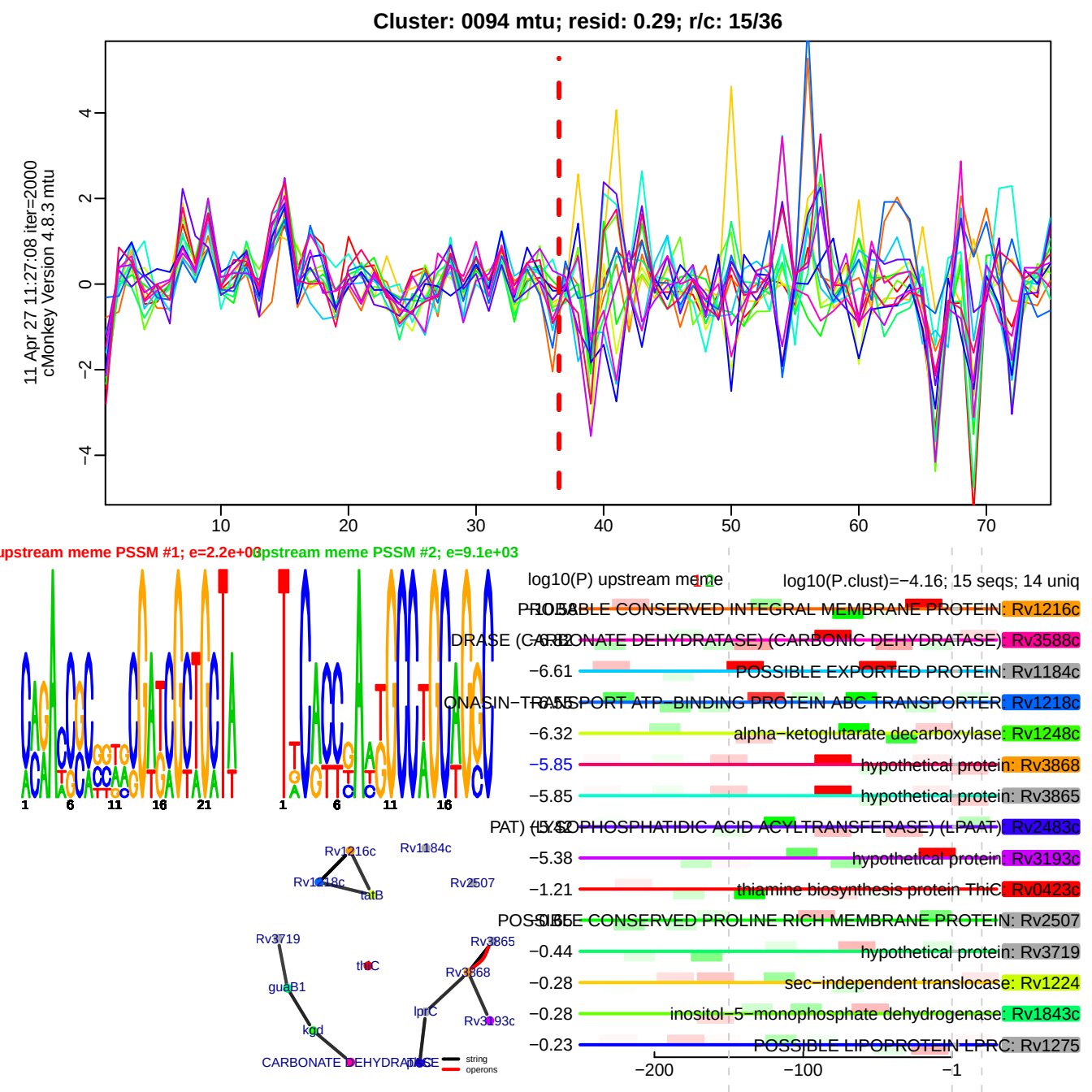
Cluster: 0094 mtu; resid: 0.29; r/c: 15/36
11 Apr 27 11:27:08 iter=2000
cMonkey Version 4.8.3 mtu
upstream meme PSSM #1; e=2.2e+03
upstream meme PSSM #2; e=9.1e+03
log10(P) upstream meme
log10(P.clust)=−4.16; 15 seqs; 14 uniq
POSSIBLE CONSERVED INTEGRAL MEMBRANE PROTEIN: Rv1216c
DRASE (CARBONATE DEHYDRATASE) (CARBONIC DEHYDRATASE): Rv3588c
POSSIBLE EXPORTED PROTEIN: Rv1184c
ONASIN−TRANSPORT ATP−BINDING PROTEIN ABC TRANSPORTER: Rv1218c
alpha−ketoglutarate decarboxylase: Rv1248c
hypothetical protein: Rv3868
hypothetical protein: Rv3865
PAT) (1−ACYLDIHYDROXYACETONE PHOSPHATE ACYLTRANSFERASE) (LPAAT): Rv2483c
hypothetical protein: Rv3193c
thiamine biosynthesis protein ThiC: Rv0423c
POSSIBLE CONSERVED PROLINE RICH MEMBRANE PROTEIN: Rv2507
hypothetical protein: Rv3719
sec−independent translocase: Rv1224
inositol−5−monophosphate dehydrogenase: Rv1843c
POSSIBLE LIPOPROTEIN LPRC: Rv1275
−200
−100
−1
Rv1216c
Rv1184c
Rv1218c
tatB
Rv2507
Rv3719
Rv3865
thiC
Rv3868
guaB1
lprC
Rv3193c
kgd
CARBONATE DEHYDRATASE
string
operons