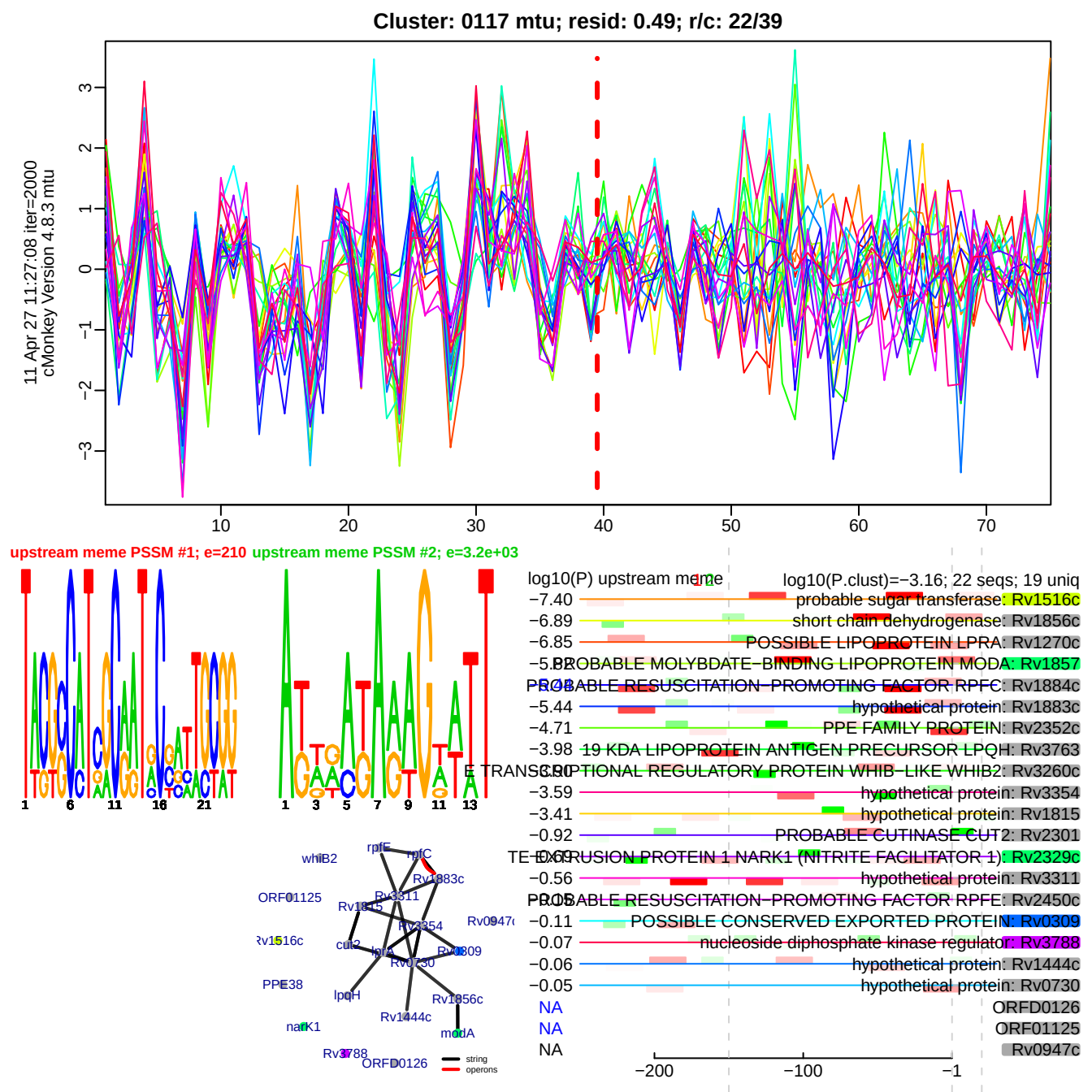
Cluster: 0117 mtu; resid: 0.49; r/c: 22/39
11 Apr 27 11:27:08 iter=2000
cMonkey Version 4.8.3 mtu
upstream meme PSSM #1; e=210
upstream meme PSSM #2; e=3.2e+03
log10(P) upstream meme
log10(P.clust)=−3.16; 22 seqs; 19 uniq
−7.40 probable sugar transferase: Rv1516c
−6.89 short chain dehydrogenase: Rv1856c
−6.85 POSSIBLE LIPOPROTEIN LPRA: Rv1270c
−5.82 PROBABLE MOLYBDATE−BINDING LIPOPROTEIN MODA: Rv1857
PROBABLE RESUSCITATION−PROMOTING FACTOR RPFC: Rv1884c
−5.44 hypothetical protein: Rv1883c
−4.71 PPE FAMILY PROTEIN: Rv2352c
−3.98 19 KDA LIPOPROTEIN ANTIGEN PRECURSOR LPQH: Rv3763
TRANSCRIPTIONAL REGULATORY PROTEIN WHIB−LIKE WHIB2: Rv3260c
−3.59 hypothetical protein: Rv3354
−3.41 hypothetical protein: Rv1815
−0.92 PROBABLE CUTINASE CUT2: Rv2301
TE−EXCLUSION PROTEIN 1 NARK1 (NITRITE FACILITATOR 1): Rv2329c
−0.56 hypothetical protein: Rv3311
PROBABLE RESUSCITATION−PROMOTING FACTOR RPFE: Rv2450c
−0.11 POSSIBLE CONSERVED EXPORTED PROTEIN: Rv0309
−0.07 nucleoside diphosphate kinase regulator: Rv3788
−0.06 hypothetical protein: Rv1444c
−0.05 hypothetical protein: Rv0730
NA ORFD0126
NA ORF01125
NA Rv0947c
−200 −100 −1
whiB2
rpfE
rpfC
Rv1883c
ORF01125
Rv3311
Rv1815
Rv3354
Rv0947c
Rv1516c
cut2
lpqA
Rv0309
Rv0730
PPE38
lpqH
Rv1856c
Rv1444c
narK1
modA
Rv3788
ORFD0126
string
operons