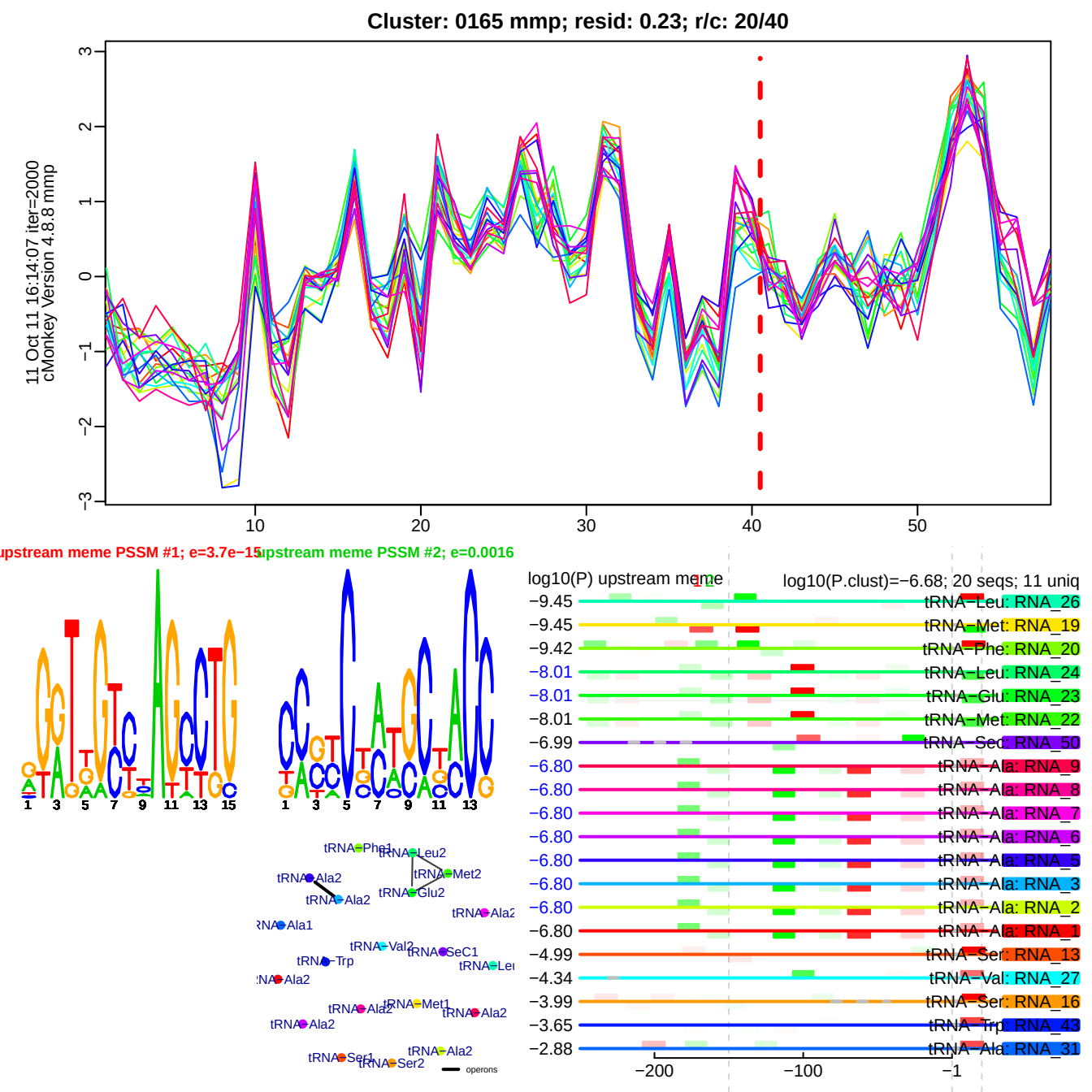
Cluster: 0165 mmp; resid: 0.23; r/c: 20/40
11 Oct 11 16:14:07 iter=2000
cMonkey Version 4.8.8 mmp
upstream meme PSSM #1; e=3.7e−15
upstream meme PSSM #2; e=0.0016
log10(P) upstream meme
log10(P.clust)=−6.68; 20 seqs; 11 uniq
−9.45 tRNA−Leu: RNA_26
−9.45 tRNA−Met: RNA_19
−9.42 tRNA−Phe: RNA_20
−8.01 tRNA−Leu: RNA_24
−8.01 tRNA−Glu: RNA_23
−8.01 tRNA−Met: RNA_22
−6.99 tRNA−Sec: RNA_50
−6.80 tRNA−Ala: RNA_9
−6.80 tRNA−Ala: RNA_8
−6.80 tRNA−Ala: RNA_7
−6.80 tRNA−Ala: RNA_6
−6.80 tRNA−Ala: RNA_5
−6.80 tRNA−Ala: RNA_3
−6.80 tRNA−Ala: RNA_2
−6.80 tRNA−Ala: RNA_1
−4.99 tRNA−Ser: RNA_13
−4.34 tRNA−Val: RNA_27
−3.99 tRNA−Ser: RNA_16
−3.65 tRNA−Trp: RNA_43
−2.88 tRNA−Ala: RNA_31
−200
−100
−1
tRNA−Phe1
tRNA−Leu2
tRNA−Ala2
tRNA−Met2
tRNA−Glu2
tRNA−Ala1
tRNA−Val2
tRNA−SeC1
tRNA−Trp
tRNA−Met1
tRNA−Ser1
tRNA−Ser2
operons