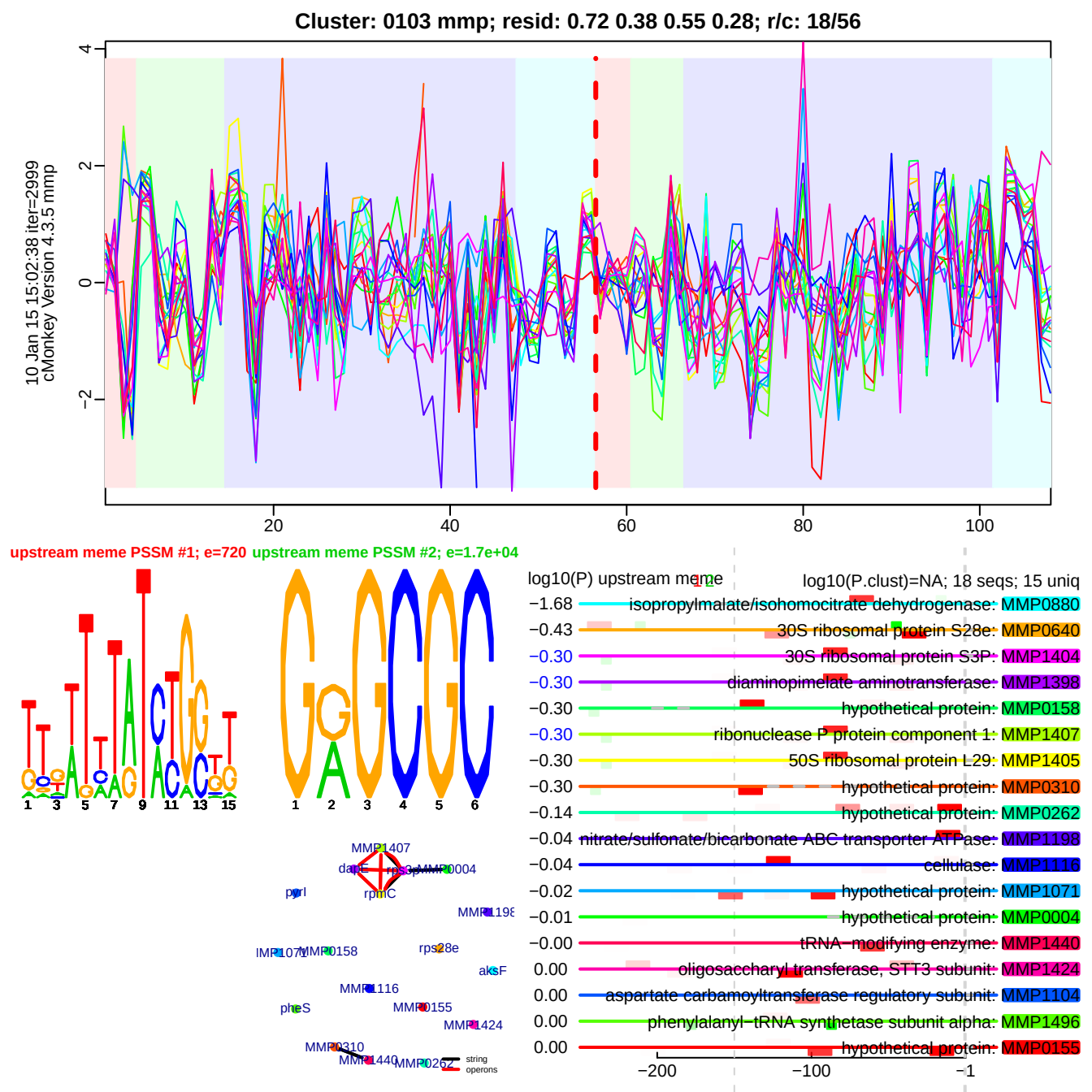

Cluster: 0103 mmp; resid: 0.72 0.38 0.55 0.28; r/c: 18/56
10 Jan 15 15:02:38 iter=2999
cMonkey Version 4.3.5 mmp
upstream meme PSSM #1; e=720
upstream meme PSSM #2; e=1.7e+04
log10(P) upstream meme
log10(P.clust)=NA; 18 seqs; 15 uniq
isopropylmalate/isohomocitrate dehydrogenase: MMP0880
30S ribosomal protein S28e: MMP0640
30S ribosomal protein S3P: MMP1404
diaminopimelate aminotransferase: MMP1398
hypothetical protein: MMP0158
ribonuclease P protein component 1: MMP1407
50S ribosomal protein L29: MMP1405
hypothetical protein: MMP0310
hypothetical protein: MMP0262
nitrate/sulfonate/bicarbonate ABC transporter ATPase: MMP1198
cellulase: MMP1116
hypothetical protein: MMP1071
hypothetical protein: MMP0004
tRNA−modifying enzyme: MMP1440
oligosaccharyl transferase, STT3 subunit: MMP1424
aspartate carbamoyltransferase regulatory subunit: MMP1104
phenylalanyl−tRNA synthetase subunit alpha: MMP1496
hypothetical protein: MMP0155
string
operons