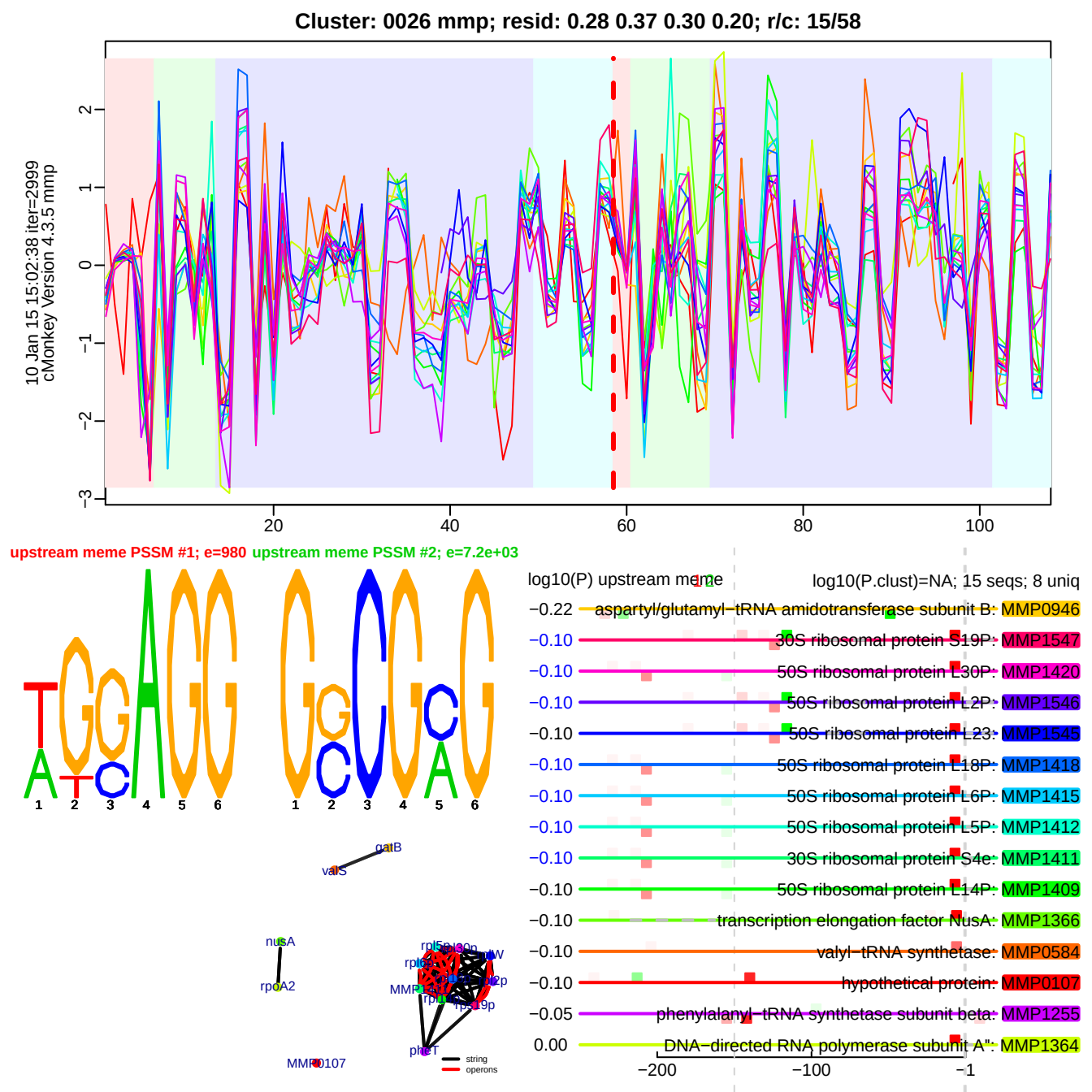
Cluster: 0026 mmp; resid: 0.28 0.37 0.30 0.20; r/c: 15/58
10 Jan 15 15:02:38 iter=2999
cMonkey Version 4.3.5 mmp
upstream meme PSSM #1; e=980
upstream meme PSSM #2; e=7.2e+03
log10(P) upstream meme
log10(P.clust)=NA; 15 seqs; 8 uniq
−0.22 aspartyl/glutamyl−tRNA amidotransferase subunit B: MMP0946
−0.10 30S ribosomal protein S19P: MMP1547
−0.10 50S ribosomal protein L30P: MMP1420
−0.10 50S ribosomal protein L2P: MMP1546
−0.10 50S ribosomal protein L23: MMP1545
−0.10 50S ribosomal protein L18P: MMP1418
−0.10 50S ribosomal protein L6P: MMP1415
−0.10 50S ribosomal protein L5P: MMP1412
−0.10 30S ribosomal protein S4e: MMP1411
−0.10 50S ribosomal protein L14P: MMP1409
−0.10 transcription elongation factor NusA: MMP1366
−0.10 valyl−tRNA synthetase: MMP0584
−0.10 hypothetical protein: MMP0107
−0.05 phenylalanyl−tRNA synthetase subunit beta: MMP1255
0.00 DNA−directed RNA polymerase subunit A": MMP1364
gatB
valS
nusA
rpoA2
MMP0107
pheT
string
operons