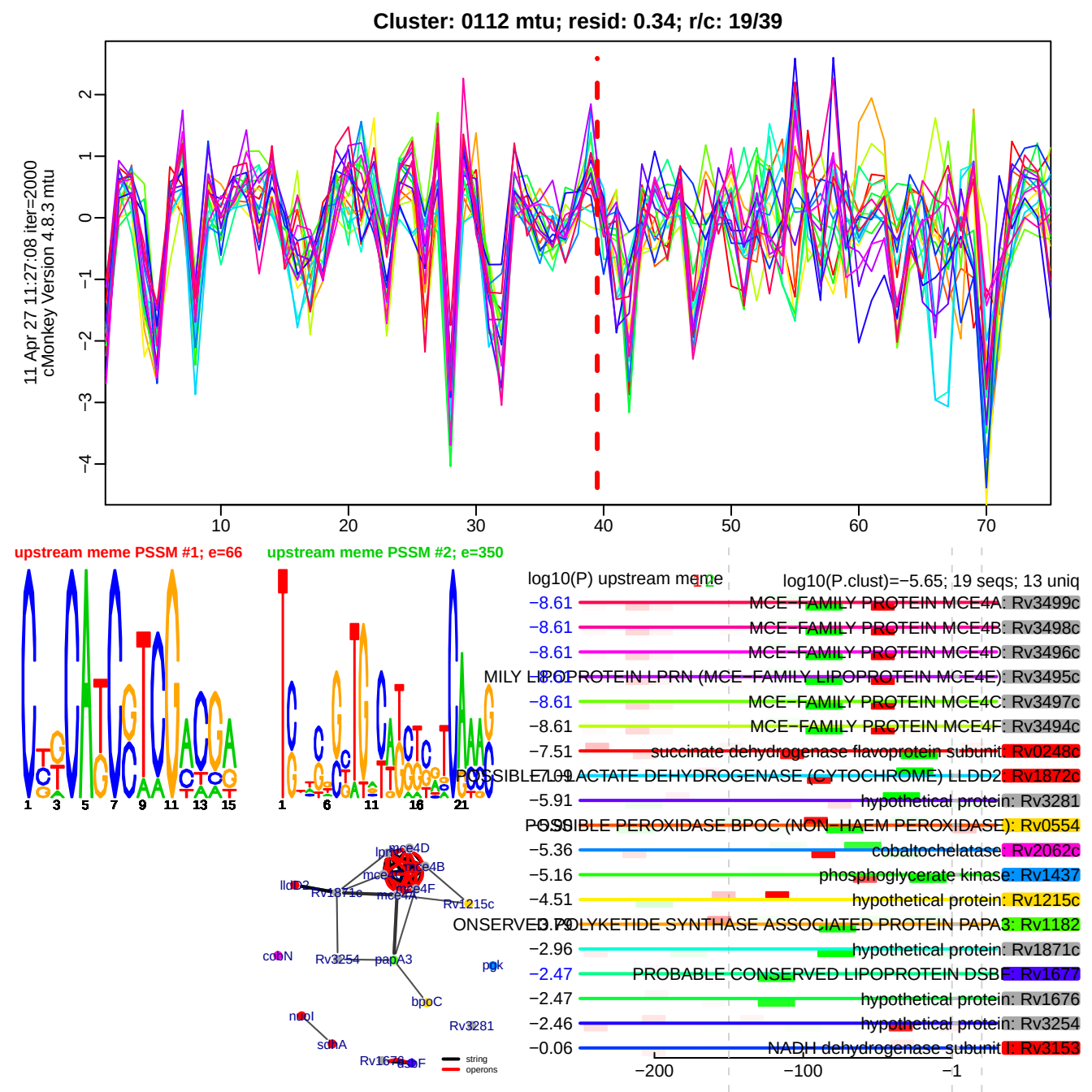

Cluster: 0112 mtu; resid: 0.34; r/c: 19/39
11 Apr 27 11:27:08 iter=2000
cMonkey Version 4.8.3 mtu
upstream meme PSSM #1; e=66
upstream meme PSSM #2; e=350
log10(P) upstream meme
log10(P.clust)=−5.65; 19 seqs; 13 uniq
−8.61 MCE−FAMILY PROTEIN MCE4A: Rv3499c
−8.61 MCE−FAMILY PROTEIN MCE4B: Rv3498c
−8.61 MCE−FAMILY PROTEIN MCE4D: Rv3496c
MILY LIPOPROTEIN LPRN (MCE−FAMILY LIPOPROTEIN MCE4E): Rv3495c
−8.61 MCE−FAMILY PROTEIN MCE4C: Rv3497c
−8.61 MCE−FAMILY PROTEIN MCE4F: Rv3494c
−7.51 succinate dehydrogenase flavoprotein subunit: Rv0248c
POSSIBLE LOLACTATE DEHYDROGENASE (CYTOCHROME) LLDD2: Rv1872c
−5.91 hypothetical protein: Rv3281
POSSIBLE PEROXIDASE BPOC (NON−HAEM PEROXIDASE): Rv0554
−5.36 cobaltochelatase: Rv2062c
−5.16 phosphoglycerate kinase: Rv1437
−4.51 hypothetical protein: Rv1215c
ONSERVED POLYKETIDE SYNTHASE ASSOCIATED PROTEIN PAPA3: Rv1182
−2.96 hypothetical protein: Rv1871c
−2.47 PROBABLE CONSERVED LIPOPROTEIN DSBF: Rv1677
−2.47 hypothetical protein: Rv1676
−2.46 hypothetical protein: Rv3254
−0.06 NADH dehydrogenase subunit I: Rv3153
string
operons