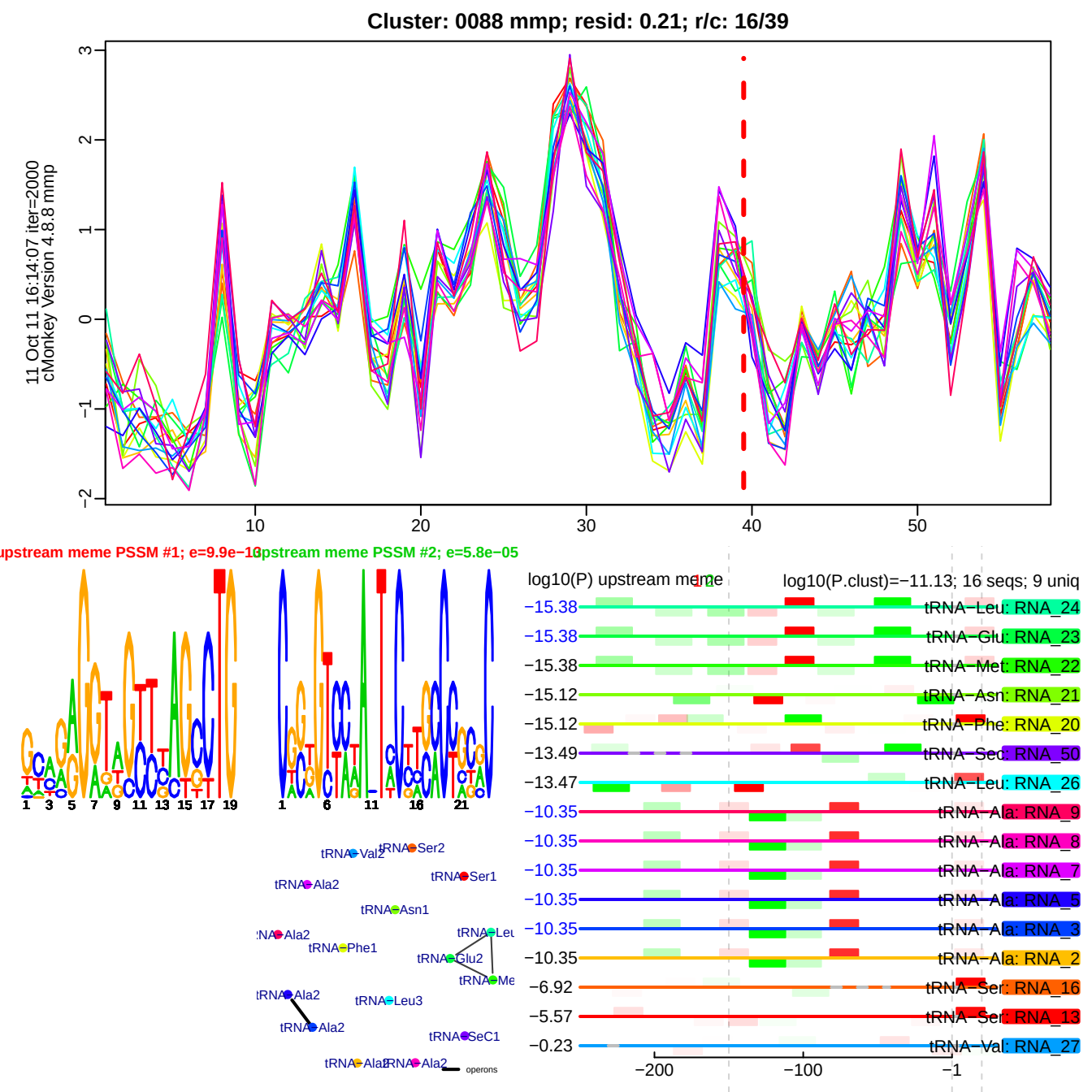
Cluster: 0088 mmp; resid: 0.21; r/c: 16/39
11 Oct 11 16:14:07 iter=2000
cMonkey Version 4.8.8 mmp
upstream meme PSSM #1; e=9.9e−16
upstream meme PSSM #2; e=5.8e−05
log10(P) upstream meme
log10(P.clust)=−11.13; 16 seqs; 9 uniq
−15.38 tRNA−Leu: RNA_24
−15.38 tRNA−Glu: RNA_23
−15.38 tRNA−Met: RNA_22
−15.12 tRNA−Asn: RNA_21
−15.12 tRNA−Phe: RNA_20
−13.49 tRNA−Sec: RNA_50
−13.47 tRNA−Leu: RNA_26
−10.35 tRNA−Ala: RNA_9
−10.35 tRNA−Ala: RNA_8
−10.35 tRNA−Ala: RNA_7
−10.35 tRNA−Ala: RNA_5
−10.35 tRNA−Ala: RNA_3
−10.35 tRNA−Ala: RNA_2
−6.92 tRNA−Ser: RNA_16
−5.57 tRNA−Ser: RNA_13
−0.23 tRNA−Val: RNA_27
−200
−100
−1
tRNA−Val2
tRNA−Ser2
tRNA−Ser1
tRNA−Ala2
tRNA−Asn1
tRNA−Phe1
tRNA−Glu2
tRNA−Leu3
tRNA−SeC1
operons