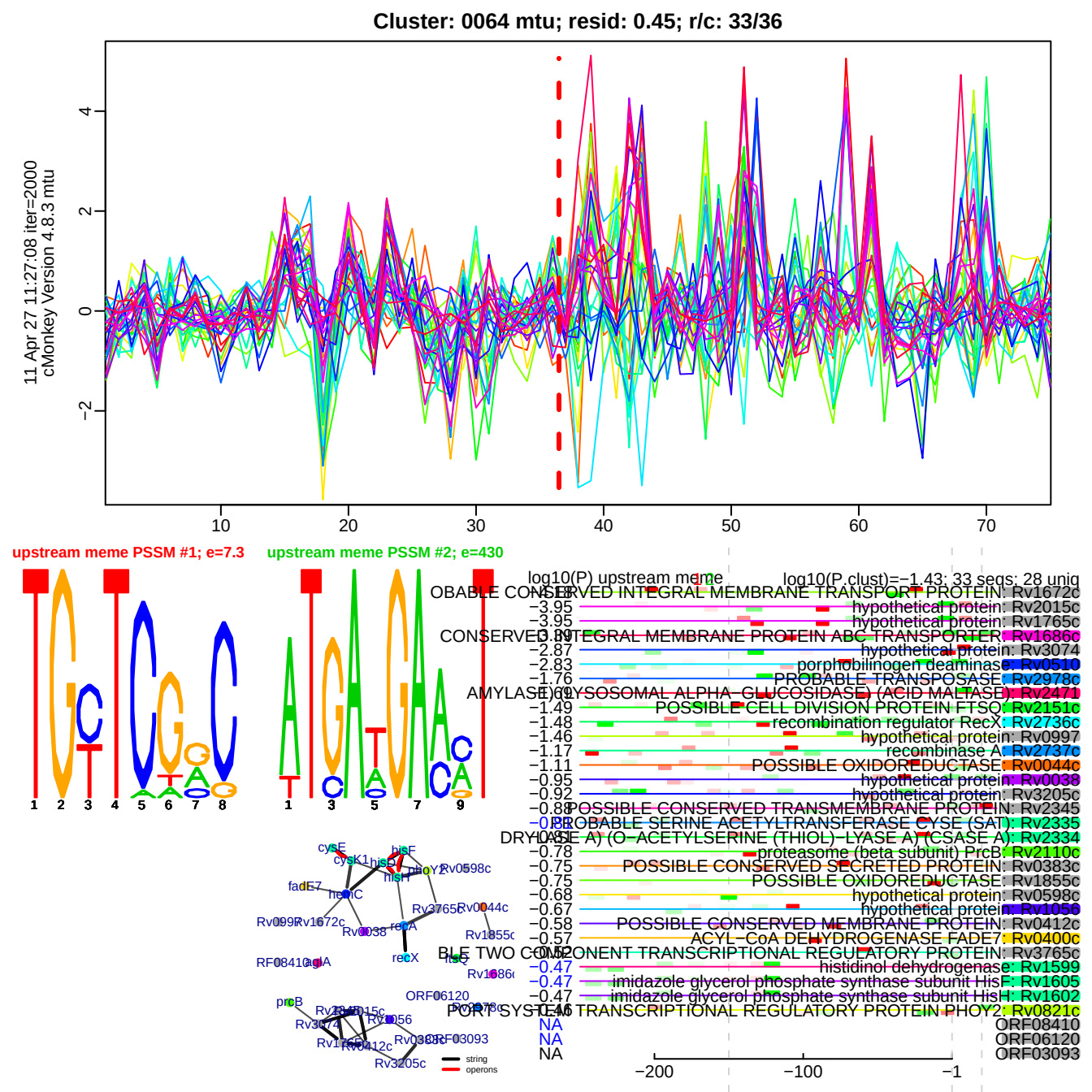
Cluster: 0064 mtu; resid: 0.45; r/c: 33/36
11 Apr 27 11:27:08 iter=2000
cMonkey Version 4.8.3 mtu
upstream meme PSSM #1; e=7.3
upstream meme PSSM #2; e=430
log10(P) upstream meme
log10(P.clust)=−1.43; 33 seqs; 28 uniq
−3.95 hypothetical protein: Rv2015c
−3.95 hypothetical protein: Rv1765c
−2.87 hypothetical protein: Rv3074
−2.83 porphobilinogen deaminase: Rv0510
−1.76 PROBABLE TRANSPOSASE: Rv2978c
−1.49 POSSIBLE CELL DIVISION PROTEIN FTSQ: Rv2151c
−1.48 recombination regulator RecX: Rv2736c
−1.46 hypothetical protein: Rv0997
−1.17 recombinase A: Rv2737c
−1.11 POSSIBLE OXIDOREDUCTASE: Rv0044c
−0.95 hypothetical protein: Rv0038
−0.92 hypothetical protein: Rv3205c
−0.78 proteasome (beta subunit) PrcB: Rv2110c
−0.75 POSSIBLE CONSERVED SECRETED PROTEIN: Rv0383c
−0.75 POSSIBLE OXIDOREDUCTASE: Rv1855c
−0.68 hypothetical protein: Rv0598c
−0.67 hypothetical protein: Rv1056
−0.58 POSSIBLE CONSERVED MEMBRANE PROTEIN: Rv0412c
−0.57 ACYL−CoA DEHYDROGENASE FADE7: Rv0400c
−0.52 TWO COMPONENT TRANSCRIPTIONAL REGULATORY PROTEIN: Rv3765c
−0.47 histidinol dehydrogenase: Rv1599
−0.47 imidazole glycerol phosphate synthase subunit HisF: Rv1605
−0.47 imidazole glycerol phosphate synthase subunit HisH: Rv1602
PORT SYSTEM TRANSCRIPTIONAL REGULATORY PROTEIN PHOY2: Rv0821c
NA ORF08410
NA ORF06120
NA ORF03093
−200 −100 −1
string
operons