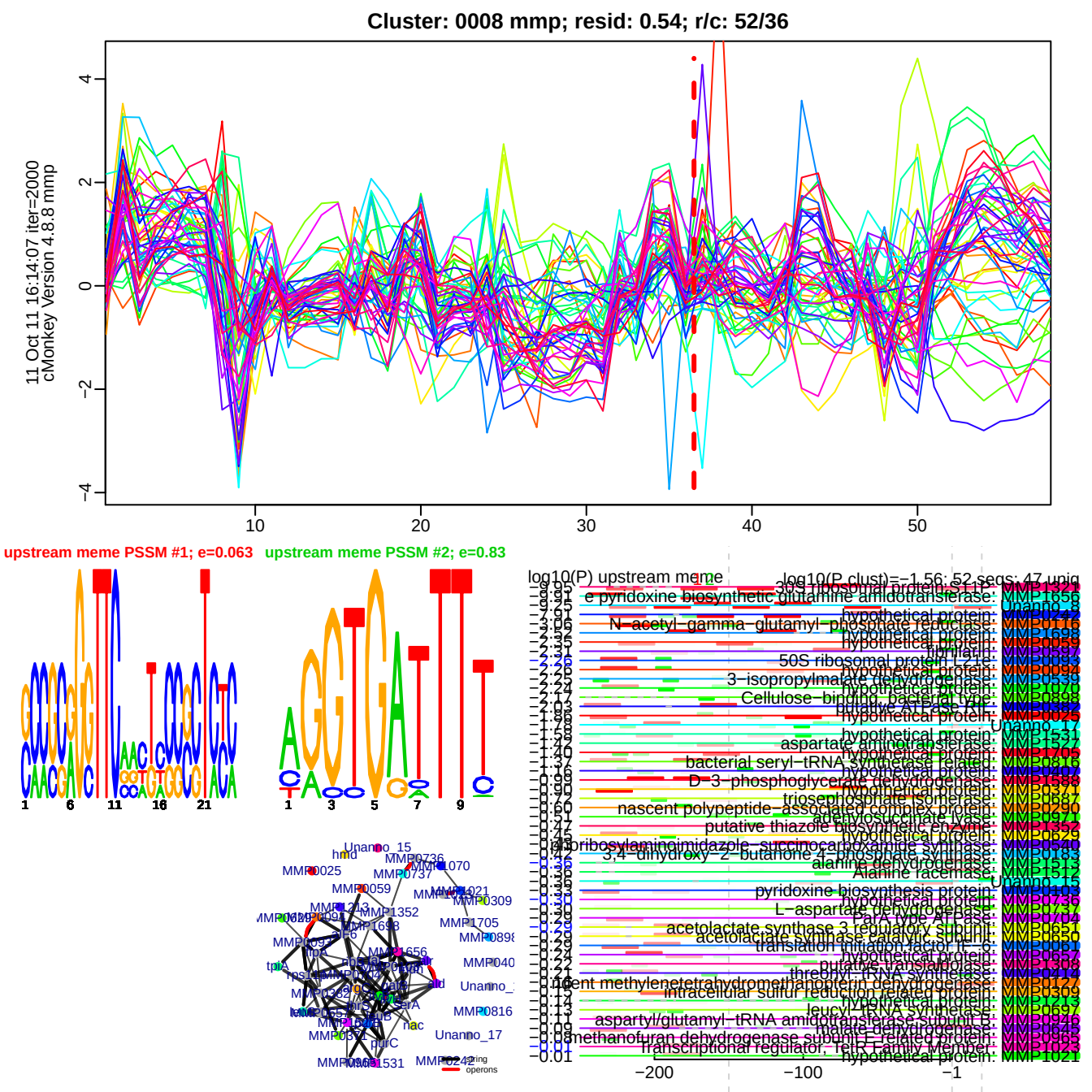
Cluster: 0008 mmp; resid: 0.54; r/c: 52/36
11 Oct 11 16:14:07 iter=2000
cMonkey Version 4.8.8 mmp
upstream meme PSSM #1; e=0.063
upstream meme PSSM #2; e=0.83
log10(P) upstream meme
log10(P clust)=−1.56; 52 seqs; 47 uniq
30S ribosomal protein S11P
e pyridoxine biosynthetic glutamine amidotransferase
hypothetical protein
N−acetyl−gamma−glutamyl−phosphate reductase
hypothetical protein
tubulin
50S ribosomal protein L21e
hypothetical protein
3−isopropylmalate dehydratase
hypothetical protein
Cellulose−binding, bacterial type
putative ATPase RIL
hypothetical protein
hypothetical protein
aspartate aminotransferase
hypothetical protein
bacterial seryl−tRNA synthetase related
hypothetical protein
D−3−phosphoglycerate dehydrogenase
hypothetical protein
triosephosphate isomerase
nascent polypeptide−associated complex protein
adenylosuccinate lyase
putative thiazole biosynthetic enzyme
hypothetical protein
phosphoribosylaminoimidazole−succinocarboxamide synthase
3,4−dihydroxy−2−butanone 4−phosphate synthase
alanine dehydrogenase
Alanine racemase
pyridoxine biosynthesis protein
hypothetical protein
L−aspartate dehydrogenase
ParA family
acetolactate synthase 3 regulatory subunit
acetolactate synthase catalytic subunit
translation initiation factor IF−6
hypothetical protein
putative transcriptase
threonyl−tRNA synthetase
nt methylenetetrahydromethanopterin dehydrogenase
intracellular sulfur reduction related protein
hypothetical protein
leucyl−tRNA synthetase
aspartyl/glutamyl−tRNA amidotransferase subunit B
malate dehydrogenase
methanofuran dehydrogenase subunit E related protein
transcriptional regulator, TetR family
hypothetical protein
−200
−100
−1
Unanno_15
hmd
MMP0736
MMP0025
MMP0737
MMP1070
MMP0059
MMP1021
MMP0309
MMP1213
MMP1352
MMP1705
MMP1656
MMP0408
Unanno_17
MMP0816
MMP0242
MMP1531
operons