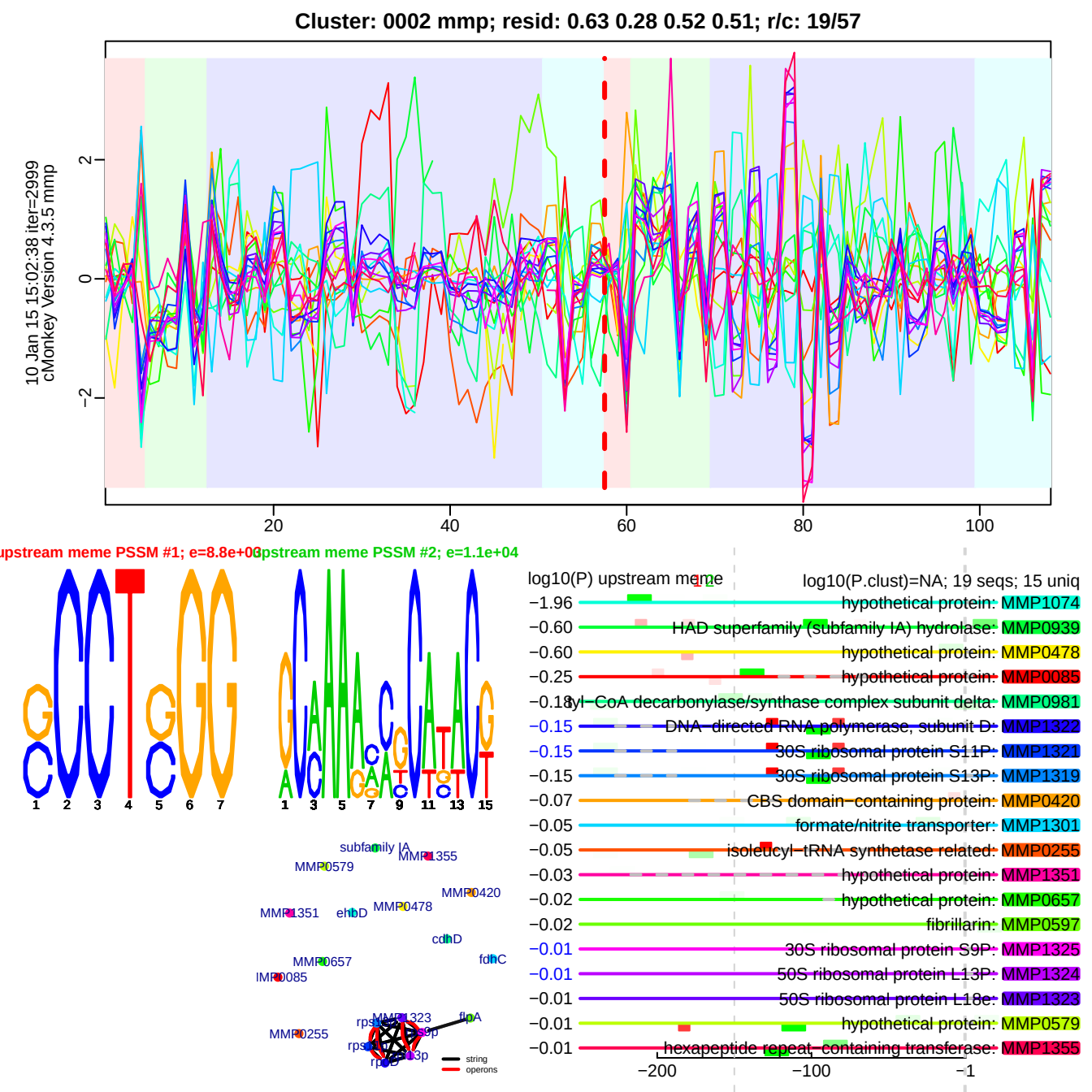
Cluster: 0002 mmp; resid: 0.63 0.28 0.52 0.51; r/c: 19/57
10 Jan 15 15:02:38 iter=2999
cMonkey Version 4.3.5 mmp
upstream meme PSSM #1; e=8.8e+03
upstream meme PSSM #2; e=1.1e+04
log10(P) upstream meme
log10(P.clust)=NA; 19 seqs; 15 uniq
−1.96 hypothetical protein: MMP1074
−0.60 HAD superfamily (subfamily IA) hydrolase: MMP0939
−0.60 hypothetical protein: MMP0478
−0.25 hypothetical protein: MMP0085
−0.18 3-hydroxyacyl−CoA decarbonylase/synthase complex subunit delta: MMP0981
−0.15 DNA−directed RNA polymerase, subunit D: MMP1322
−0.15 30S ribosomal protein S11P: MMP1321
−0.15 30S ribosomal protein S13P: MMP1319
−0.07 CBS domain−containing protein: MMP0420
−0.05 formate/nitrite transporter: MMP1301
−0.05 isoleucyl−tRNA synthetase related: MMP0255
−0.03 hypothetical protein: MMP1351
−0.02 hypothetical protein: MMP0657
−0.02 fibrillarin: MMP0597
−0.01 30S ribosomal protein S9P: MMP1325
−0.01 50S ribosomal protein L13P: MMP1324
−0.01 50S ribosomal protein L18e: MMP1323
−0.01 hypothetical protein: MMP0579
−0.01 hexapeptide repeat−containing transferase: MMP1355
−200
−100
−1
subfamily IA
MMP1355
MMP0579
MMP0420
MMP1351
ehbD
MMP0478
cdhD
fdhC
MMP0657
MMP0085
MMP1323
flgA
MMP0255
string
operons