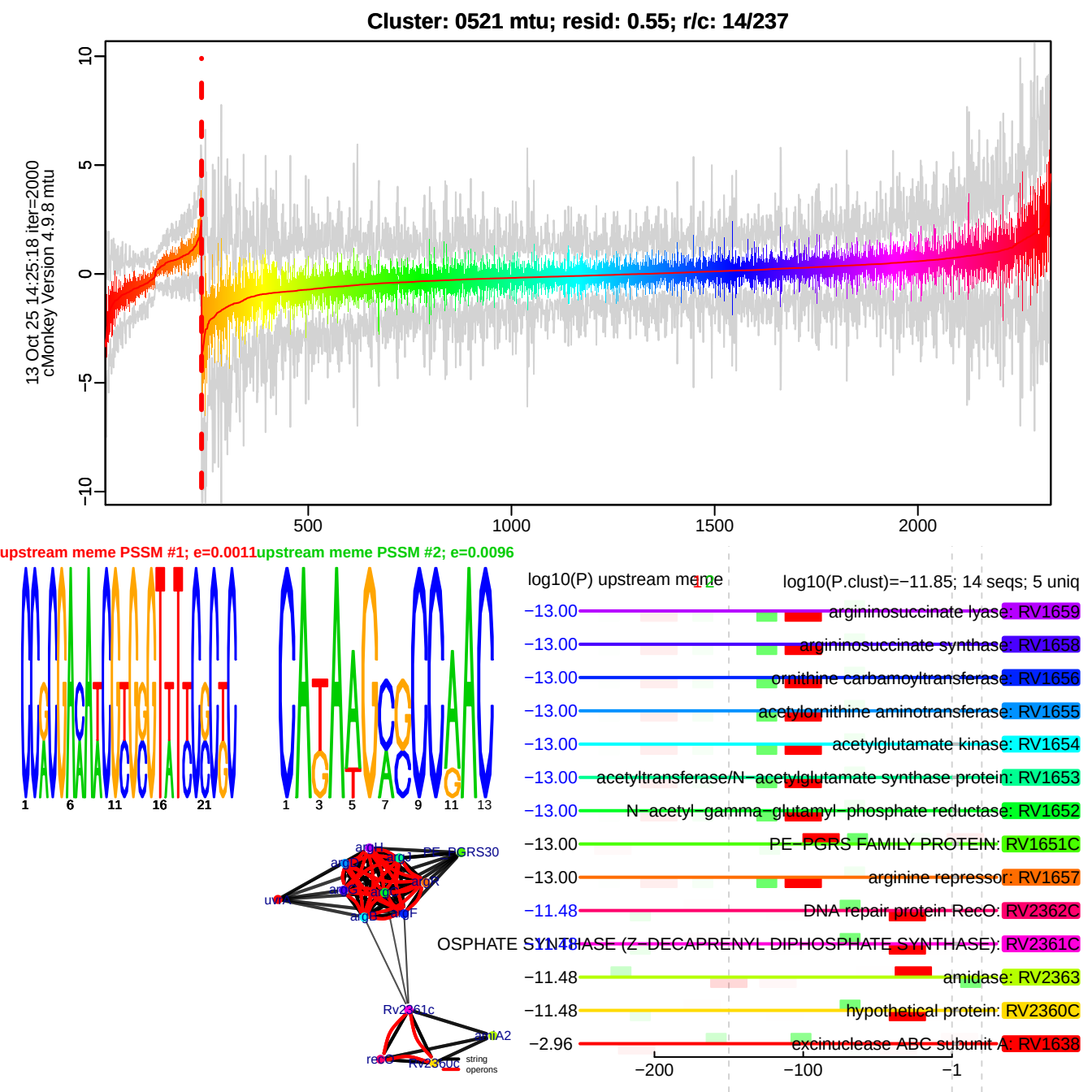
Cluster: 0521 mtu; resid: 0.55; r/c: 14/237
13 Oct 25 14:25:18 iter=2000
cMonkey Version 4.9.8 mtu
upstream meme PSSM #1; e=0.0011
upstream meme PSSM #2; e=0.0096
log10(P) upstream meme 1
log10(P.clust)=−11.85; 14 seqs; 5 uniq
−13.00 argininosuccinate lyase: RV1659
−13.00 argininosuccinate synthase: RV1658
−13.00 ornithine carbamoyltransferase: RV1656
−13.00 acetylornithine aminotransferase: RV1655
−13.00 acetylglutamate kinase: RV1654
−13.00 acetyltransferase/N−acetylglutamate synthase protein: RV1653
−13.00 N−acetyl−gamma−glutamyl−phosphate reductase: RV1652
−13.00 PE−PGRS FAMILY PROTEIN: RV1651C
−13.00 arginine repressor: RV1657
−11.48 DNA repair protein RecO: RV2362C
OSPHATE SYNTHASE (Z−DECAPRENYL DIPHOSPHATE SYNTHASE): RV2361C
−11.48 amidase: RV2363
−11.48 hypothetical protein: RV2360C
−2.96 excinuclease ABC subunit A: RV1638
−200
−100
−1
Rv2361c
recO
Rv2360c
string operons