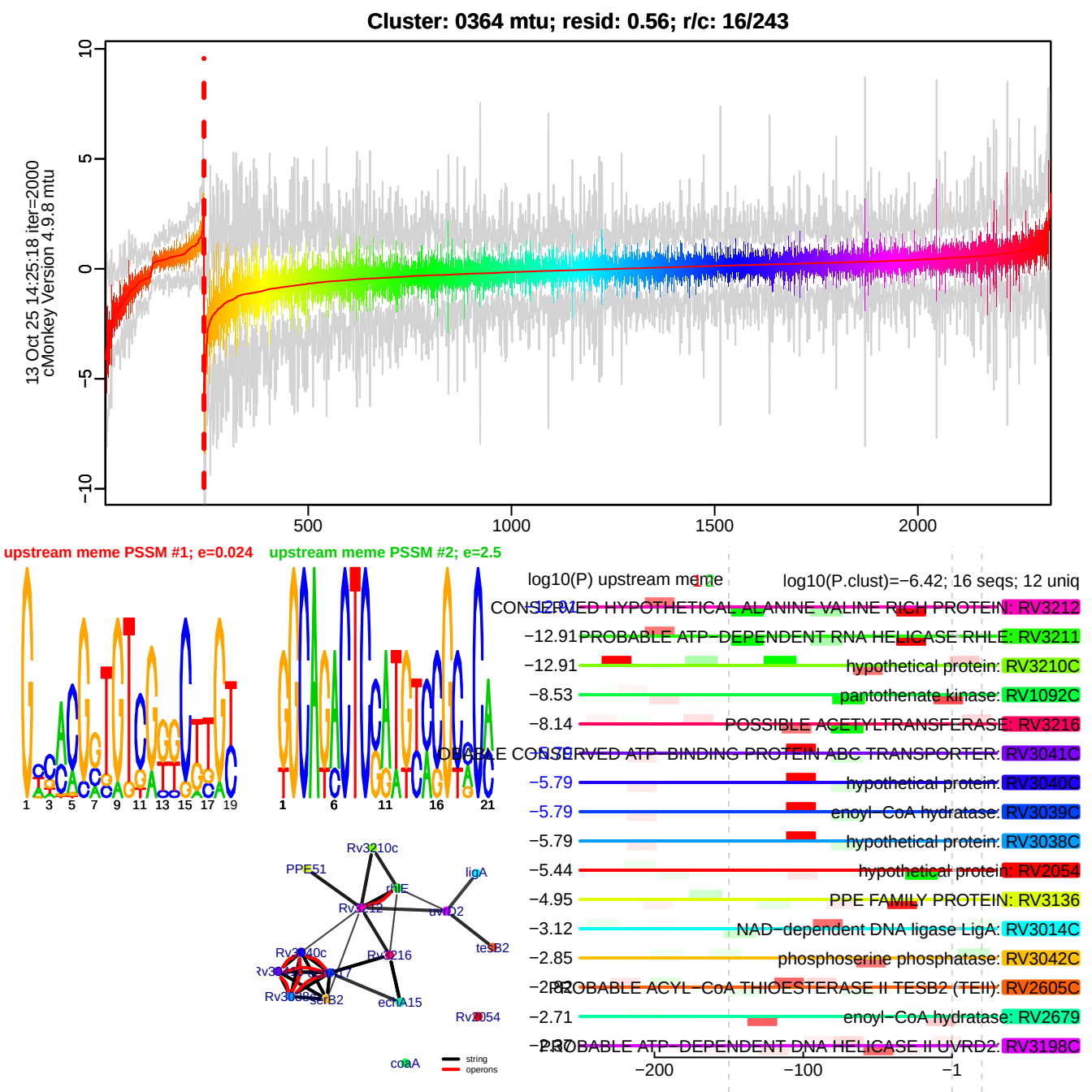

Cluster: 0364 mtu; resid: 0.56; r/c: 16/243
13 Oct 25 14:25:18 iter=2000
cMonkey Version 4.9.8 mtu
upstream meme PSSM #1; e=0.024
upstream meme PSSM #2; e=2.5
log10(P) upstream meme
log10(P.clust)=−6.42; 16 seqs; 12 uniq
CONSERVED HYPOTHETICAL ALANINE VALINE RICH PROTEIN: RV3212
PROBABLE ATP−DEPENDENT RNA HELICASE RHLE: RV3211
hypothetical protein: RV3210C
pantothenate kinase: RV1092C
POSSIBLE ACETYLTRANSFERASE: RV3216
PROBABLE CONSERVED ATP−BINDING PROTEIN ABC TRANSPORTER: RV3041C
hypothetical protein: RV3040C
enoyl−CoA hydratase: RV3039C
hypothetical protein: RV3038C
hypothetical protein: RV2054
PPE FAMILY PROTEIN: RV3136
NAD−dependent DNA ligase LigA: RV3014C
phosphoserine phosphatase: RV3042C
PROBABLE ACYL−CoA THIOESTERASE II TESB2 (TEII): RV2605C
enoyl−CoA hydratase: RV2679
PROBABLE ATP−DEPENDENT DNA HELICASE II UVRD2: RV3198C
Rv3210c
PPE51
ligA
rhlE
Rv3212
uvrD2
tesB2
Rv3040c
Rv3216
Rv3041c
cenA17
Rv3038c
serB2
echA15
Rv2054
coaA
string
operons