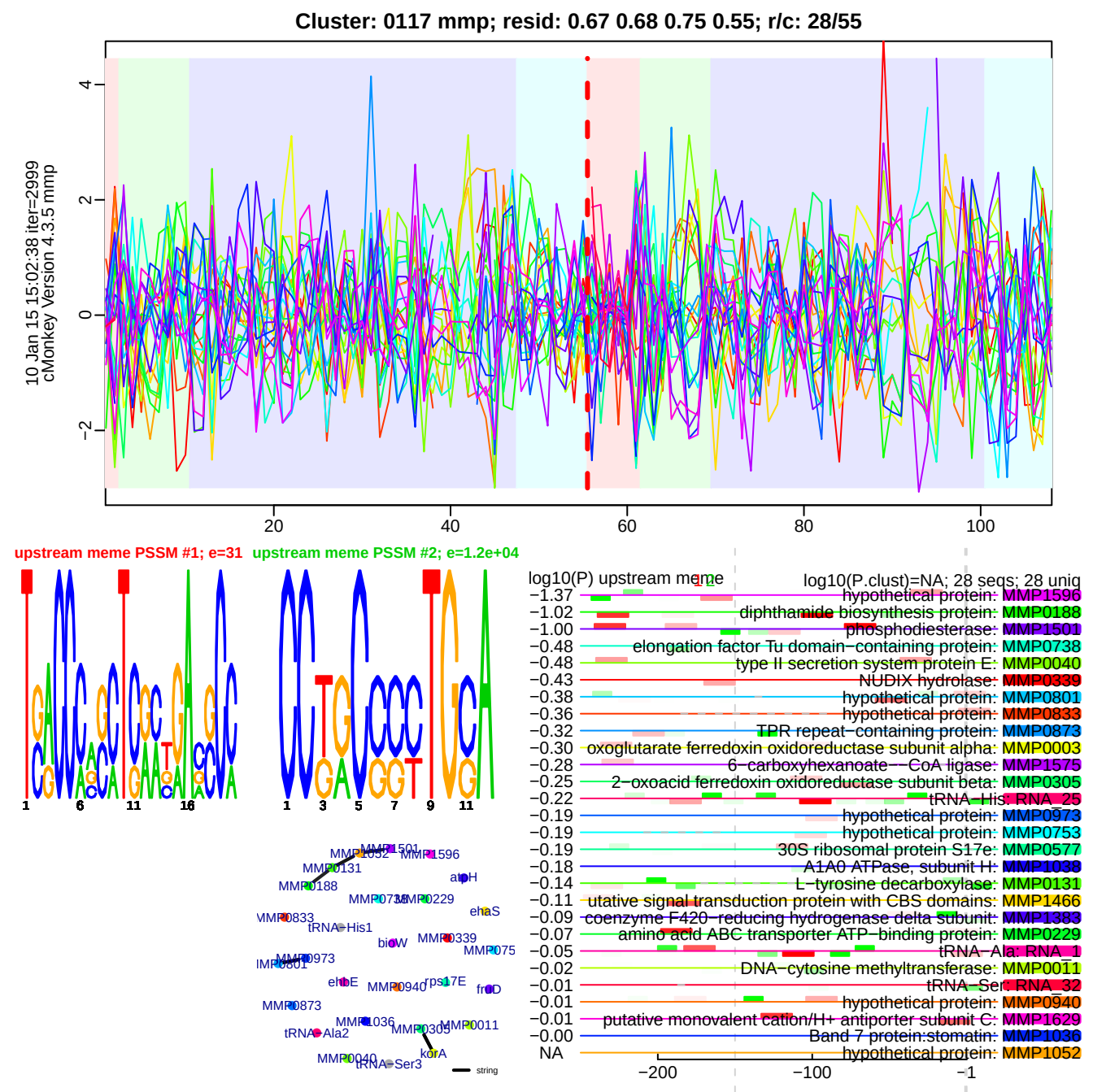
Cluster: 0117 mmp; resid: 0.67 0.68 0.75 0.55; r/c: 28/55
10 Jan 15 15:02:38 iter=2999
cMonkey Version 4.3.5 mmp
upstream meme PSSM #1; e=31
upstream meme PSSM #2; e=1.2e+04
log10(P) upstream meme
log10(P.clust)=NA; 28 seqs; 28 uniq
−1.37 hypothetical protein: MMP1596
−1.02 diphthamide biosynthesis protein: MMP0188
−1.00 phosphodiesterase: MMP1501
−0.48 elongation factor Tu domain−containing protein: MMP0738
−0.48 type II secretion system protein E: MMP0040
−0.43 NUDIX hydrolase: MMP0339
−0.38 hypothetical protein: MMP0801
−0.36 hypothetical protein: MMP0833
−0.32 TPR repeat−containing protein: MMP0873
−0.30 oxoglutarate ferredoxin oxidoreductase subunit alpha: MMP0003
−0.28 6−carboxyhexanoate−−CoA ligase: MMP1575
−0.25 2−oxoacid ferredoxin oxidoreductase subunit beta: MMP0305
−0.22 tRNA−His: RNA_25
−0.19 hypothetical protein: MMP0973
−0.19 hypothetical protein: MMP0753
−0.19 30S ribosomal protein S17e: MMP0577
−0.18 A1A0 ATPase, subunit H: MMP1038
−0.14 L−tyrosine decarboxylase: MMP0131
−0.11 utative signal transduction protein with CBS domains: MMP1466
−0.09 coenzyme F420−reducing hydrogenase delta subunit: MMP1383
−0.07 amino acid ABC transporter ATP−binding protein: MMP0229
−0.05 tRNA−Ala: RNA_1
−0.02 DNA−cytosine methyltransferase: MMP0011
−0.01 tRNA−Ser: RNA_32
−0.01 hypothetical protein: MMP0940
−0.01 putative monovalent cation/H+ antiporter subunit C: MMP1629
−0.00 Band 7 protein:stomatin: MMP1036
NA hypothetical protein: MMP1052
MMP1501
MMP1052
MMP1596
MMP0131
atpH
MMP0188
MMP0738
MMP0229
MMP0833
ehaS
tRNA−His1
bioW
MMP0339
MMP0075
MMP0973
MMP0801
ehbE
MMP0940
rps17E
fmdD
MMP0873
MMP1036
MMP0011
tRNA−Ala2
MMP0305
korA
MMP0040
tRNA−Ser3
string