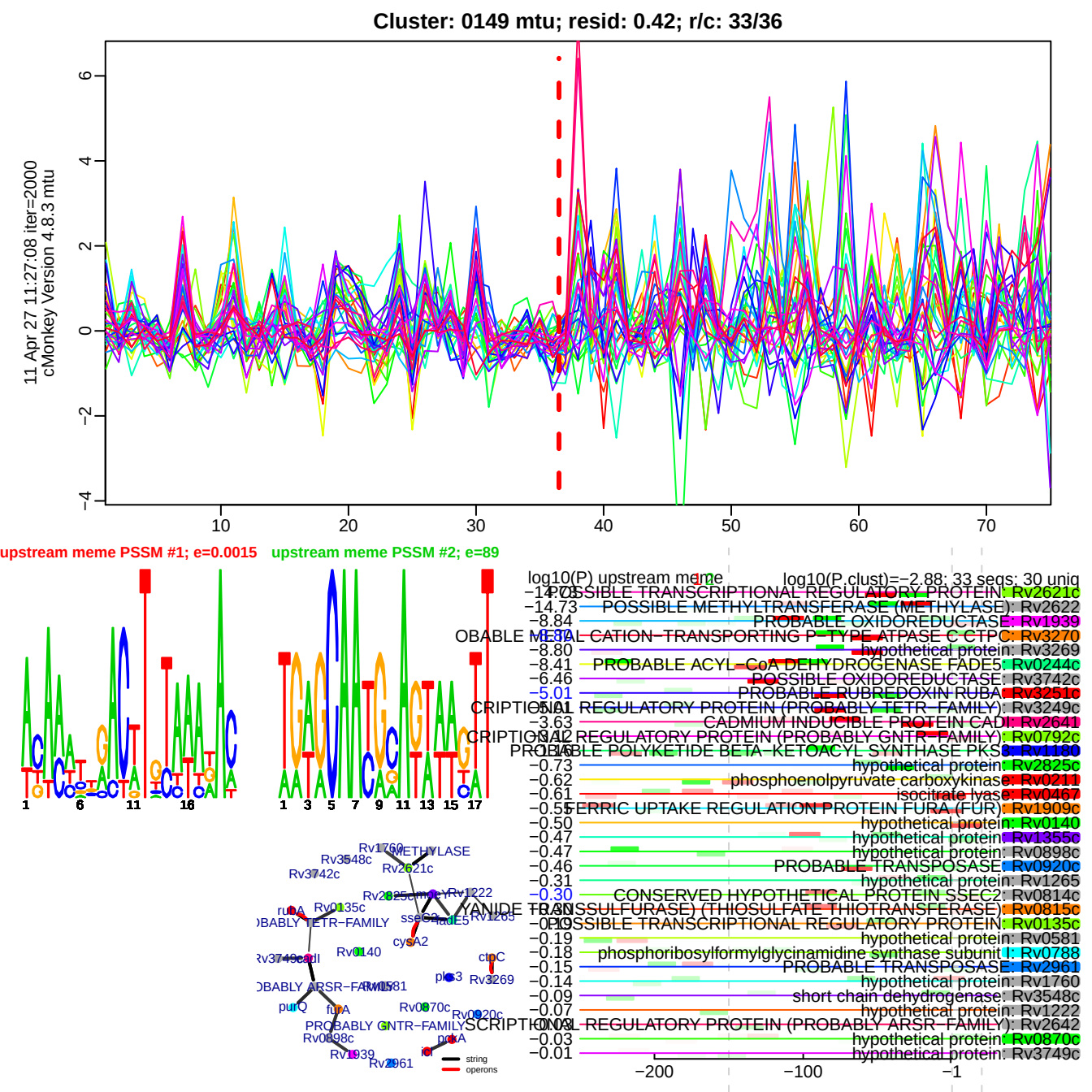
Cluster: 0149 mtu; resid: 0.42; r/c: 33/36
11 Apr 27 11:27:08 iter=2000
cMonkey Version 4.8.3 mtu
upstream meme PSSM #1; e=0.0015
upstream meme PSSM #2; e=89
log10(P) upstream meme
log10(P.clust)=−2.88; 33 seqs; 30 uniq
POSSIBLE TRANSCRIPTIONAL REGULATORY PROTEIN: Rv2621c
POSSIBLE METHYLTRANSFERASE (METHYLASE): Rv2622
PROBABLE OXIDOREDUCTASE: Rv1939
hypothetical protein: Rv3269
PROBABLE ACYL−CoA DEHYDROGENASE FADE5: Rv0244c
POSSIBLE OXIDOREDUCTASE: Rv3742c
PROBABLE RUBREDOXIN RUBA: Rv3251c
CADMIUM INDUCIBLE PROTEIN CADI: Rv2641
hypothetical protein: Rv2825c
phosphoenolpyruvate carboxykinase: Rv0211
isocitrate lyase: Rv0467
FERRIC UPTAKE REGULATION PROTEIN FURA (FUR): Rv1909c
hypothetical protein: Rv0140
hypothetical protein: Rv1355c
hypothetical protein: Rv0898c
PROBABLE TRANSPOSASE: Rv0920c
hypothetical protein: Rv1265
CONSERVED HYPOTHETICAL PROTEIN SSEC2: Rv0814c
hypothetical protein: Rv0581
hypothetical protein: Rv1760
short chain dehydrogenase: Rv3548c
hypothetical protein: Rv1222
hypothetical protein: Rv0870c
hypothetical protein: Rv3749c
−200
−100
−1
string
operons