALANINE RICH PROTEIN: Rv2744c
OSSIBLE CONSERVED TRANSMEMBRANE ALANINE RICH PROTEIN: Rv2743c
proteasome (beta subunit) PrcB: Rv2110c
hypothetical protein: Rv0201c
AND ALANINE AND VALINE RICH PROTEIN ABC TRANSPORTER: Rv2686c
hypothetical protein: Rv0028
HYPOTHETICAL TRANSCRIPTIONAL REGULATORY PROTEIN: Rv3095
hypothetical protein: Rv0664
hypothetical protein: Rv0333
ORF187_in_recF_region
ORF08109
NA
NA
−200
−100
−1
string
operons