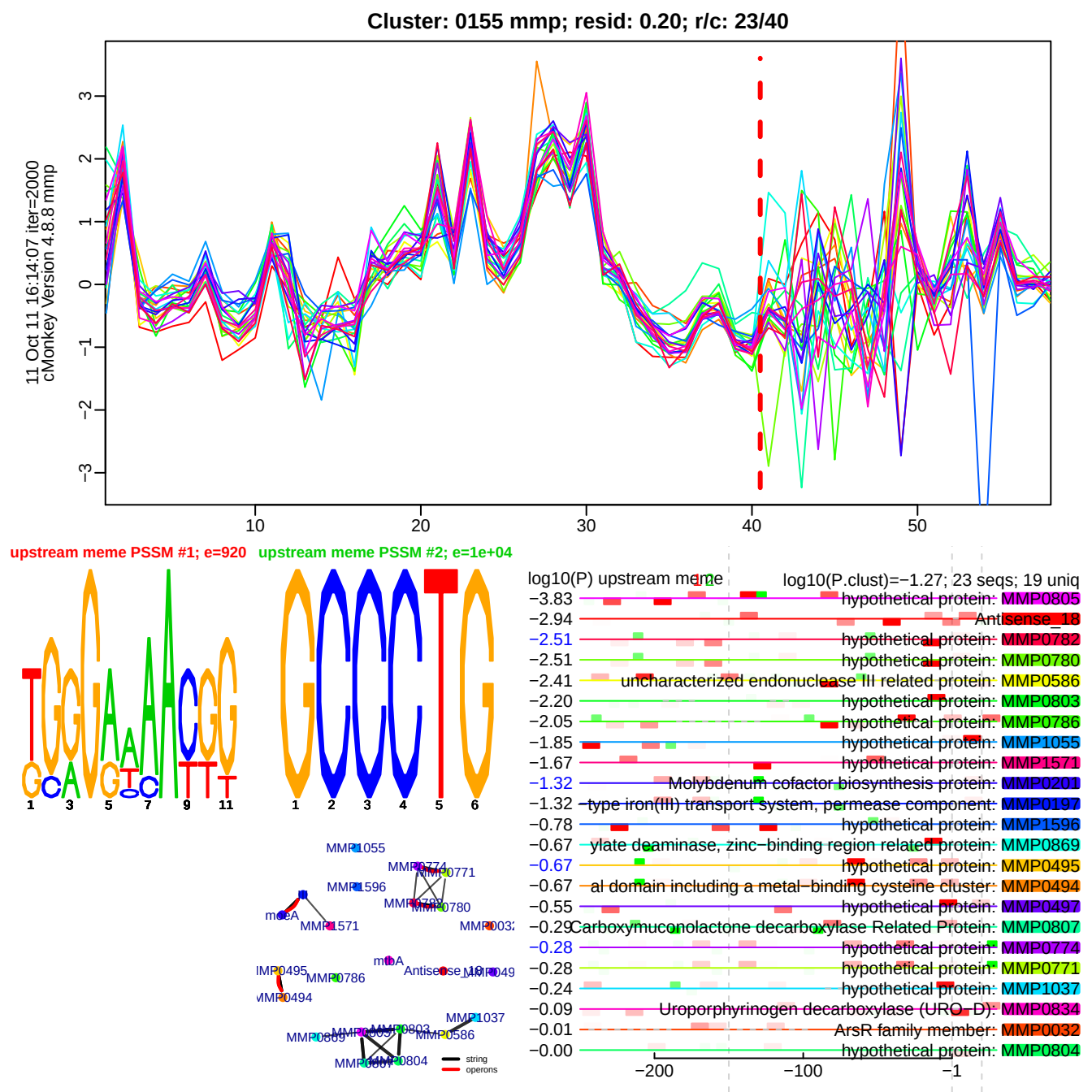
Cluster: 0155 mmp; resid: 0.20; r/c: 23/40
11 Oct 11 16:14:07 iter=2000
cMonkey Version 4.8.8 mmp
upstream meme PSSM #1; e=920
upstream meme PSSM #2; e=1e+04
log10(P) upstream meme
log10(P.clust)=−1.27; 23 seqs; 19 uniq
−3.83 hypothetical protein: MMP0805
−2.94 Antisense_18
−2.51 hypothetical protein: MMP0782
−2.51 hypothetical protein: MMP0780
−2.41 uncharacterized endonuclease III related protein: MMP0586
−2.20 hypothetical protein: MMP0803
−2.05 hypothetical protein: MMP0786
−1.85 hypothetical protein: MMP1055
−1.67 hypothetical protein: MMP1571
−1.32 Molybdenum cofactor biosynthesis protein: MMP0201
−1.32 type iron(III) transport system, permease component: MMP0197
−0.78 hypothetical protein: MMP1596
−0.67 ylate deaminase, zinc−binding region related protein: MMP0869
−0.67 hypothetical protein: MMP0495
−0.67 al domain including a metal−binding cysteine cluster: MMP0494
−0.55 hypothetical protein: MMP0497
−0.29 Carboxymuconolactone decarboxylase Related Protein: MMP0807
−0.28 hypothetical protein: MMP0774
−0.28 hypothetical protein: MMP0771
−0.24 hypothetical protein: MMP1037
−0.09 Uroporphyrinogen decarboxylase (URO−D): MMP0834
−0.01 ArsR family member: MMP0032
−0.00 hypothetical protein: MMP0804
MMP1055
MMP0774
MMP0771
MMP1596
MMP0782
MMP0780
MMP1571
MMP0786
MMP1037
MMP0803
MMP0586
MMP0804
string
operons