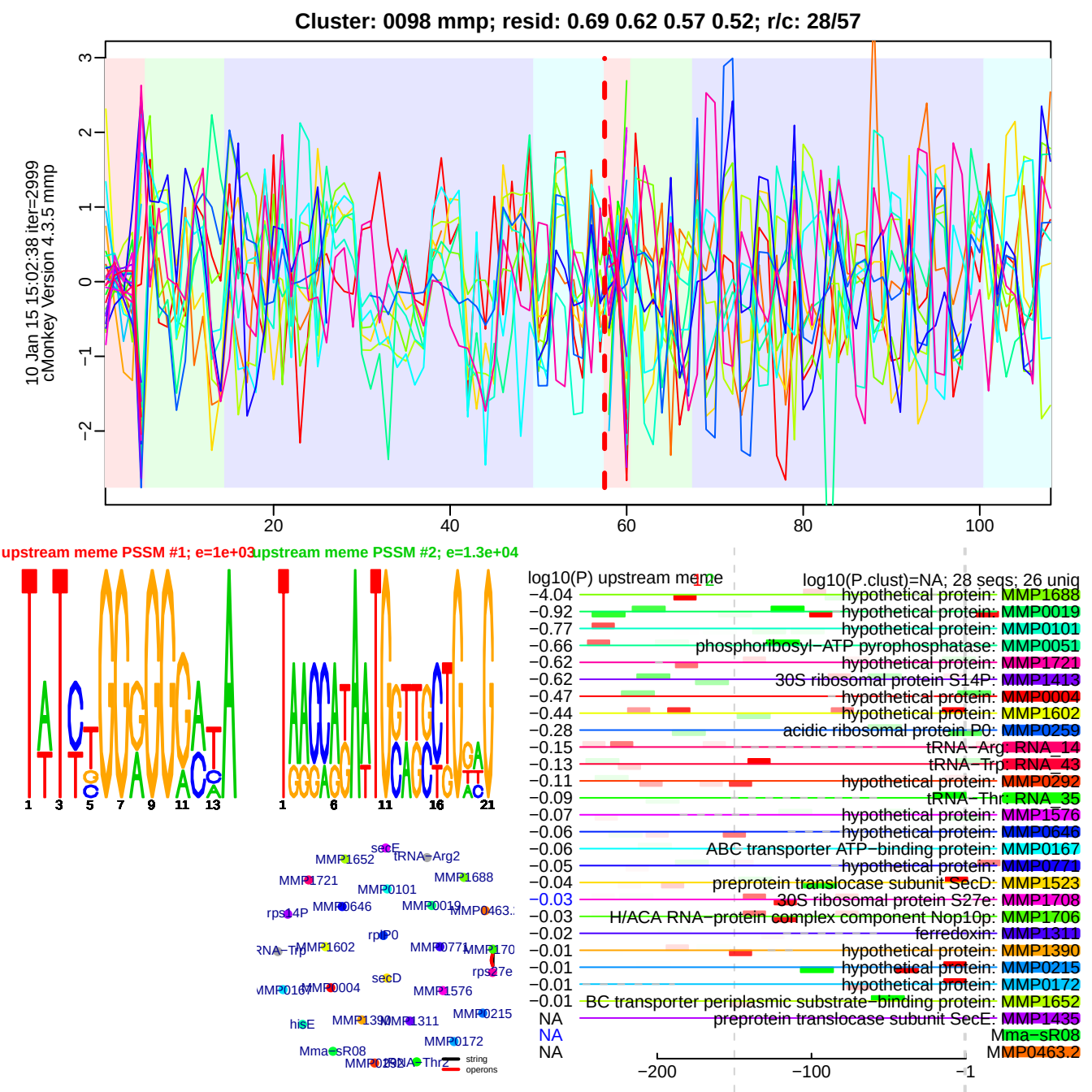
Cluster: 0098 mmp; resid: 0.69 0.62 0.57 0.52; r/c: 28/57
10 Jan 15 15:02:38 iter=2999
cMonkey Version 4.3.5 mmp
upstream meme PSSM #1; e=1e+03
upstream meme PSSM #2; e=1.3e+04
log10(P) upstream meme
log10(P.clust)=NA; 28 seqs; 26 uniq
−4.04 hypothetical protein: MMP1688
−0.92 hypothetical protein: MMP0019
−0.77 hypothetical protein: MMP0101
−0.66 phosphoribosyl−ATP pyrophosphatase: MMP0051
−0.62 hypothetical protein: MMP1721
−0.62 30S ribosomal protein S14P: MMP1413
−0.47 hypothetical protein: MMP0004
−0.44 hypothetical protein: MMP1602
−0.28 acidic ribosomal protein P0: MMP0259
−0.15 tRNA−Arg: RNA_14
−0.13 tRNA−Trp: RNA_43
−0.11 hypothetical protein: MMP0292
−0.09 tRNA−Thr: RNA_35
−0.07 hypothetical protein: MMP1576
−0.06 hypothetical protein: MMP0646
−0.06 ABC transporter ATP−binding protein: MMP0167
−0.05 hypothetical protein: MMP0771
−0.04 preprotein translocase subunit SecD: MMP1523
−0.03 30S ribosomal protein S27e: MMP1708
−0.03 H/ACA RNA−protein complex component Nop10p: MMP1706
−0.02 ferredoxin: MMP1311
−0.01 hypothetical protein: MMP1390
−0.01 hypothetical protein: MMP0215
−0.01 hypothetical protein: MMP0172
−0.01 BC transporter periplasmic substrate−binding protein: MMP1652
NA preprotein translocase subunit SecE: MMP1435
NA Mma−sR08
NA MMP0463.2
−200
−100
−1
secE
tRNA−Arg2
MMP1652
MMP1721
MMP1688
MMP0101
rps14P
MMP0646
MMP0019
rplP0
tRNA−Trp
MMP1602
MMP0771
rps27e
secD
MMP0004
MMP1576
hisE
MMP1390
MMP1311
MMP0215
MMP0172
Mma−sR08
tRNA−Thr2
string
operons