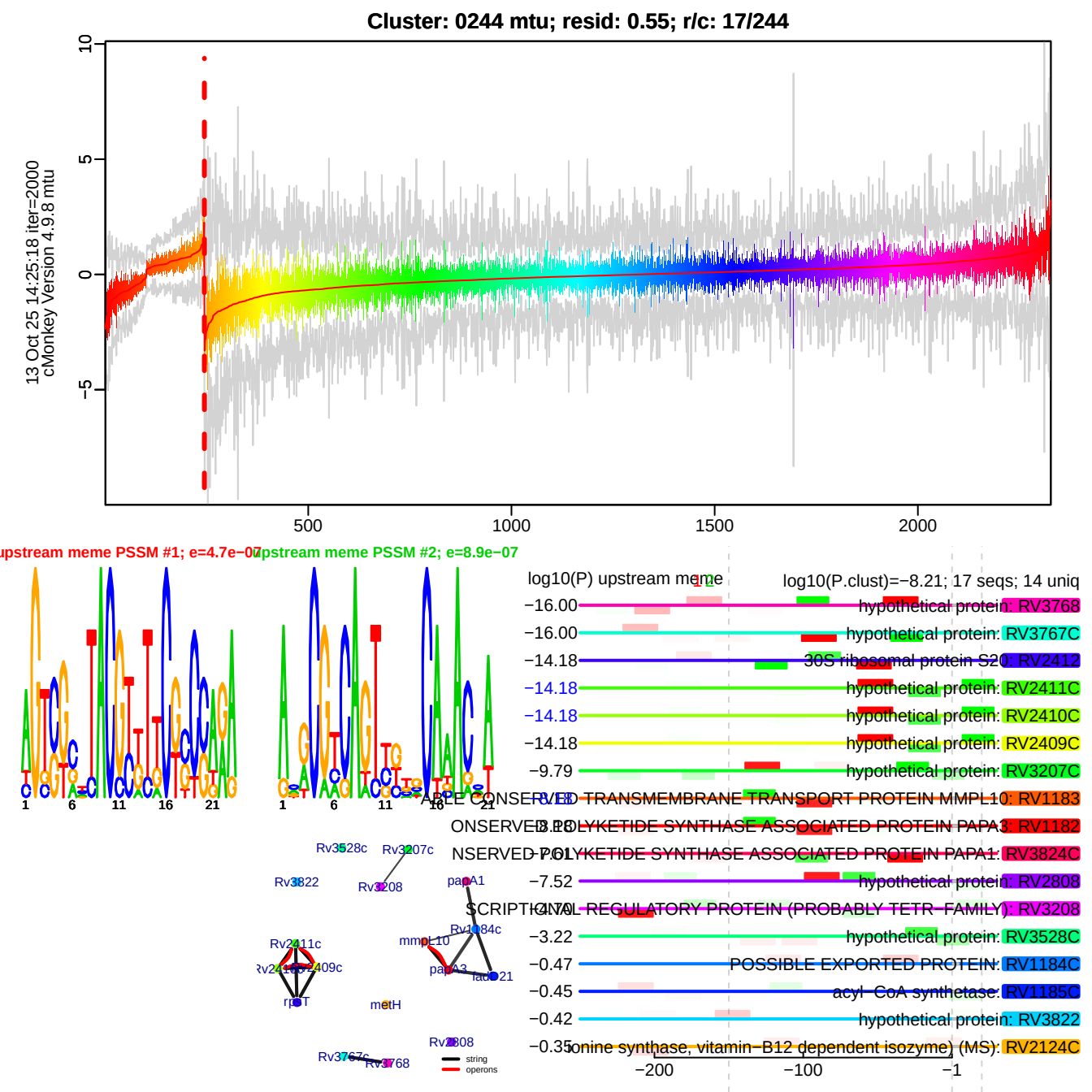
Cluster: 0244 mtu; resid: 0.55; r/c: 17/244
13 Oct 25 14:25:18 iter=2000
cMonkey Version 4.9.8 mtu
upstream meme PSSM #1; e=4.7e−07
upstream meme PSSM #2; e=8.9e−07
log10(P) upstream meme
log10(P.clust)=−8.21; 17 seqs; 14 uniq
−16.00 hypothetical protein: RV3768
−16.00 hypothetical protein: RV3767C
−14.18 30S ribosomal protein S20: RV2412
−14.18 hypothetical protein: RV2411C
−14.18 hypothetical protein: RV2410C
−14.18 hypothetical protein: RV2409C
−9.79 hypothetical protein: RV3207C
ABLE CONSERVED TRANSMEMBRANE TRANSPORT PROTEIN MMPL10: RV1183
ONSERVED POLYKETIDE SYNTHASE ASSOCIATED PROTEIN PAPA3: RV1182
NSERVED POLYKETIDE SYNTHASE ASSOCIATED PROTEIN PAPA1: RV3824C
−7.52 hypothetical protein: RV2808
SCRIPTIONAL REGULATORY PROTEIN (PROBABLY TETR−FAMILY): RV3208
−3.22 hypothetical protein: RV3528C
−0.47 POSSIBLE EXPORTED PROTEIN: RV1184C
−0.45 acyl−CoA synthetase: RV1185C
−0.42 hypothetical protein: RV3822
−0.35 onine synthase, vitamin−B12 dependent isozyme) (MS): RV2124C
−200
−100
−1
Rv3528c
Rv3207c
Rv3822
Rv3208
papA1
Rv1184c
mmpL10
papA3
fadD21
Rv2411c
rplT
metH
Rv2808
Rv3767c
Rv3768
string
operons