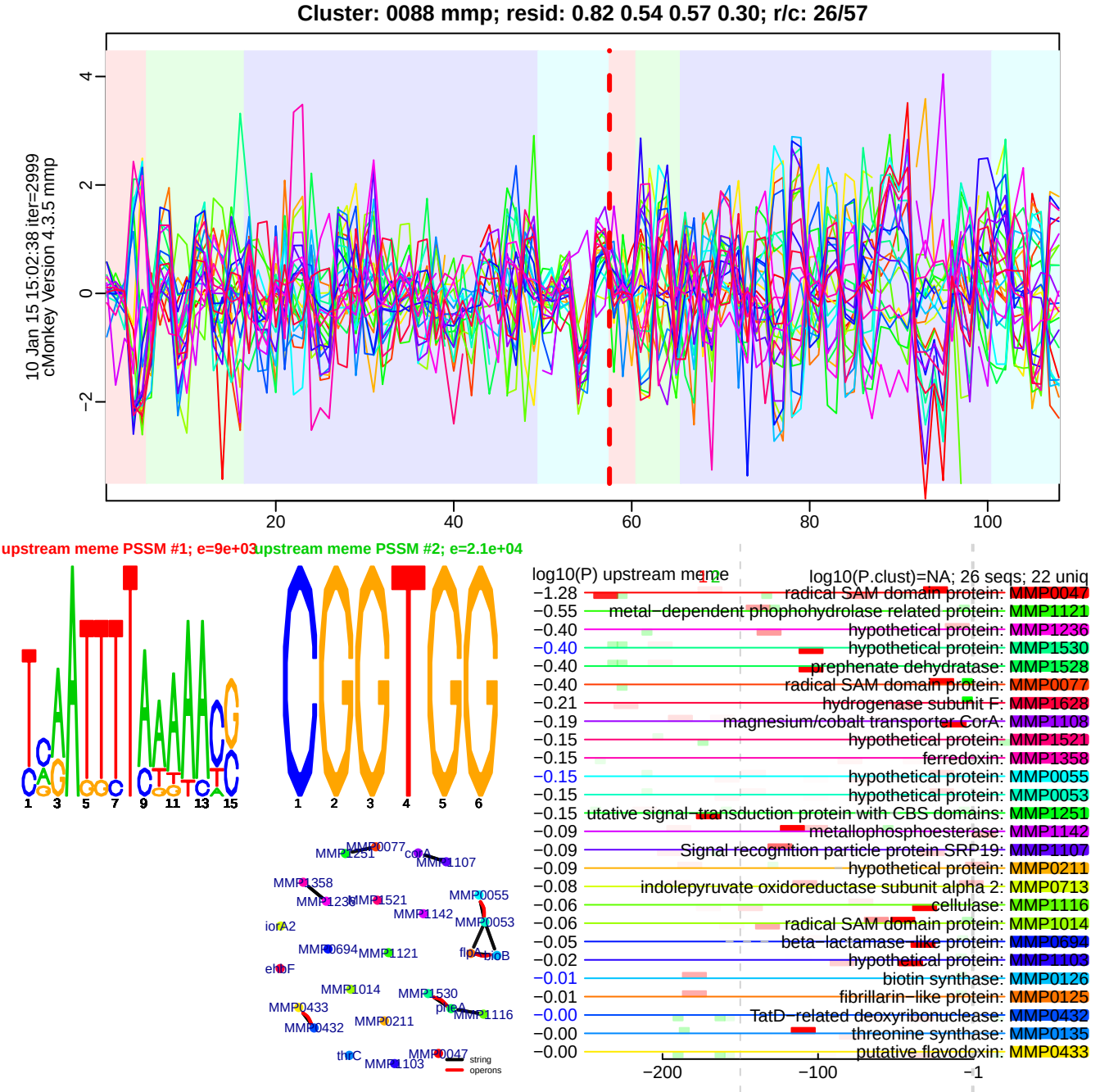
Cluster: 0088 mmp; resid: 0.82 0.54 0.57 0.30; r/c: 26/57
10 Jan 15 15:02:38 iter=2999
cMonkey Version 4.3.5 mmp
upstream meme PSSM #1; e=9e+03
upstream meme PSSM #2; e=2.1e+04
log10(P) upstream meme
log10(P.clust)=NA; 26 seqs; 22 uniq
−1.28 radical SAM domain protein: MMP0047
−0.55 metal−dependent phophohydrolase related protein: MMP1121
−0.40 hypothetical protein: MMP1236
−0.40 hypothetical protein: MMP1530
−0.40 prephenate dehydratase: MMP1528
−0.40 radical SAM domain protein: MMP0077
−0.21 hydrogenase subunit F: MMP1628
−0.19 magnesium/cobalt transporter CorA: MMP1108
−0.15 hypothetical protein: MMP1521
−0.15 ferredoxin: MMP1358
−0.15 hypothetical protein: MMP0055
−0.15 hypothetical protein: MMP0053
−0.15 utative signal−transduction protein with CBS domains: MMP1251
−0.09 metallophosphoesterase: MMP1142
−0.09 Signal recognition particle protein SRP19: MMP1107
−0.09 hypothetical protein: MMP0211
−0.08 indolepyruvate oxidoreductase subunit alpha 2: MMP0713
−0.06 cellulase: MMP1116
−0.06 radical SAM domain protein: MMP1014
−0.05 beta−lactamase−like protein: MMP0694
−0.02 hypothetical protein: MMP1103
−0.01 biotin synthase: MMP0126
−0.01 fibrillarin−like protein: MMP0125
−0.00 TatD−related deoxyribonuclease: MMP0432
−0.00 threonine synthase: MMP0135
−0.00 putative flavodoxin: MMP0433
MMP0077
MMP1251
corA
MMP1107
MMP1358
MMP1236
MMP1521
MMP0055
MMP1142
MMP0053
iorA2
MMP0694
MMP1121
flgA
bioB
ehbF
MMP1014
MMP1530
MMP0433
phbA
MMP1116
MMP0211
MMP0432
thrC
MMP0047
MMP1103
string
operons