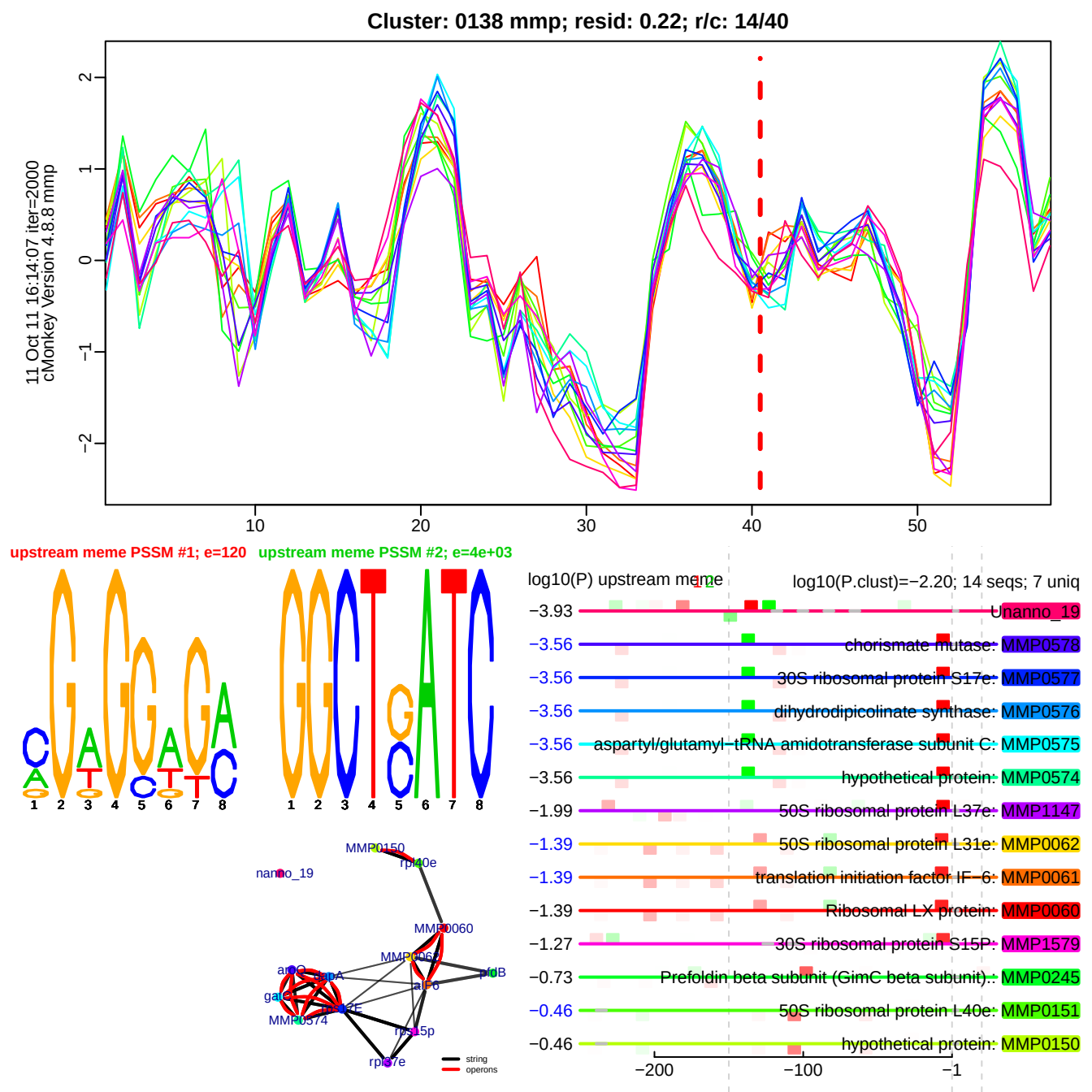
Cluster: 0138 mmp; resid: 0.22; r/c: 14/40
11 Oct 11 16:14:07 iter=2000
cMonkey Version 4.8.8 mmp
upstream meme PSSM #1; e=120
upstream meme PSSM #2; e=4e+03
log10(P) upstream meme
log10(P.clust)=−2.20; 14 seqs; 7 uniq
−3.93 Unanno_19
−3.56 chorismate mutase: MMP0578
−3.56 30S ribosomal protein S17e: MMP0577
−3.56 dihydrodipicolinate synthase: MMP0576
−3.56 aspartyl/glutamyl−tRNA amidotransferase subunit C: MMP0575
−3.56 hypothetical protein: MMP0574
−1.99 50S ribosomal protein L37e: MMP1147
−1.39 50S ribosomal protein L31e: MMP0062
−1.39 translation initiation factor IF−6: MMP0061
−1.39 Ribosomal LX protein: MMP0060
−1.27 30S ribosomal protein S15P: MMP1579
−0.73 Prefoldin beta subunit (GimC beta subunit).: MMP0245
−0.46 50S ribosomal protein L40e: MMP0151
−0.46 hypothetical protein: MMP0150
MMP0150
rpl40e
nanno_19
MMP0060
MMP0061
aIF6
pfdB
MMP0574
rpl15p
rpl37e
string
operons