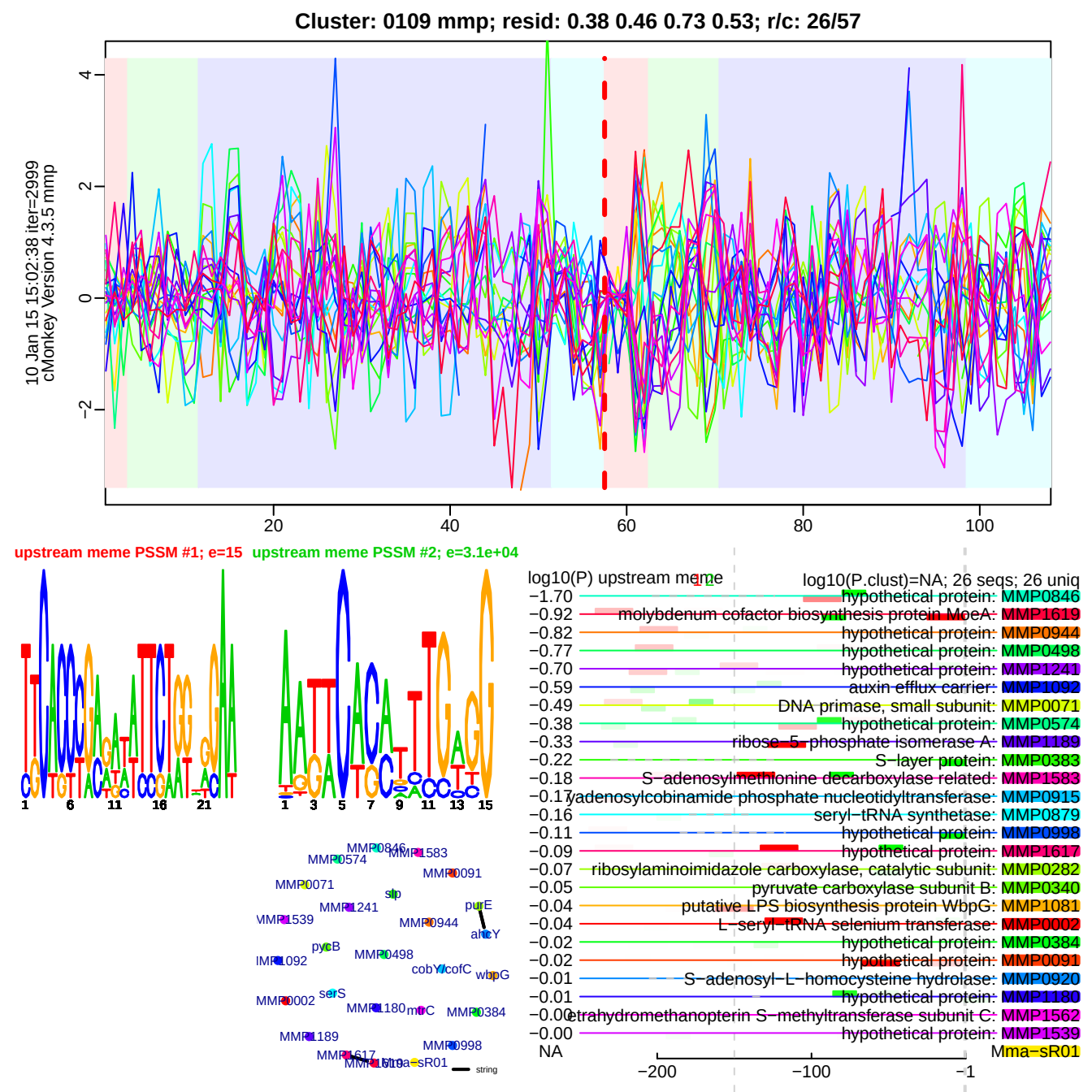
Cluster: 0109 mmp; resid: 0.38 0.46 0.73 0.53; r/c: 26/57
10 Jan 15 15:02:38 iter=2999
cMonkey Version 4.3.5 mmp
upstream meme PSSM #1; e=15
upstream meme PSSM #2; e=3.1e+04
log10(P) upstream meme
log10(P.clust)=NA; 26 seqs; 26 uniq
−1.70 hypothetical protein: MMP0846
−0.92 molybdenum cofactor biosynthesis protein MoeA: MMP1619
−0.82 hypothetical protein: MMP0944
−0.77 hypothetical protein: MMP0498
−0.70 hypothetical protein: MMP1241
−0.59 auxin efflux carrier: MMP1092
−0.49 DNA primase, small subunit: MMP0071
−0.38 hypothetical protein: MMP0574
−0.33 ribose−5−phosphate isomerase A: MMP1189
−0.22 S−layer protein: MMP0383
−0.18 S−adenosylmethionine decarboxylase related: MMP1583
−0.17 adenosylcobinamide phosphate nucleotidyltransferase: MMP0915
−0.16 seryl−tRNA synthetase: MMP0879
−0.11 hypothetical protein: MMP0998
−0.09 hypothetical protein: MMP1617
−0.07 ribosylaminoimidazole carboxylase, catalytic subunit: MMP0282
−0.05 pyruvate carboxylase subunit B: MMP0340
−0.04 putative LPS biosynthesis protein WbpG: MMP1081
−0.04 L−seryl−tRNA selenium transferase: MMP0002
−0.02 hypothetical protein: MMP0384
−0.02 hypothetical protein: MMP0091
−0.01 S−adenosyl−L−homocysteine hydrolase: MMP0920
−0.01 hypothetical protein: MMP1180
−0.00 tetrahydromethanopterin S−methyltransferase subunit C: MMP1562
−0.00 hypothetical protein: MMP1539
NA Mma−sR01
MMP0846
MMP0574
MMP1583
MMP0091
MMP0071
slp
MMP1241
purE
MMP1539
MMP0944
ahcY
pycB
MMP1092
MMP0498
cobY/cofC
wbpG
serS
MMP0002
MMP1180
mtC
MMP0384
MMP1189
MMP0998
MMP1617
string