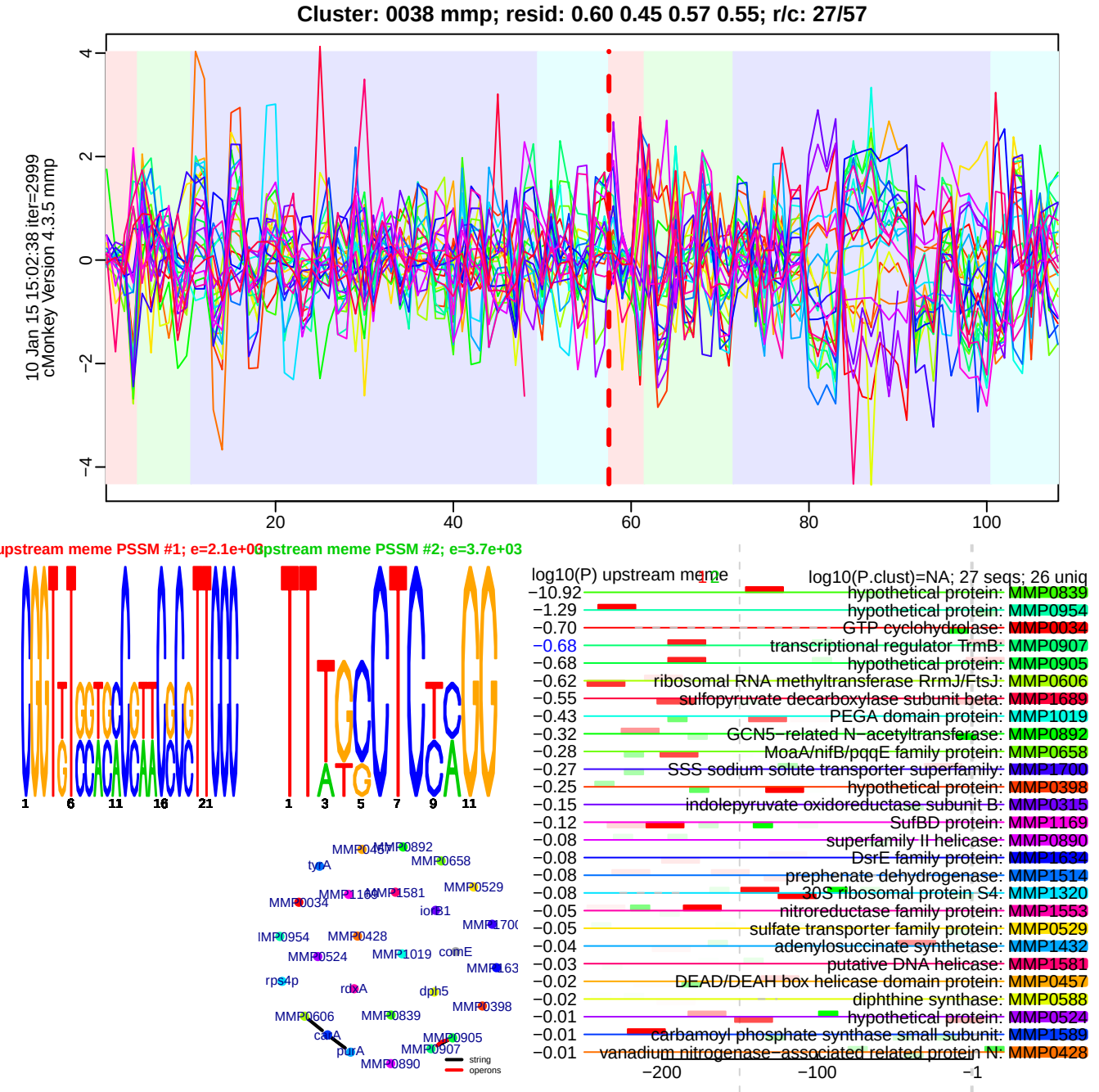
Cluster: 0038 mmp; resid: 0.60 0.45 0.57 0.55; r/c: 27/57
10 Jan 15 15:02:38 iter=2999
cMonkey Version 4.3.5 mmp
upstream meme PSSM #1; e=2.1e+03
upstream meme PSSM #2; e=3.7e+03
log10(P) upstream meme
log10(P.clust)=NA; 27 seqs; 26 uniq
−10.92 hypothetical protein: MMP0839
−1.29 hypothetical protein: MMP0954
−0.70 GTP cyclohydrolase: MMP0034
−0.68 transcriptional regulator TrmB: MMP0907
−0.68 hypothetical protein: MMP0905
−0.62 ribosomal RNA methyltransferase RrmJ/FtsJ: MMP0606
−0.55 sulfopyruvate decarboxylase subunit beta: MMP1689
−0.43 PEGA domain protein: MMP1019
−0.32 GCN5−related N−acetyltransferase: MMP0892
−0.28 MoaA/nifB/pqqE family protein: MMP0658
−0.27 SSS sodium solute transporter superfamily: MMP1700
−0.25 hypothetical protein: MMP0398
−0.15 indolepyruvate oxidoreductase subunit B: MMP0315
−0.12 SufBD protein: MMP1169
−0.08 superfamily II helicase: MMP0890
−0.08 DsrE family protein: MMP1634
−0.08 prephenate dehydrogenase: MMP1514
−0.08 30S ribosomal protein S4: MMP1320
−0.05 nitroreductase family protein: MMP1553
−0.05 sulfate transporter family protein: MMP0529
−0.04 adenylosuccinate synthetase: MMP1432
−0.03 putative DNA helicase: MMP1581
−0.02 DEAD/DEAH box helicase domain protein: MMP0457
−0.02 diphthine synthase: MMP0588
−0.01 hypothetical protein: MMP0524
−0.01 carbamoyl phosphate synthase small subunit: MMP1589
−0.01 vanadium nitrogenase−associated related protein N: MMP0428
string
operons