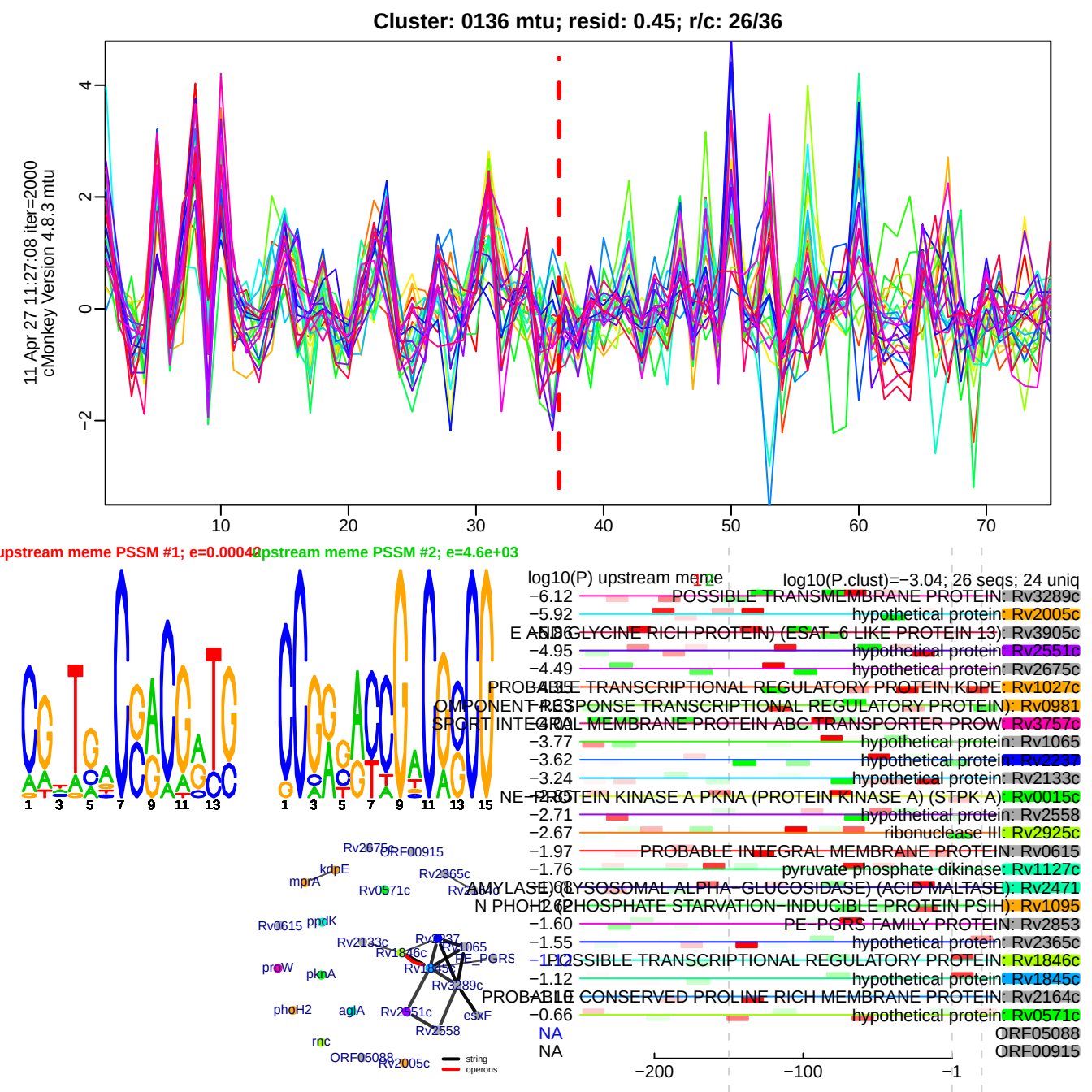
Cluster: 0136 mtu; resid: 0.45; r/c: 26/36
11 Apr 27 11:27:08 iter=2000
cMonkey Version 4.8.3 mtu
upstream meme PSSM #1; e=0.00042
upstream meme PSSM #2; e=4.6e+03
log10(P) upstream meme
log10(P.clust)=−3.04; 26 seqs; 24 uniq
−6.12 POSSIBLE TRANSMEMBRANE PROTEIN: Rv3289c
−5.92 hypothetical protein: Rv2005c
−5.06 E AND GLYCINE RICH PROTEIN) (ESAT−6 LIKE PROTEIN 13): Rv3905c
−4.95 hypothetical protein: Rv2551c
−4.49 hypothetical protein: Rv2675c
PROBABLE TRANSCRIPTIONAL REGULATORY PROTEIN KDPE: Rv1027c
OMPONENT-RESPONSE TRANSCRIPTIONAL REGULATORY PROTEIN): Rv0981
SPORT INTEGRAL MEMBRANE PROTEIN ABC TRANSPORTER PROW: Rv3757c
−3.77 hypothetical protein: Rv1065
−3.62 hypothetical protein: Rv2237
−3.24 hypothetical protein: Rv2133c
NE−PROTEIN KINASE A PKNA (PROTEIN KINASE A) (STPK A): Rv0015c
−2.71 hypothetical protein: Rv2558
−2.67 ribonuclease III: Rv2925c
−1.97 PROBABLE INTEGRAL MEMBRANE PROTEIN: Rv0615
−1.76 pyruvate phosphate dikinase: Rv1127c
AMYLASE) (LYSOSOMAL ALPHA−GLUCOSIDASE) (ACID MALTASE): Rv2471
N PHOH2 (PHOSPHATE STARVATION−INDUCIBLE PROTEIN PSIH): Rv1095
−1.60 PE−PGRS FAMILY PROTEIN: Rv2853
−1.55 hypothetical protein: Rv2365c
POSSIBLE TRANSCRIPTIONAL REGULATORY PROTEIN: Rv1846c
−1.12 hypothetical protein: Rv1845c
PROBABLE CONSERVED PROLINE RICH MEMBRANE PROTEIN: Rv2164c
−0.66 hypothetical protein: Rv0571c
NA ORF05088
NA ORF00915
−200 −100 −1
string
operons