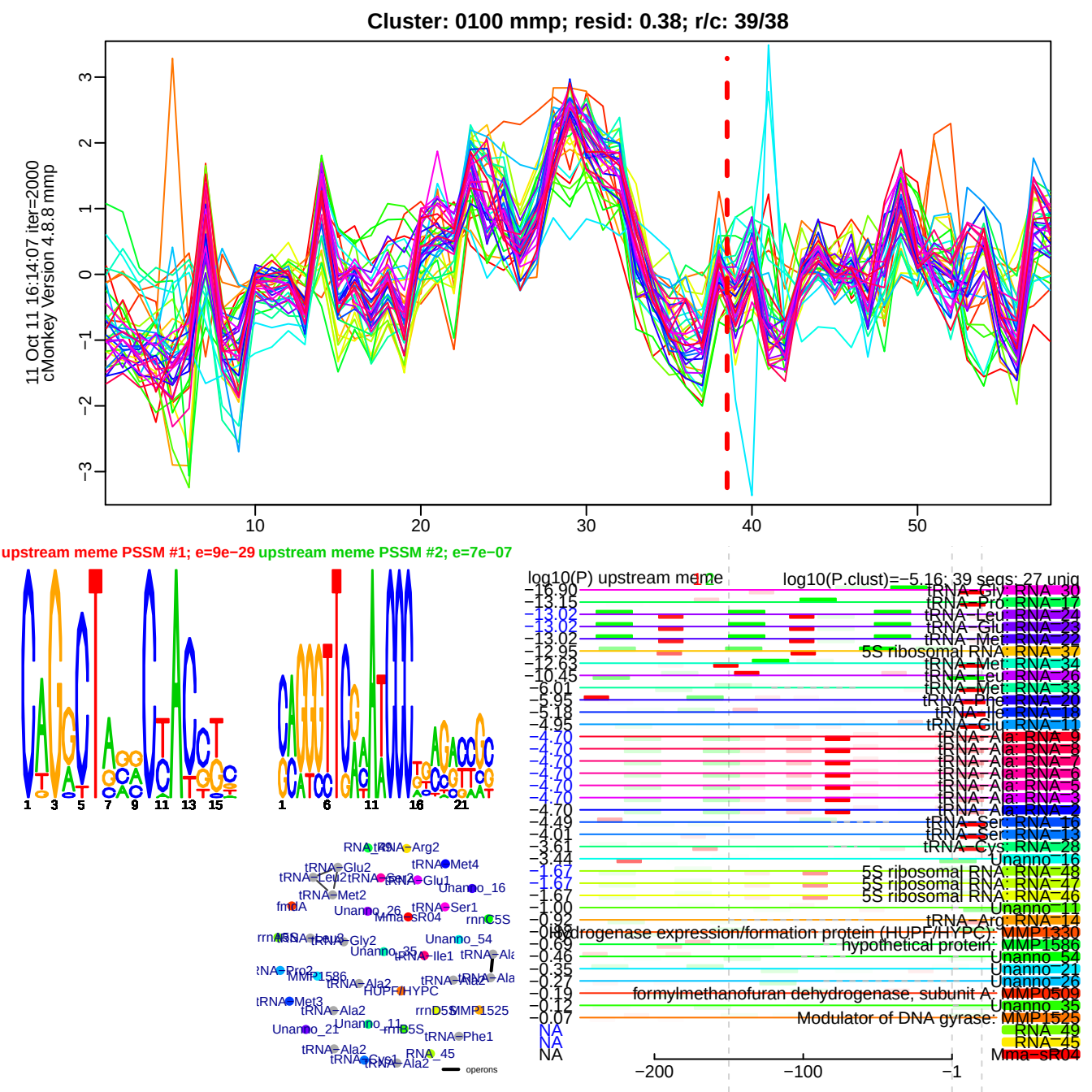
Cluster: 0100 mmp; resid: 0.38; r/c: 39/38
11 Oct 11 16:14:07 iter=2000
cMonkey Version 4.8.8 mmp
upstream meme PSSM #1; e=9e−29
upstream meme PSSM #2; e=7e−07
log10(P) upstream meme
log10(P.clust)=−5.16; 39 seqs; 27 uniq
tRNA-Gly: RNA_30
tRNA-Pro: RNA_17
tRNA-Leu: RNA_24
tRNA-Glu: RNA_23
tRNA-Met: RNA_22
5S ribosomal RNA: RNA_37
tRNA-Met: RNA_34
tRNA-Leu: RNA_26
tRNA-Met: RNA_33
tRNA-Phe: RNA_20
tRNA-Ile: RNA_18
tRNA-Glu: RNA_11
tRNA-Ala: RNA_9
tRNA-Ala: RNA_8
tRNA-Ala: RNA_7
tRNA-Ala: RNA_6
tRNA-Ala: RNA_5
tRNA-Ala: RNA_3
tRNA-Ala: RNA_2
tRNA-Ser: RNA_16
tRNA-Ser: RNA_13
tRNA-Cys: RNA_28
Unanno_16
5S ribosomal RNA: RNA_48
5S ribosomal RNA: RNA_47
5S ribosomal RNA: RNA_46
Unanno_11
tRNA-Arg: RNA_14
hydrogenase expression/formation protein (HUPF/HYPC): MMP1330
hypothetical protein: MMP1586
Unanno_54
Unanno_21
Unanno_26
formylmethanofuran dehydrogenase, subunit A: MMP0509
Unanno_35
Modulator of DNA gyrase: MMP1525
RNA_49
RNA_45
Mma-sR04
operons