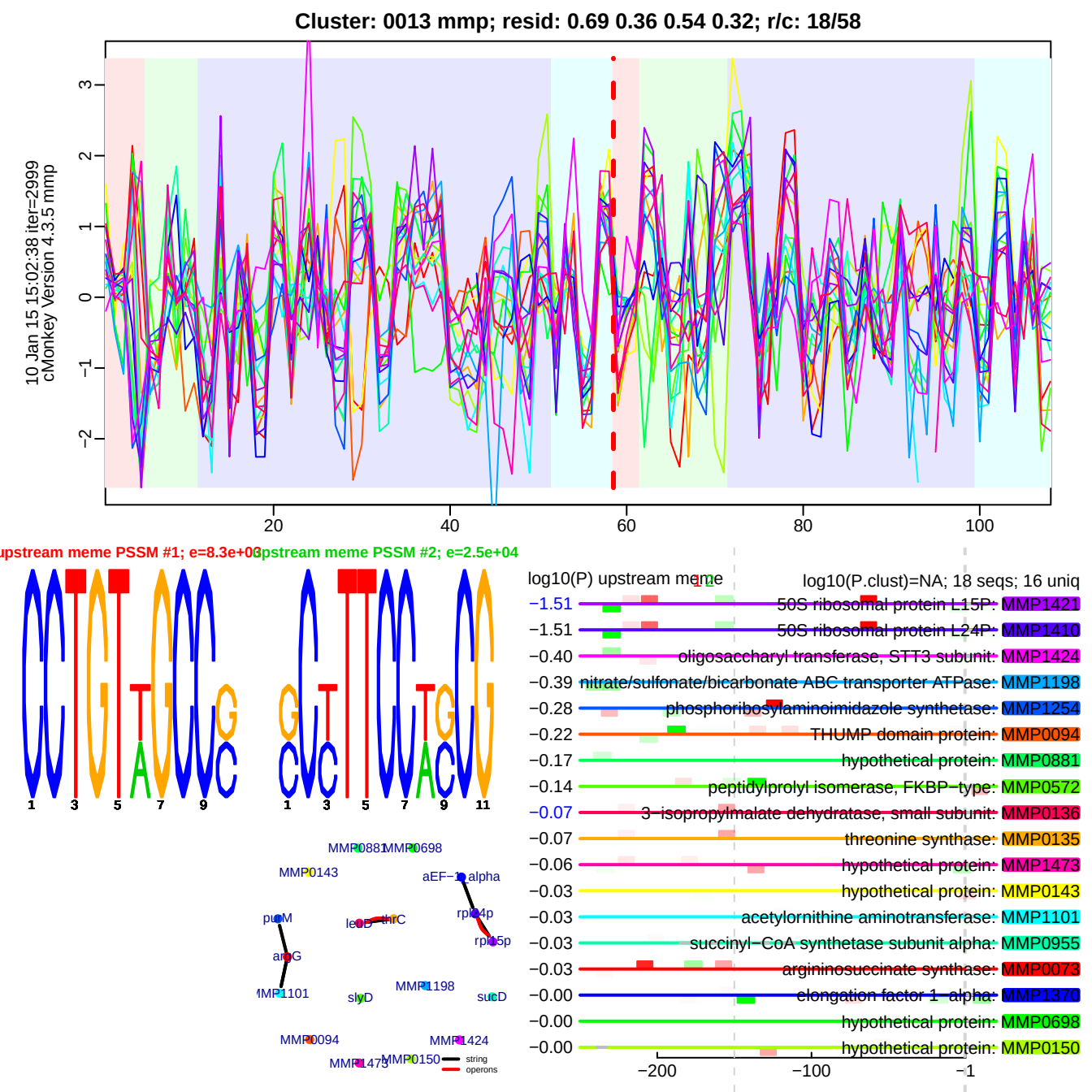
Cluster: 0013 mmp; resid: 0.69 0.36 0.54 0.32; r/c: 18/58
10 Jan 15 15:02:38 iter=2999
cMonkey Version 4.3.5 mmp
upstream meme PSSM #1; e=8.3e+03
upstream meme PSSM #2; e=2.5e+04
log10(P) upstream meme
log10(P.clust)=NA; 18 seqs; 16 uniq
−1.51 50S ribosomal protein L15P: MMP1421
−1.51 50S ribosomal protein L24P: MMP1410
−0.40 oligosaccharyl transferase, STT3 subunit: MMP1424
−0.39 nitrate/sulfonate/bicarbonate ABC transporter ATPase: MMP1198
−0.28 phosphoribosylaminoimidazole synthetase: MMP1254
−0.22 THUMP domain protein: MMP0094
−0.17 hypothetical protein: MMP0881
−0.14 peptidylprolyl isomerase, FKBP−type: MMP0572
−0.07 3−isopropylmalate dehydratase, small subunit: MMP0136
−0.07 threonine synthase: MMP0135
−0.06 hypothetical protein: MMP1473
−0.03 hypothetical protein: MMP0143
−0.03 acetylornithine aminotransferase: MMP1101
−0.03 succinyl−CoA synthetase subunit alpha: MMP0955
−0.03 argininosuccinate synthase: MMP0073
−0.00 elongation factor 1−alpha: MMP1370
−0.00 hypothetical protein: MMP0698
−0.00 hypothetical protein: MMP0150
MMP0881
MMP0698
MMP0143
aEF−1a alpha
rpl24p
rpl15p
purM
leuB
thrC
argG
MMP1101
MMP1198
sdhD
sucD
MMP0094
MMP1424
MMP1473
MMP0150
string
operons