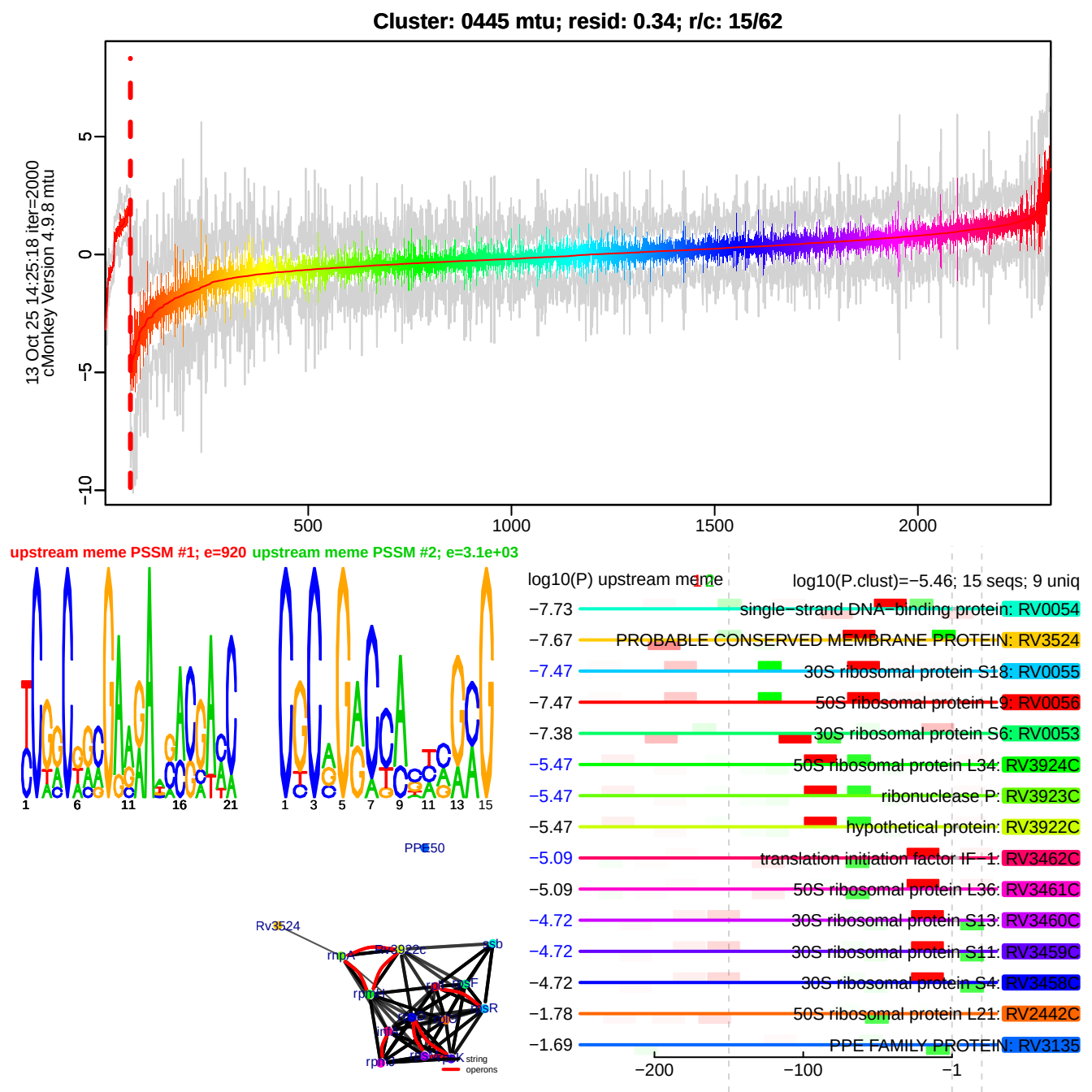
Cluster: 0445 mtu; resid: 0.34; r/c: 15/62
13 Oct 25 14:25:18 iter=2000
cMonkey Version 4.9.8 mtu
upstream meme PSSM #1; e=920
upstream meme PSSM #2; e=3.1e+03
log10(P) upstream meme
log10(P.clust)=−5.46; 15 seqs; 9 uniq
−7.73 single−strand DNA−binding protein: RV0054
−7.67 PROBABLE CONSERVED MEMBRANE PROTEIN: RV3524
−7.47 30S ribosomal protein S18: RV0055
−7.47 50S ribosomal protein L9: RV0056
−7.38 30S ribosomal protein S6: RV0053
−5.47 50S ribosomal protein L34: RV3924C
−5.47 ribonuclease P: RV3923C
−5.47 hypothetical protein: RV3922C
−5.09 translation initiation factor IF−1: RV3462C
−5.09 50S ribosomal protein L36: RV3461C
−4.72 30S ribosomal protein S13: RV3460C
−4.72 30S ribosomal protein S11: RV3459C
−4.72 30S ribosomal protein S4: RV3458C
−1.78 50S ribosomal protein L21: RV2442C
−1.69 PPE FAMILY PROTEIN: RV3135
PPE50
Rv3524
ssb
string
operons