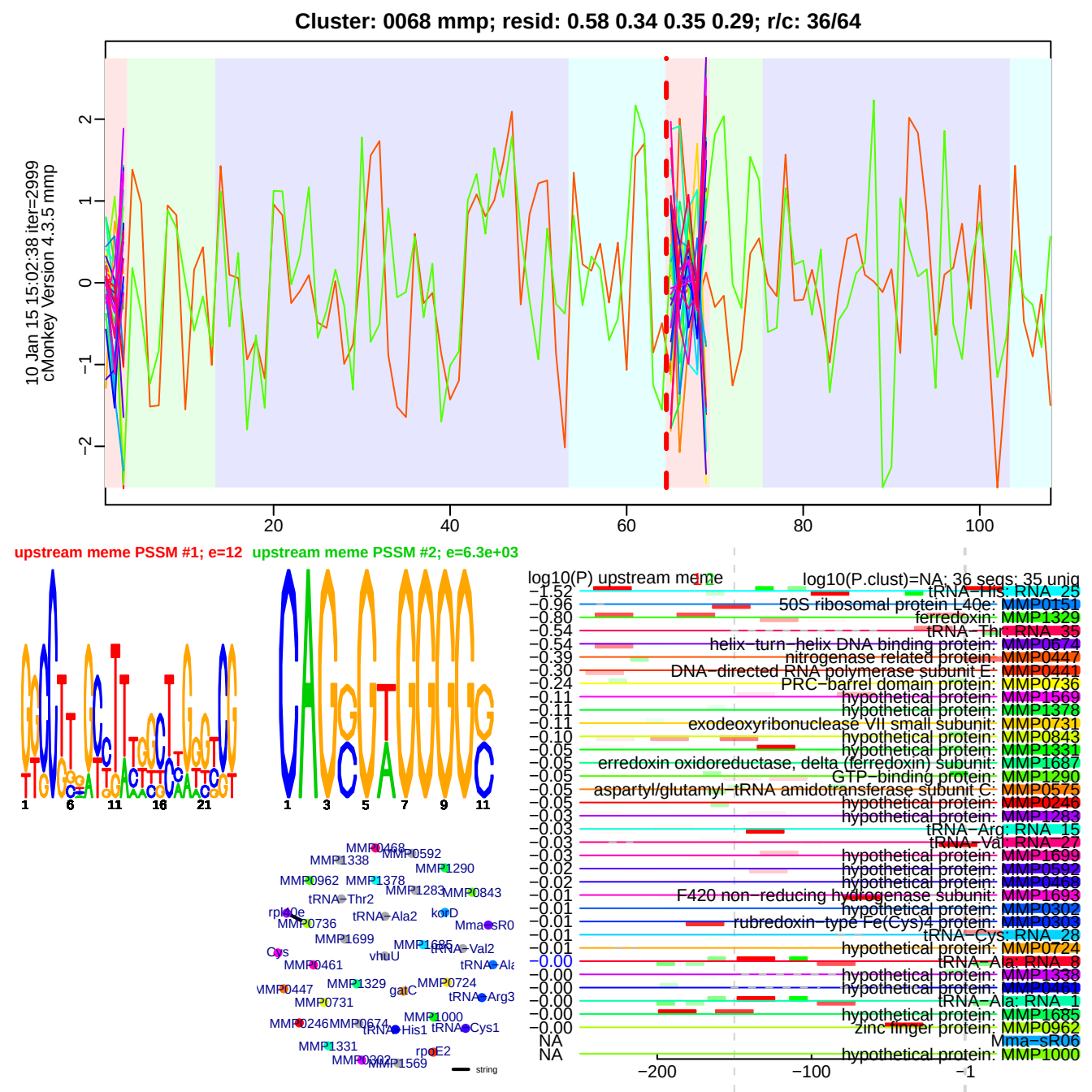
Cluster: 0068 mmp; resid: 0.58 0.34 0.35 0.29; r/c: 36/64
10 Jan 15 15:02:38 iter=2999
cMonkey Version 4.3.5 mmp
upstream meme PSSM #1; e=12
upstream meme PSSM #2; e=6.3e+03
log10(P) upstream meme
log10(P.clust)=NA; 36 segs; 35 uniq
tRNA−His: RNA_25
50S ribosomal protein L40e: MMP0151
ferredoxin: MMP1329
tRNA−Thr: RNA_35
helix−turn−helix DNA binding protein: MMP0674
nitrogenase related protein: MMP0447
DNA−directed RNA polymerase subunit E: MMP0441
PRC−barrel domain protein: MMP0736
hypothetical protein: MMP1569
hypothetical protein: MMP1378
exodeoxyribonuclease VII small subunit: MMP0731
hypothetical protein: MMP0843
hypothetical protein: MMP1331
erredoxin oxidoreductase, delta (ferredoxin) subunit: MMP1687
GTP−binding protein: MMP1290
aspartyl/glutamyl−tRNA amidotransferase subunit C: MMP0575
hypothetical protein: MMP0246
hypothetical protein: MMP1283
tRNA−Arg: RNA_15
tRNA−Val: RNA_27
hypothetical protein: MMP1699
hypothetical protein: MMP0592
hypothetical protein: MMP0468
F420 non−reducing hydrogenase subunit: MMP1693
hypothetical protein: MMP0302
rubredoxin−type Fe(Cys)4 protein: MMP0303
tRNA−Cys: RNA_28
hypothetical protein: MMP0724
tRNA−Ala: RNA_8
hypothetical protein: MMP1338
hypothetical protein: MMP0461
tRNA−Ala: RNA_1
hypothetical protein: MMP1685
zinc finger protein: MMP0962
Mma−sR06
hypothetical protein: MMP1000
MMP0468
MMP0592
MMP1338
MMP1290
MMP0962
MMP1378
MMP1283
MMP0843
tRNA−Thr2
rplQe
tRNA−Ala2
kdpD
MMP0736
MmasR0
MMP1699
MMP1685
Cys
vhuU
tRNA−Val2
MMP0461
tRNA−Ala
MMP0447
MMP1329
gatC
MMP0724
MMP0731
tRNA−Arg3
MMP0246
MMP0674
MMP1000
tRNA−His1
tRNA−Cys1
MMP1331
rpoE2
MMP0302
MMP1569
string