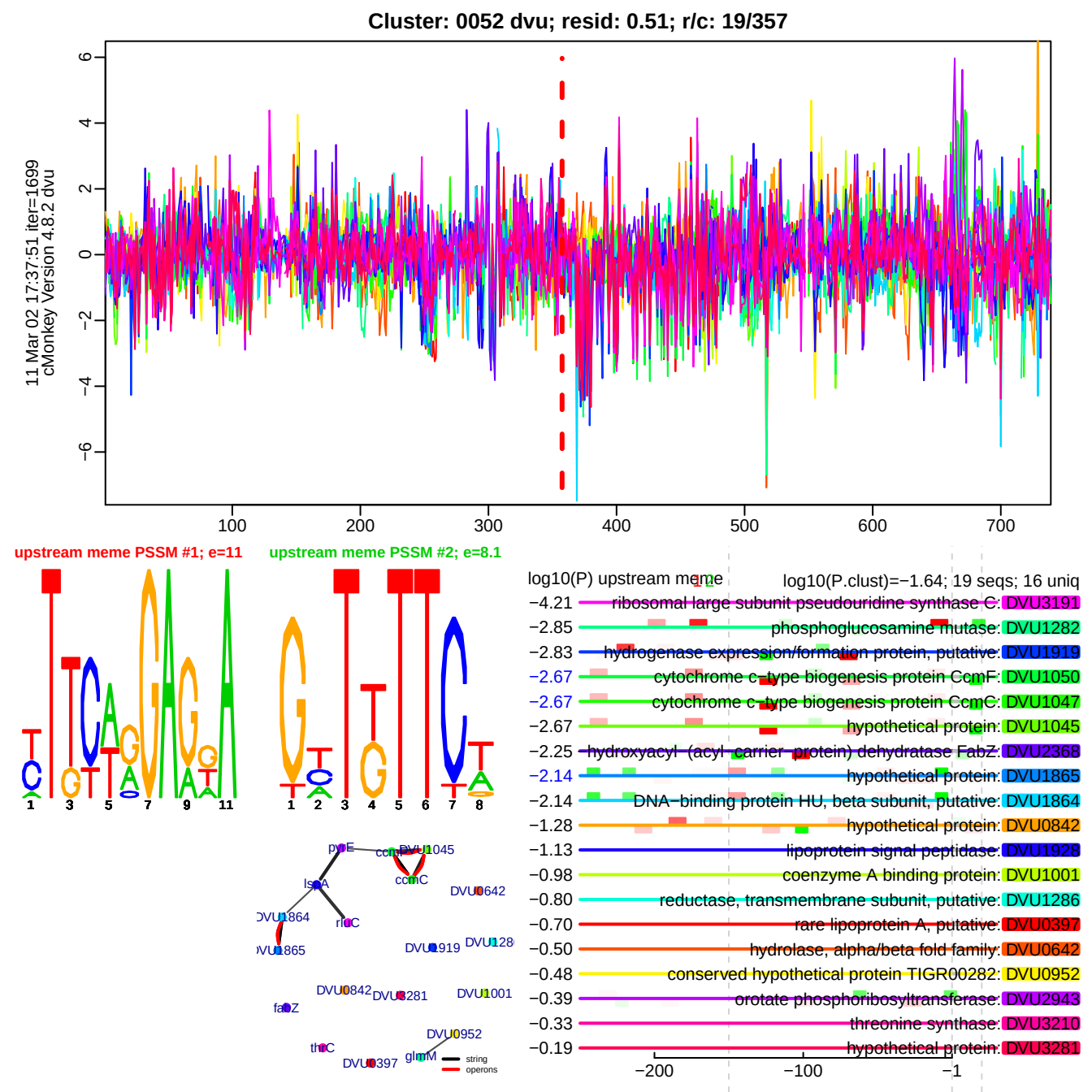
Cluster: 0052 dvu; resid: 0.51; r/c: 19/357
11 Mar 02 17:37:51 iter=1699
cMonkey Version 4.8.2 dvu
upstream meme PSSM #1; e=11
upstream meme PSSM #2; e=8.1
log10(P) upstream meme
log10(P.clust)=−1.64; 19 seqs; 16 uniq
−4.21 ribosomal large subunit pseudouridine synthase C: DVU3191
−2.85 phosphoglucosamine mutase: DVU1282
−2.83 hydrogenase expression/formation protein, putative: DVU1919
−2.67 cytochrome c-type biogenesis protein CcmF: DVU1050
−2.67 cytochrome c-type biogenesis protein CcmC: DVU1047
−2.67 hypothetical protein: DVU1045
−2.25 hydroxyacyl−(acyl−carrier−protein) dehydratase FabZ: DVU2368
−2.14 hypothetical protein: DVU1865
−2.14 DNA−binding protein HU, beta subunit, putative: DVU1864
−1.28 hypothetical protein: DVU0842
−1.13 lipoprotein signal peptidase: DVU1928
−0.98 coenzyme A binding protein: DVU1001
−0.80 reductase, transmembrane subunit, putative: DVU1286
−0.70 rare lipoprotein A, putative: DVU0397
−0.50 hydrolase, alpha/beta fold family: DVU0642
−0.48 conserved hypothetical protein TIGR00282: DVU0952
−0.39 orotate phosphoribosyltransferase: DVU2943
−0.33 threonine synthase: DVU3210
−0.19 hypothetical protein: DVU3281
string
operons