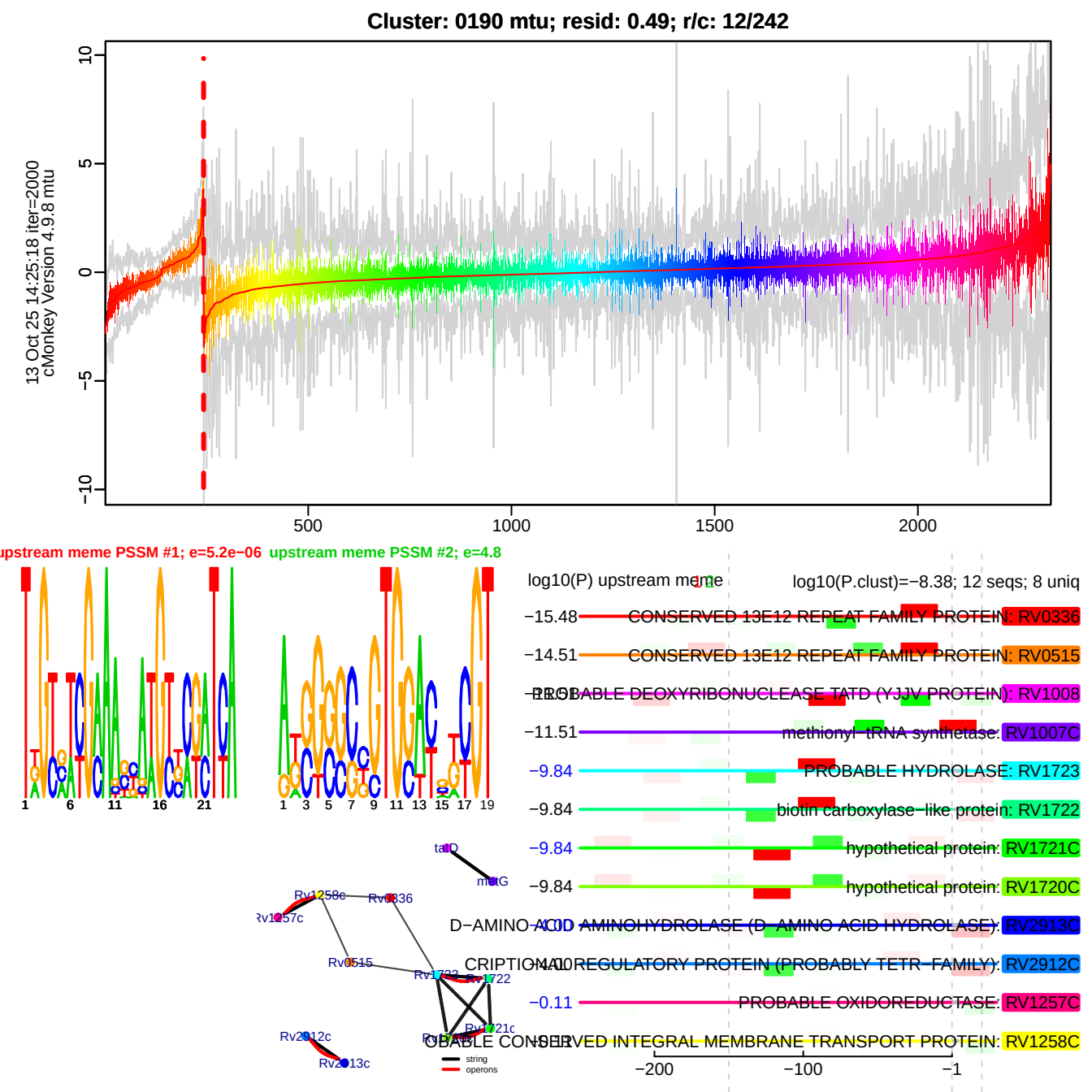
Cluster: 0190 mtu; resid: 0.49; r/c: 12/242
13 Oct 25 14:25:18 iter=2000
cMonkey Version 4.9.8 mtu
upstream meme PSSM #1; e=5.2e−06
upstream meme PSSM #2; e=4.8
log10(P) upstream meme
log10(P.clust)=−8.38; 12 seqs; 8 uniq
−15.48 CONSERVED 13E12 REPEAT FAMILY PROTEIN: RV0336
−14.51 CONSERVED 13E12 REPEAT FAMILY PROTEIN: RV0515
−12.03 PROBABLE DEOXYRIBONUCLEASE TATD (YJJV PROTEIN): RV1008
−11.51 methionyl−tRNA synthetase: RV1007C
−9.84 PROBABLE HYDROLASE: RV1723
−9.84 biotin carboxylase−like protein: RV1722
−9.84 hypothetical protein: RV1721C
−9.84 hypothetical protein: RV1720C
−4.10 D−AMINO ACID AMINOHYDROLASE (D−AMINO ACID HYDROLASE): RV2913C
−4.10 CRIPTION REGULATORY PROTEIN (PROBABLY TETR−FAMILY): RV2912C
−0.11 PROBABLE OXIDOREDUCTASE: RV1257C
−0.13 CONSERVED INTEGRAL MEMBRANE TRANSPORT PROTEIN: RV1258C
−200
−100
−1
tatD
mutG
Rv1258c
Rv0336
Rv1257c
Rv0515
Rv1723
Rv1722
Rv1721c
Rv2912c
Rv2913c
string
operons