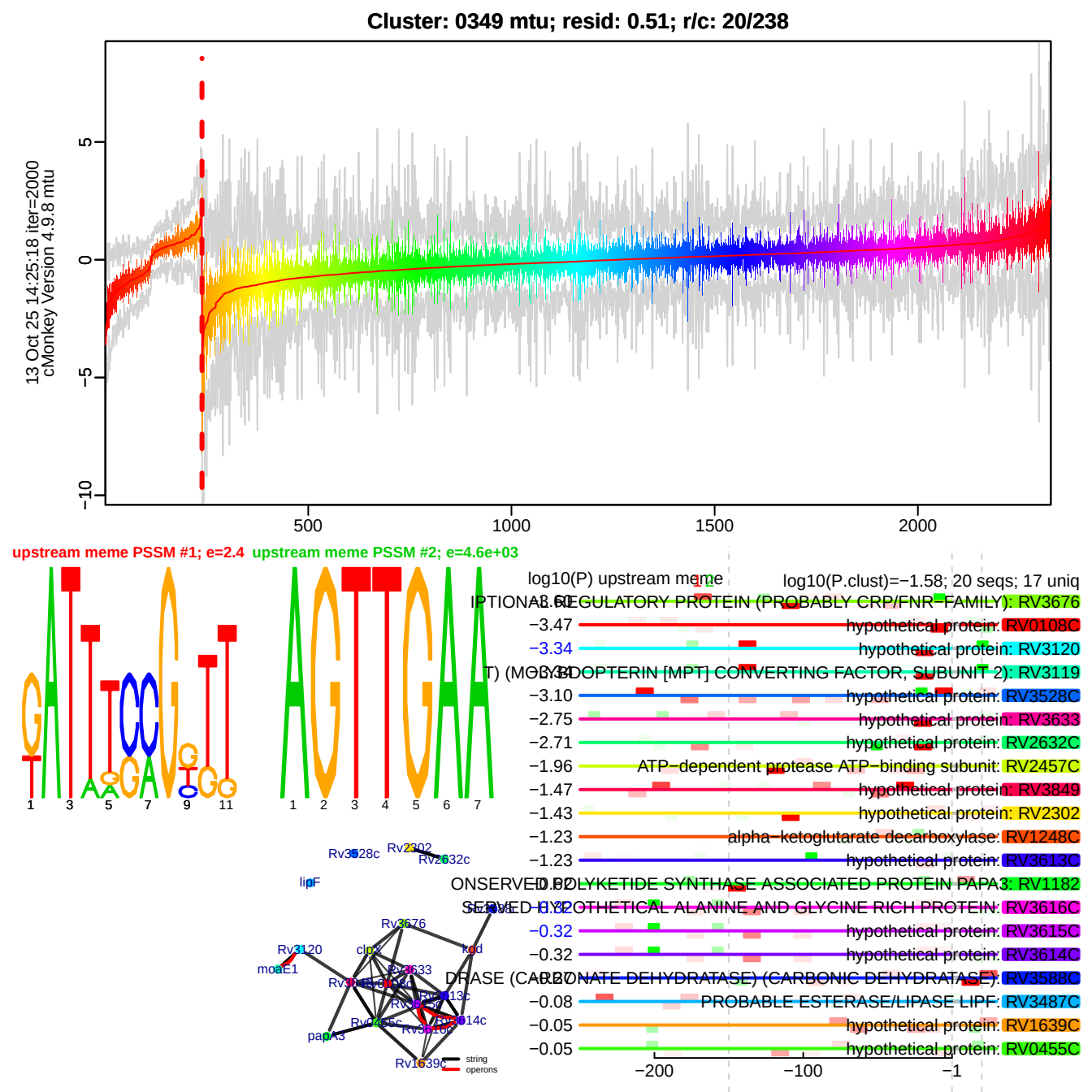
Cluster: 0349 mtu; resid: 0.51; r/c: 20/238
13 Oct 25 14:25:18 iter=2000
cMonkey Version 4.9.8 mtu
upstream meme PSSM #1; e=2.4
upstream meme PSSM #2; e=4.6e+03
log10(P) upstream meme
log10(P.clust)=−1.58; 20 seqs; 17 uniq
−3.47 hypothetical protein: RV0108C
−3.34 hypothetical protein: RV3120
−3.10 hypothetical protein: RV3528C
−2.75 hypothetical protein: RV3633
−2.71 hypothetical protein: RV2632C
−1.96 ATP−dependent protease ATP−binding subunit: RV2457C
−1.47 hypothetical protein: RV3849
−1.43 hypothetical protein: RV2302
−1.23 alpha−ketoglutarate decarboxylase: RV1248C
−1.23 hypothetical protein: RV3613C
−0.32 hypothetical protein: RV3615C
−0.32 hypothetical protein: RV3614C
−0.08 PROBABLE ESTERASE/LIPASE LIPF: RV3487C
−0.05 hypothetical protein: RV1639C
−0.05 hypothetical protein: RV0455C
string
operons