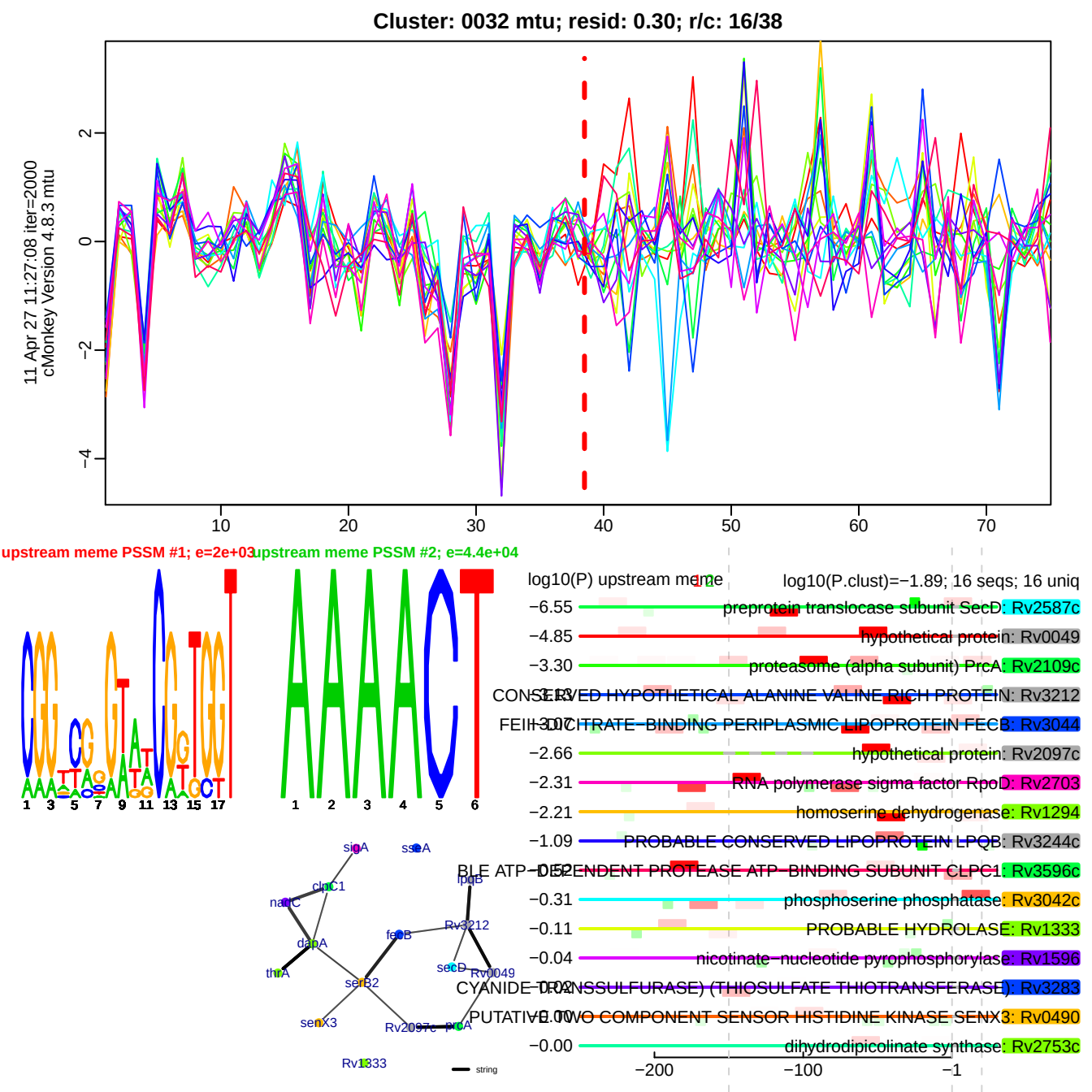
Cluster: 0032 mtu; resid: 0.30; r/c: 16/38
11 Apr 27 11:27:08 iter=2000
cMonkey Version 4.8.3 mtu
upstream meme PSSM #1; e=2e+03
upstream meme PSSM #2; e=4.4e+04
log10(P) upstream meme
log10(P.clust)=−1.89; 16 seqs; 16 uniq
−6.55 preprotein translocase subunit SecD: Rv2587c
−4.85 hypothetical protein: Rv0049
−3.30 proteasome (alpha subunit) PrcA: Rv2109c
CONSERVED HYPOTHETICAL ALANINE VALINE RICH PROTEIN: Rv3212
FEIH-3.07 CITRATE BINDING PERIPLASMIC LIPOPROTEIN FECB: Rv3044
−2.66 hypothetical protein: Rv2097c
−2.31 RNA polymerase sigma factor RpoD: Rv2703
−2.21 homoserine dehydrogenase: Rv1294
−1.09 PROBABLE CONSERVED LIPOPROTEIN LPQB: Rv3244c
BLE ATP DEPENDENT PROTEASE ATP BINDING SUBUNIT CLPC1: Rv3596c
−0.31 phosphoserine phosphatase: Rv3042c
−0.11 PROBABLE HYDROLASE: Rv1333
−0.04 nicotinate nucleotide pyrophosphorylase: Rv1596
CYANIDE DEHYDRATASE (THIOSULFATE THIOTRANSFERASE): Rv3283
PUTATIVE TWO COMPONENT SENSOR HISTIDINE KINASE SENX3: Rv0490
−0.00 dihydrodipicolinate synthase: Rv2753c
−200
−100
−1
sigA
sseA
clpC1
nadC
lpqB
Rv3212
feoB
dapA
thrA
secD
Rv0049
serB2
senX3
Rv2097c
prcA
Rv1333
string