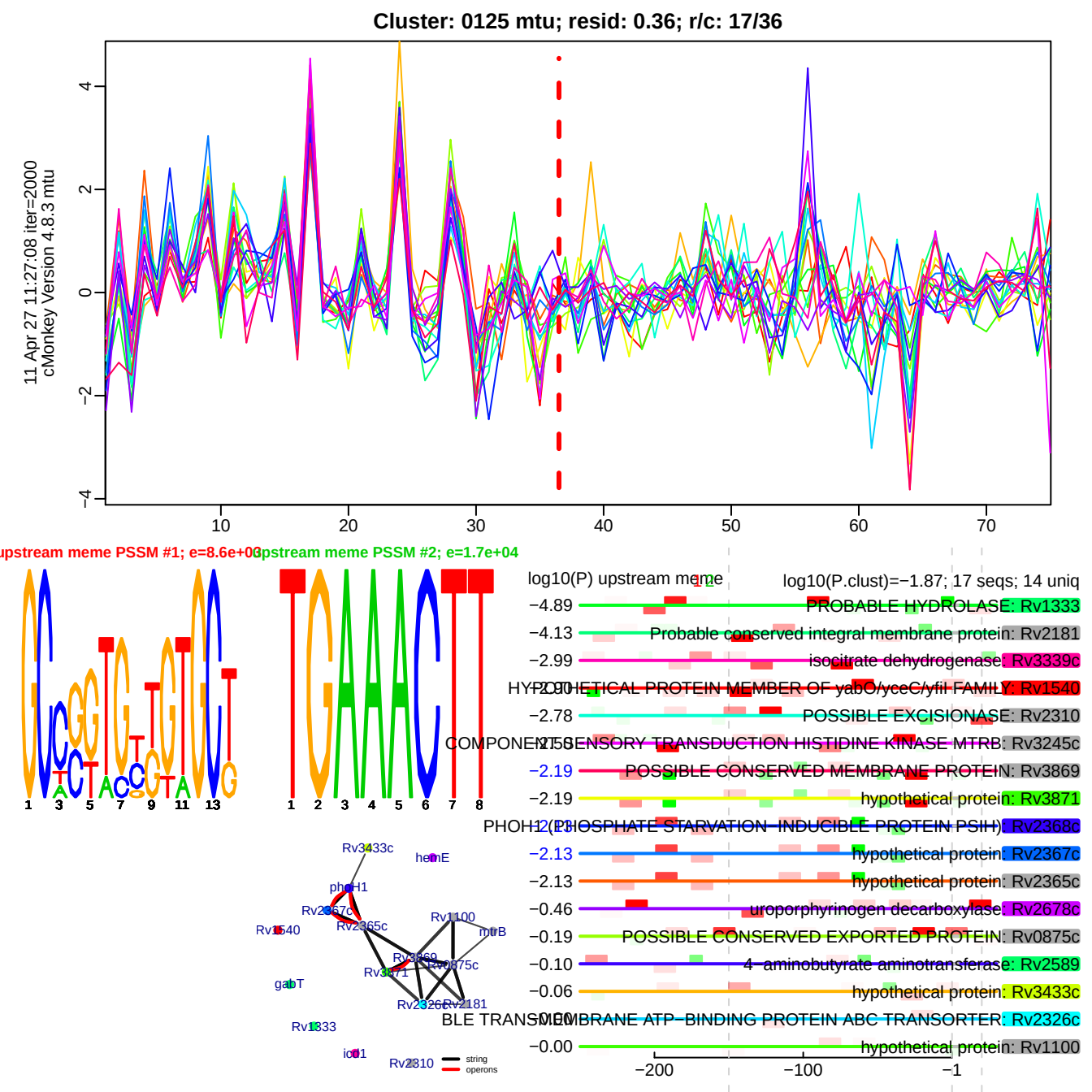
Cluster: 0125 mtu; resid: 0.36; r/c: 17/36
11 Apr 27 11:27:08 iter=2000
cMonkey Version 4.8.3 mtu
upstream meme PSSM #1; e=8.6e+03
upstream meme PSSM #2; e=1.7e+04
log10(P) upstream meme
log10(P.clust)=−1.87; 17 seqs; 14 uniq
−4.89 PROBABLE HYDROLASE: Rv1333
−4.13 Probable conserved integral membrane protein: Rv2181
−2.99 isocitrate dehydrogenase: Rv3339c
−2.90 HYPOTHETICAL PROTEIN MEMBER OF yabO/yceC/yfiI FAMILY: Rv1540
−2.78 POSSIBLE EXCISIONASE: Rv2310
−2.56 COMPONENT SENSORY TRANSDUCTION HISTIDINE KINASE MTRB: Rv3245c
−2.19 POSSIBLE CONSERVED MEMBRANE PROTEIN: Rv3869
−2.19 hypothetical protein: Rv3871
−2.14 PHOH2 (PHOSPHATE STARVATION−INDUCIBLE PROTEIN PSIH1): Rv2368c
−2.13 hypothetical protein: Rv2367c
−2.13 hypothetical protein: Rv2365c
−0.46 uroporphyrinogen decarboxylase: Rv2678c
−0.19 POSSIBLE CONSERVED EXPORTED PROTEIN: Rv0875c
−0.10 4−aminobutyrate aminotransferase: Rv2589
−0.06 hypothetical protein: Rv3433c
−0.20 BLE TRANSMEMBRANE ATP−BINDING PROTEIN ABC TRANSORTER: Rv2326c
−0.00 hypothetical protein: Rv1100
−200
−100
−1
Rv3433c
hemE
phoH1
Rv2367c
Rv2365c
Rv1540
Rv1100
mtrB
Rv3869
Rv3871
Rv0875c
gabT
Rv2326c
Rv2181
Rv1333
icd1
Rv2310
string
operons