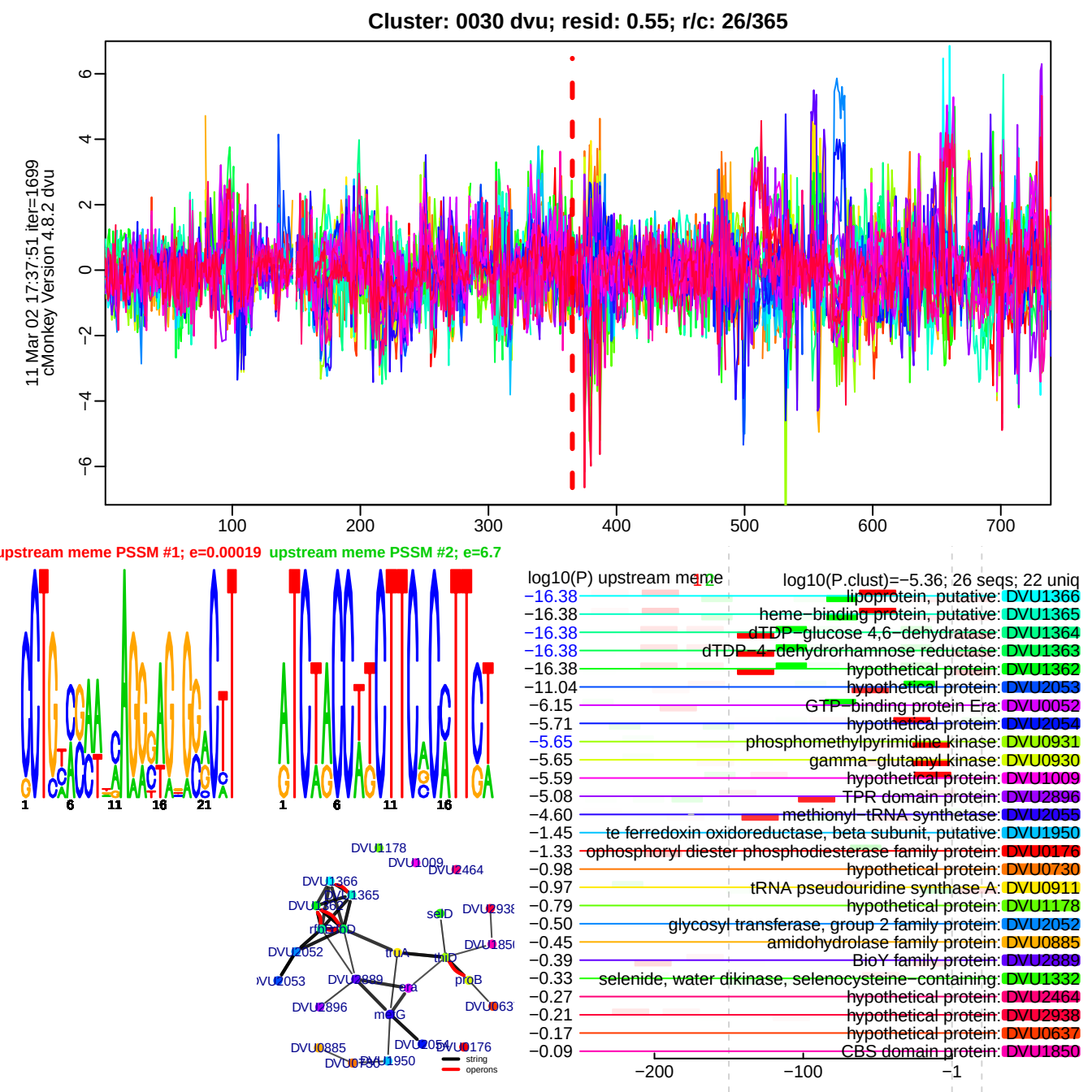
Cluster: 0030 dvu; resid: 0.55; r/c: 26/365
11 Mar 02 17:37:51 iter=1699
cMonkey Version 4.8.2 dvu
upstream meme PSSM #1; e=0.00019
upstream meme PSSM #2; e=6.7
log10(P) upstream meme
log10(P.clust)=−5.36; 26 seqs; 22 uniq
−16.38 lipoprotein, putative: DVU1366
−16.38 heme−binding protein, putative: DVU1365
−16.38 dTDP−glucose 4,6−dehydratase: DVU1364
−16.38 dTDP−4−dehydrorhamnose reductase: DVU1363
−16.38 hypothetical protein: DVU1362
−11.04 hypothetical protein: DVU2053
−6.15 GTP−binding protein Era: DVU0052
−5.71 hypothetical protein: DVU2054
−5.65 phosphomethylpyrimidine kinase: DVU0931
−5.65 gamma−glutamyl kinase: DVU0930
−5.59 hypothetical protein: DVU1009
−5.08 TPR domain protein: DVU2896
−4.60 methionyl−tRNA synthetase: DVU2055
−1.45 te ferredoxin oxidoreductase, beta subunit, putative: DVU1950
−1.33 ophosphoryl diester phosphodiesterase family protein: DVU0176
−0.98 hypothetical protein: DVU0730
−0.97 tRNA pseudouridine synthase A: DVU0911
−0.79 hypothetical protein: DVU1178
−0.50 glycosyl transferase, group 2 family protein: DVU2052
−0.45 amidohydrolase family protein: DVU0885
−0.39 BioY family protein: DVU2889
−0.33 selenide, water dikinase, selenocysteine−containing: DVU1332
−0.27 hypothetical protein: DVU2464
−0.21 hypothetical protein: DVU2938
−0.17 hypothetical protein: DVU0637
−0.09 CBS domain protein: DVU1850
DVU1178
DVU1009
DVU2464
DVU1366
DVU1365
selD
DVU2938
DVU2052
truA
thiD
DVU1850
DVU2053
DVU2889
era
proB
DVU2896
DVU0637
mnmG
DVU0885
DVU2054
DVU0176
DVU1950
string
operons