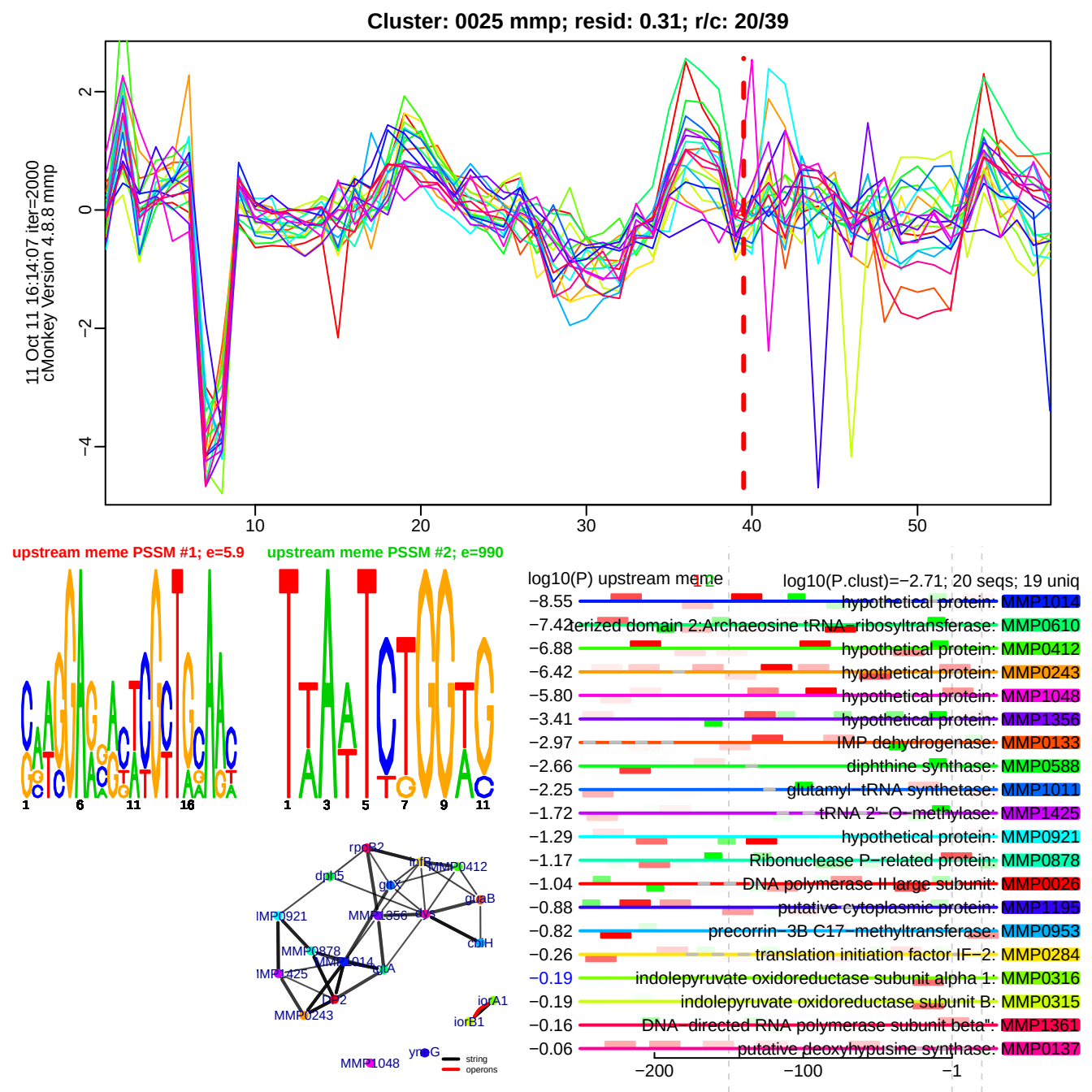

Cluster: 0025 mmp; resid: 0.31; r/c: 20/39
11 Oct 11 16:14:07 iter=2000
cMonkey Version 4.8.8 mmp
upstream meme PSSM #1; e=5.9
upstream meme PSSM #2; e=990
log10(P) upstream meme
log10(P.clust)=−2.71; 20 seqs; 19 uniq
−8.55 hypothetical protein: MMP1014
−7.42 terized domain 2:Archaeosine tRNA−ribosyltransferase: MMP0610
−6.88 hypothetical protein: MMP0412
−6.42 hypothetical protein: MMP0243
−5.80 hypothetical protein: MMP1048
−3.41 hypothetical protein: MMP1356
−2.97 IMP dehydrogenase: MMP0133
−2.66 diphthine synthase: MMP0588
−2.25 glutamyl−tRNA synthetase: MMP1011
−1.72 tRNA 2'−O−methylase: MMP1425
−1.29 hypothetical protein: MMP0921
−1.17 Ribonuclease P−related protein: MMP0878
−1.04 DNA polymerase II large subunit: MMP0026
−0.88 putative cytoplasmic protein: MMP1195
−0.82 precorrin−3B C17−methyltransferase: MMP0953
−0.26 translation initiation factor IF−2: MMP0284
−0.19 indolepyruvate oxidoreductase subunit alpha 1: MMP0316
−0.19 indolepyruvate oxidoreductase subunit B: MMP0315
−0.16 DNA−directed RNA polymerase subunit beta": MMP1361
−0.06 putative deoxyhypusine synthase: MMP0137
string
operons