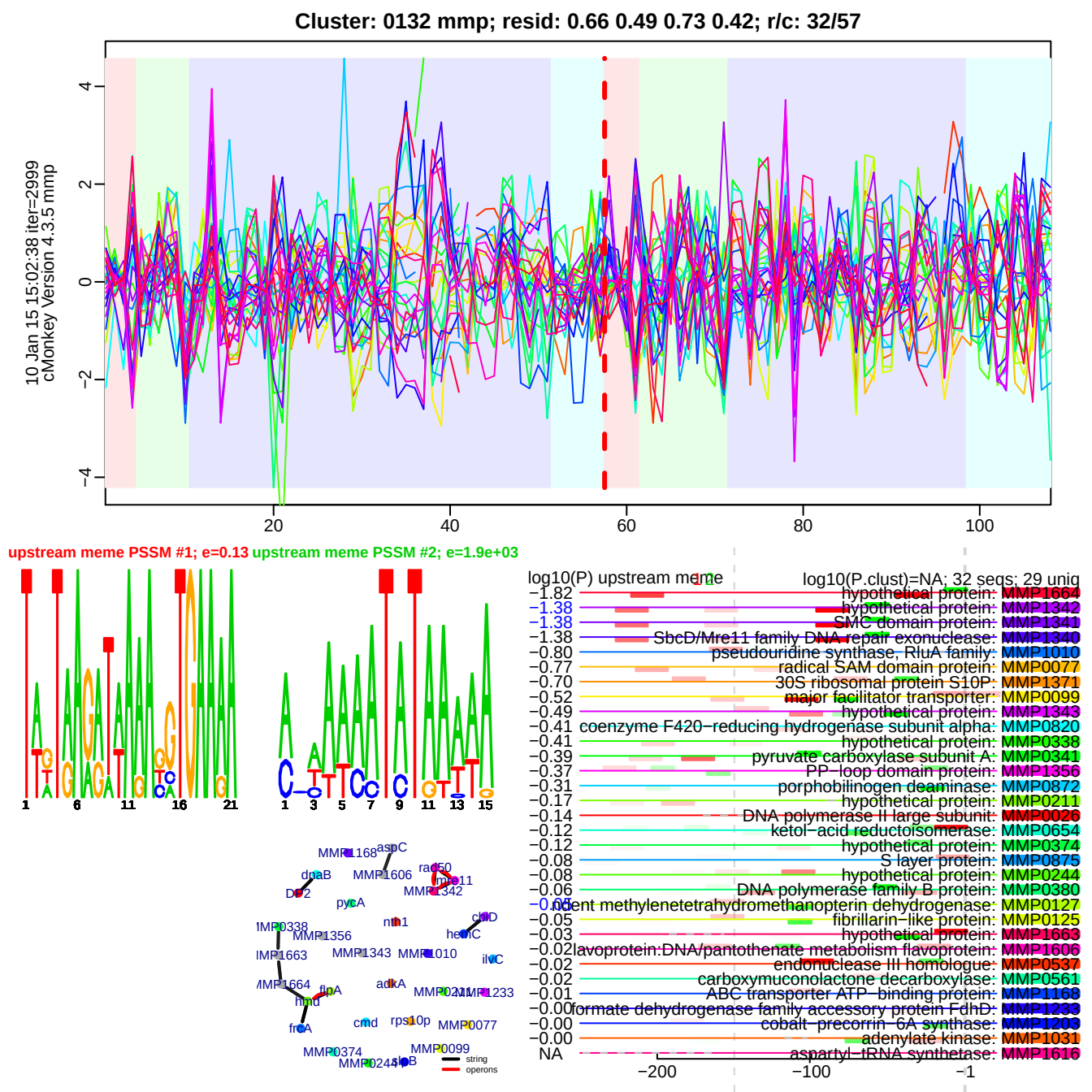

Cluster: 0132 mmp; resid: 0.66 0.49 0.73 0.42; r/c: 32/57
10 Jan 15 15:02:38 iter=2999
cMonkey Version 4.3.5 mmp
upstream meme PSSM #1; e=0.13
upstream meme PSSM #2; e=1.9e+03
log10(P) upstream meme
log10(P.clust)=NA; 32 seqs; 29 uniq
hypothetical protein: MMP1664
hypothetical protein: MMP1342
SMC domain protein: MMP1341
SbcD/Mre11 family DNA repair exonuclease: MMP1340
pseudouridine synthase, RluA family: MMP1010
radical SAM domain protein: MMP0077
30S ribosomal protein S10P: MMP1371
major facilitator transporter: MMP0099
hypothetical protein: MMP1343
coenzyme F420−reducing hydrogenase subunit alpha: MMP0820
hypothetical protein: MMP0338
pyruvate carboxylase subunit A: MMP0341
PP−loop domain protein: MMP1356
porphobilinogen deaminase: MMP0872
hypothetical protein: MMP0211
DNA polymerase II large subunit: MMP0026
ketol−acid reductoisomerase: MMP0654
hypothetical protein: MMP0374
S layer protein: MMP0875
hypothetical protein: MMP0244
DNA polymerase family B protein: MMP0380
ident methylenetetrahydromethanopterin dehydrogenase: MMP0127
fibrillarin−like protein: MMP0125
hypothetical protein: MMP1663
lavoprotein:DNA/pantothenate metabolism flavoprotein: MMP1606
endonuclease III homologue: MMP0537
carboxymuconolactone decarboxylase: MMP0561
ABC transporter ATP−binding protein: MMP1168
formate dehydrogenase family accessory protein FdhD: MMP1233
cobalt−precorrin−6A synthase: MMP1203
adenylate kinase: MMP1031
aspartyl−tRNA synthetase: MMP1616
string
operons