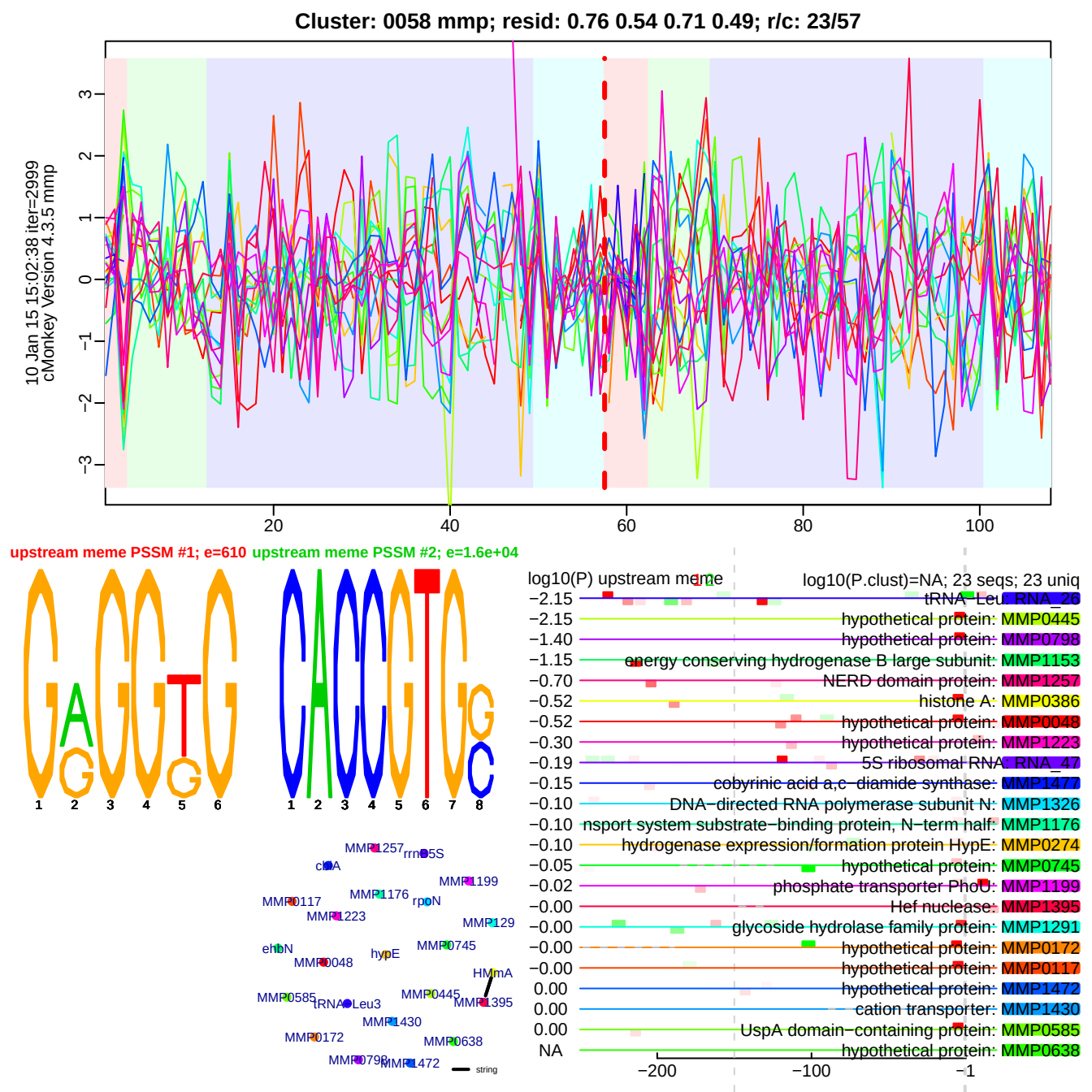
Cluster: 0058 mmp; resid: 0.76 0.54 0.71 0.49; r/c: 23/57
10 Jan 15 15:02:38 iter=2999
cMonkey Version 4.3.5 mmp
upstream meme PSSM #1; e=610
upstream meme PSSM #2; e=1.6e+04
log10(P) upstream meme
log10(P.clust)=NA; 23 seqs; 23 uniq
−2.15 tRNA−Leu: RNA_26
−2.15 hypothetical protein: MMP0445
−1.40 hypothetical protein: MMP0798
−1.15 energy conserving hydrogenase B large subunit: MMP1153
−0.70 NERD domain protein: MMP1257
−0.52 histone A: MMP0386
−0.52 hypothetical protein: MMP0048
−0.30 hypothetical protein: MMP1223
−0.19 5S ribosomal RNA: RNA_47
−0.15 cobyrinic acid a,c−diamide synthase: MMP1477
−0.10 DNA−directed RNA polymerase subunit N: MMP1326
−0.10 nsport system substrate−binding protein, N−term half: MMP1176
−0.10 hydrogenase expression/formation protein HypE: MMP0274
−0.05 hypothetical protein: MMP0745
−0.02 phosphate transporter PhoU: MMP1199
−0.00 Hef nuclease: MMP1395
−0.00 glycoside hydrolase family protein: MMP1291
−0.00 hypothetical protein: MMP0172
−0.00 hypothetical protein: MMP0117
0.00 hypothetical protein: MMP1472
0.00 cation transporter: MMP1430
0.00 UspA domain−containing protein: MMP0585
NA hypothetical protein: MMP0638
MMP1257
rrnB 5S
cbiA
MMP1199
MMP1176
rpoN
MMP0117
MMP1223
MMP1291
ehbN
MMP0745
hypE
MMP0048
HMmA
MMP0585
tRNA−Leu3
MMP0445
MMP1395
MMP1430
MMP0172
MMP0638
MMP0798
MMP1472
string