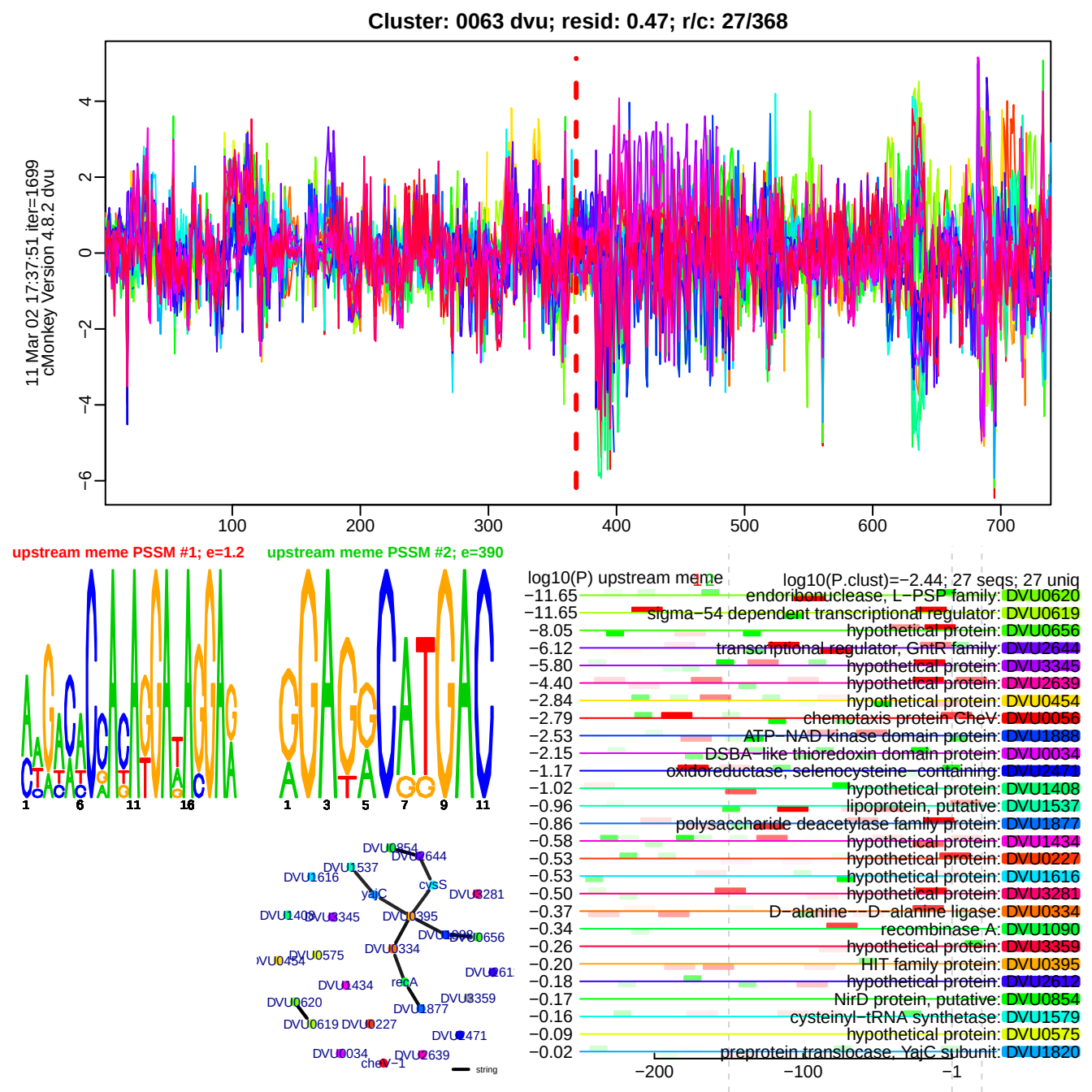
Cluster: 0063 dvu; resid: 0.47; r/c: 27/368
11 Mar 02 17:37:51 iter=1699
cMonkey Version 4.8.2 dvu
upstream meme PSSM #1; e=1.2
upstream meme PSSM #2; e=390
log10(P) upstream meme
log10(P.clust)=−2.44; 27 seqs; 27 uniq
−11.65 endoribonuclease, L−PSP family: DVU0620
−11.65 sigma−54 dependent transcriptional regulator: DVU0619
−8.05 hypothetical protein: DVU0656
−6.12 transcriptional regulator, GntR family: DVU2644
−5.80 hypothetical protein: DVU3345
−4.40 hypothetical protein: DVU2639
−2.84 hypothetical protein: DVU0454
−2.79 chemotaxis protein CheV: DVU0056
−2.53 ATP−NAD kinase domain protein: DVU1888
−2.15 DSBA−like thioredoxin domain protein: DVU0034
−1.17 oxidoreductase, selenocysteine−containing: DVU2471
−1.02 hypothetical protein: DVU1408
−0.96 lipoprotein, putative: DVU1537
−0.86 polysaccharide deacetylase family protein: DVU1877
−0.58 hypothetical protein: DVU1434
−0.53 hypothetical protein: DVU0227
−0.53 hypothetical protein: DVU1616
−0.50 hypothetical protein: DVU3281
−0.37 D−alanine−−D−alanine ligase: DVU0334
−0.34 recombinase A: DVU1090
−0.26 hypothetical protein: DVU3359
−0.20 HIT family protein: DVU0395
−0.18 hypothetical protein: DVU2612
−0.17 NirD protein, putative: DVU0854
−0.16 cysteinyl−tRNA synthetase: DVU1579
−0.09 hypothetical protein: DVU0575
−0.02 preprotein translocase, YajC subunit: DVU1820
DVU0854
DVU2644
DVU1537
DVU1616
cysS
yajC
DVU3281
DVU1408
DVU3345
DVU0395
DVU0656
DVU0334
DVU0575
DVU1434
recA
DVU3359
DVU0620
DVU1877
DVU0619
DVU0227
DVU2471
DVU0034
DVU2639
cheV−1
string