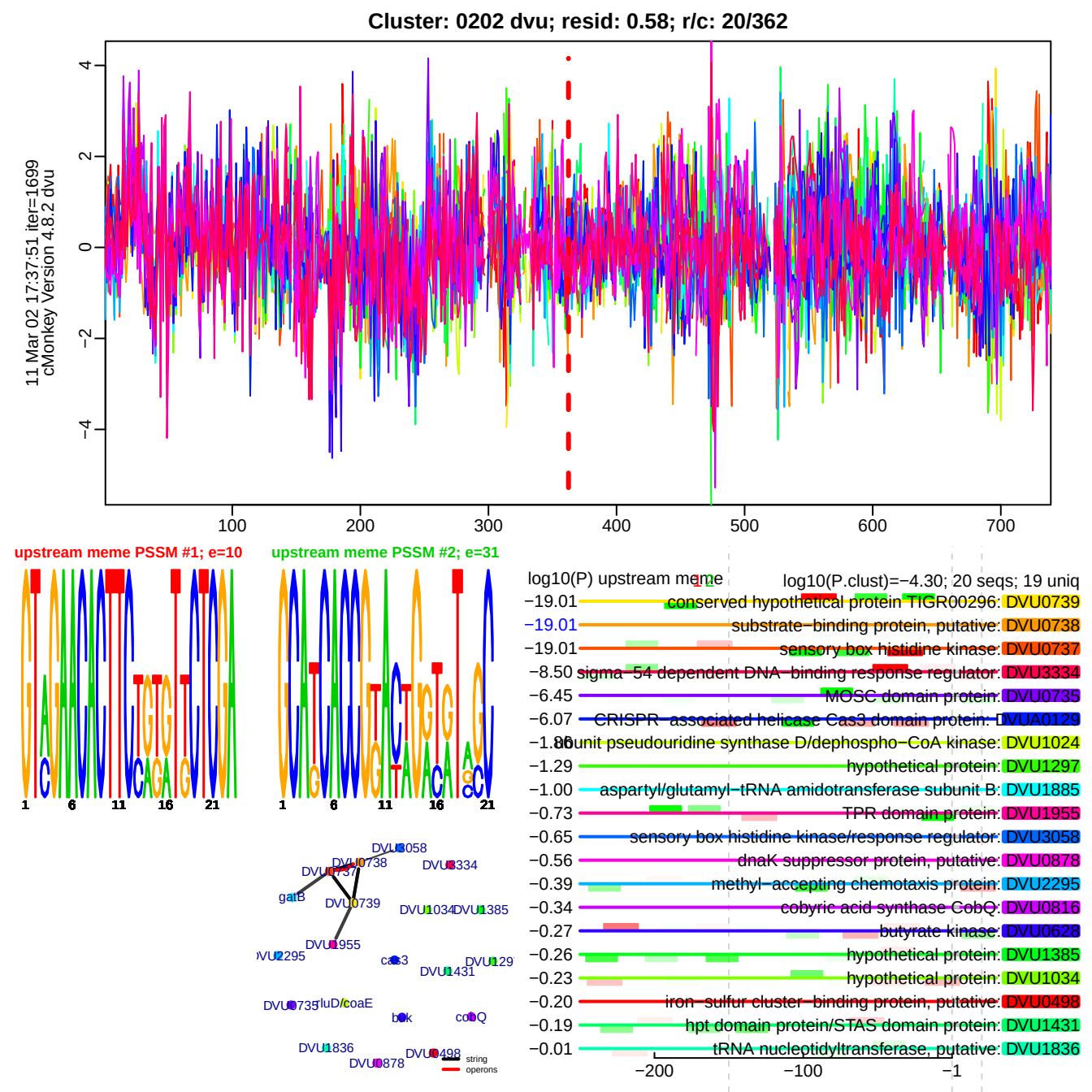
Cluster: 0202 dvu; resid: 0.58; r/c: 20/362
11 Mar 02 17:37:51 iter=1699
cMonkey Version 4.8.2 dvu
upstream meme PSSM #1; e=10
upstream meme PSSM #2; e=31
log10(P) upstream meme
log10(P.clust)=−4.30; 20 seqs; 19 uniq
−19.01 conserved hypothetical protein TIGR00296: DVU0739
−19.01 substrate−binding protein, putative: DVU0738
−19.01 sensory box histidine kinase: DVU0737
−8.50 sigma−54 dependent DNA−binding response regulator: DVU3334
−6.45 MOSC domain protein: DVU0735
−6.07 CRISPR−associated helicase Cas3 domain protein: DVU0129
−1.86 subunit pseudouridine synthase D/dephospho−CoA kinase: DVU1024
−1.29 hypothetical protein: DVU1297
−1.00 aspartyl/glutamyl−tRNA amidotransferase subunit B: DVU1885
−0.73 TPR domain protein: DVU1955
−0.65 sensory box histidine kinase/response regulator: DVU3058
−0.56 dnaK suppressor protein, putative: DVU0878
−0.39 methyl−accepting chemotaxis protein: DVU2295
−0.34 cobyric acid synthase CobQ: DVU0816
−0.27 butyrate kinase: DVU0628
−0.26 hypothetical protein: DVU1385
−0.23 hypothetical protein: DVU1034
−0.20 iron−sulfur cluster−binding protein, putative: DVU0498
−0.19 hpt domain protein/STAS domain protein: DVU1431
−0.01 tRNA nucleotidyltransferase, putative: DVU1836
string
operons