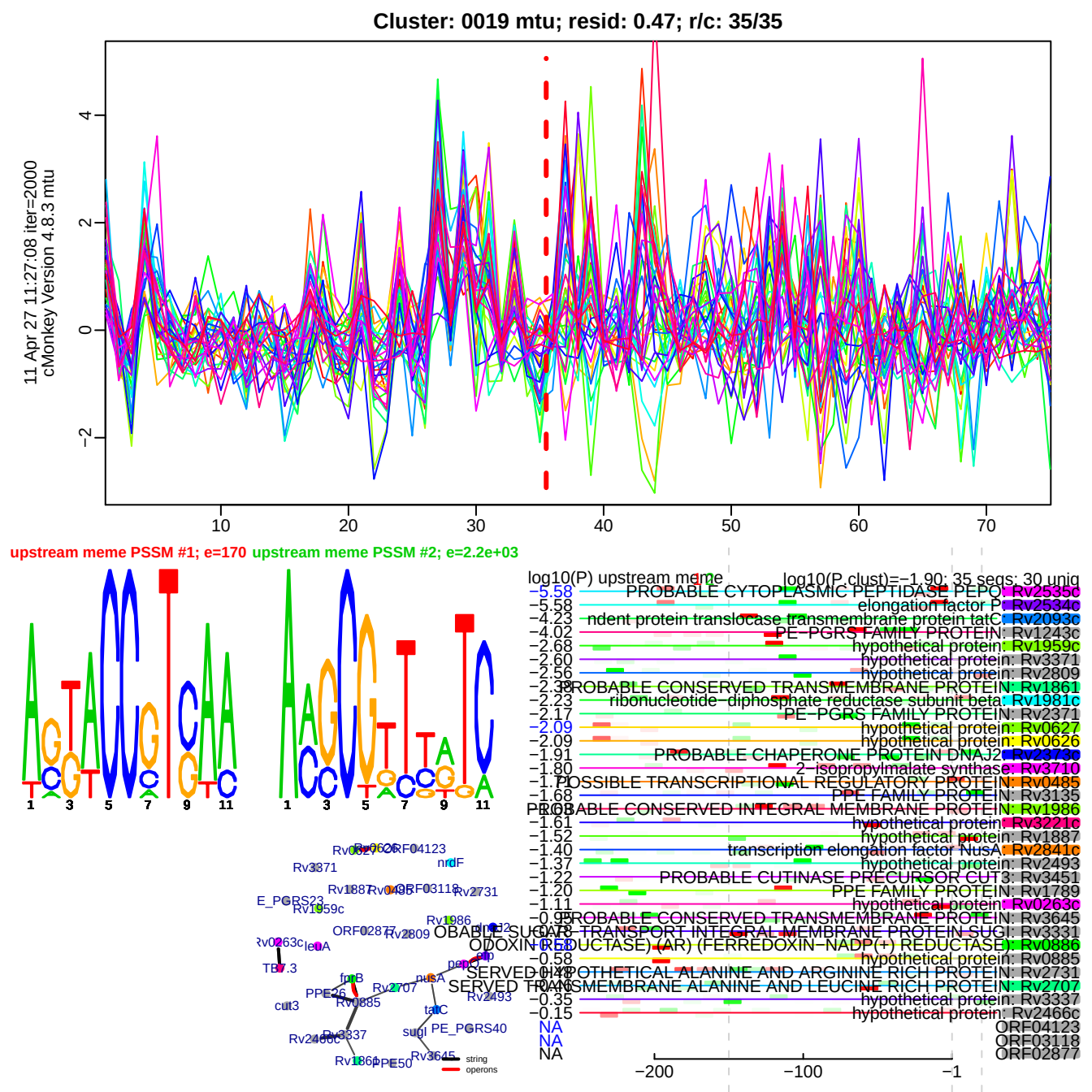
Cluster: 0019 mtu; resid: 0.47; r/c: 35/35
11 Apr 27 11:27:08 iter=2000
cMonkey Version 4.8.3 mtu
upstream meme PSSM #1; e=170
upstream meme PSSM #2; e=2.2e+03
log10(P) upstream meme
log10(P clust)=−1.90; 35 segs; 30 uniq
PROBABLE CYTOPLASMIC PEPTIDASE PEPO: Rv2535c
elongation factor P: Rv2534c
ndent protein translocase transmembrane protein tatC: Rv2093c
PE−PGRS FAMILY PROTEIN: Rv1243c
hypothetical protein: Rv1959c
hypothetical protein: Rv3371
hypothetical protein: Rv2809
PROBABLE CONSERVED TRANSMEMBRANE PROTEIN: Rv1861
ribonucleotide−diphosphate reductase subunit beta: Rv1981c
PE−PGRS FAMILY PROTEIN: Rv2371
hypothetical protein: Rv0627
hypothetical protein: Rv0626
PROBABLE CHAPERONE PROTEIN DNAJ2: Rv2373c
2−isopropylmalate synthase: Rv3710
POSSIBLE TRANSCRIPTIONAL REGULATORY PROTEIN: Rv0485
PPE FAMILY PROTEIN: Rv3135
PROBABLE CONSERVED INTEGRAL MEMBRANE PROTEIN: Rv1986
hypothetical protein: Rv3221c
hypothetical protein: Rv1887
transcription elongation factor NusA: Rv2841c
hypothetical protein: Rv2493
PROBABLE CUTINASE PRECURSOR CUT3: Rv3451
PPE FAMILY PROTEIN: Rv1789
hypothetical protein: Rv0263c
PROBABLE CONSERVED TRANSMEMBRANE PROTEIN: Rv3645
ODOXIN−REDUCTASE) (AR) (FERREDOXIN−NADP(+) REDUCTASE): Rv0886
hypothetical protein: Rv0885
SERVED HYPOTHETICAL ALANINE AND ARGININE RICH PROTEIN: Rv2731
SERVED TRANSMEMBRANE ALANINE AND LEUCINE RICH PROTEIN: Rv2707
hypothetical protein: Rv3337
hypothetical protein: Rv2466c
ORF04123
ORF03118
ORF02877
−200
−100
−1
string
operons