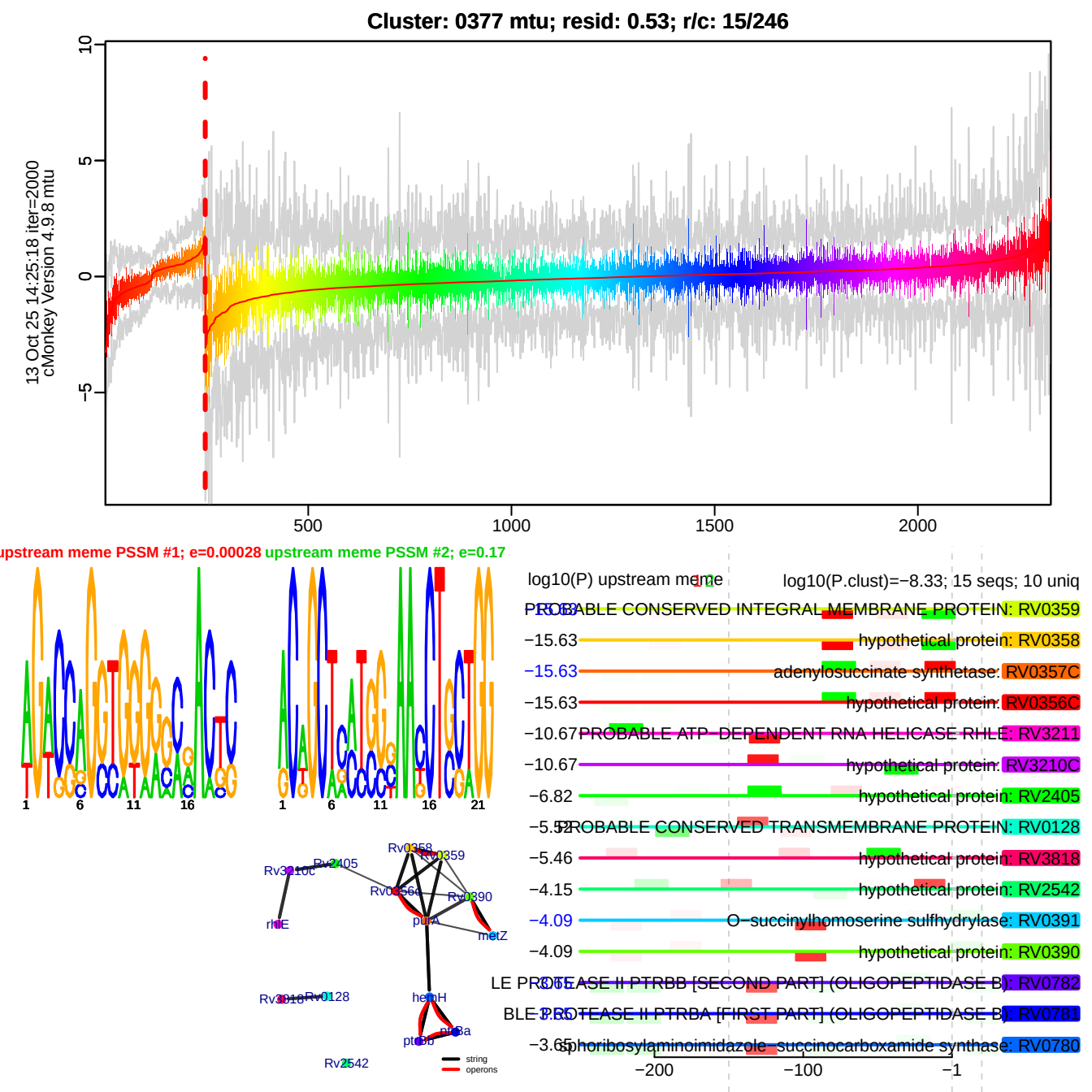
Cluster: 0377 mtu; resid: 0.53; r/c: 15/246
13 Oct 25 14:25:18 iter=2000
cMonkey Version 4.9.8 mtu
upstream meme PSSM #1; e=0.00028
upstream meme PSSM #2; e=0.17
log10(P) upstream meme
log10(P.clust)=−8.33; 15 seqs; 10 uniq
PROBABLE CONSERVED INTEGRAL MEMBRANE PROTEIN: RV0359
hypothetical protein: RV0358
adenylosuccinate synthetase: RV0357C
hypothetical protein: RV0356C
PROBABLE ATP−DEPENDENT RNA HELICASE RHLE: RV3211
hypothetical protein: RV3210C
hypothetical protein: RV2405
PROBABLE CONSERVED TRANSMEMBRANE PROTEIN: RV0128
hypothetical protein: RV3818
hypothetical protein: RV2542
O−succinylhomoserine sulfhydrylase: RV0391
hypothetical protein: RV0390
OLIGOPEPTIDASE B): RV0782
OLIGOPEPTIDASE B): RV0781
phoribosylaminoimidazole−succinocarboxamide synthase: RV0780
string
operons