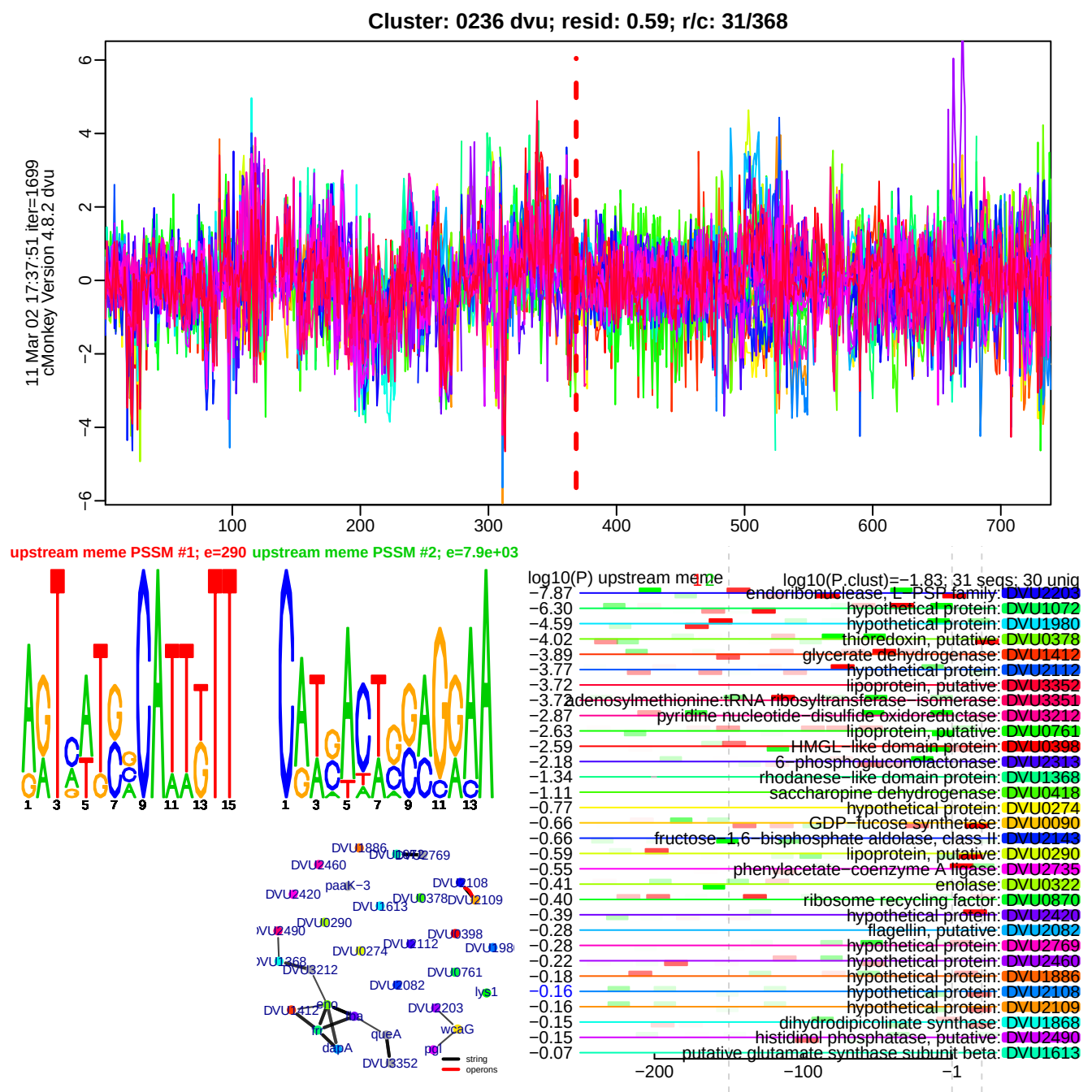
Cluster: 0236 dvu; resid: 0.59; r/c: 31/368
11 Mar 02 17:37:51 iter=1699
cMonkey Version 4.8.2 dvu
upstream meme PSSM #1; e=290
upstream meme PSSM #2; e=7.9e+03
log10(P) upstream meme
log10(P.clust)=−1.83; 31 seqs; 30 uniq
−7.87 endoribonuclease, L−PSP family: DVU2203
−6.30 hypothetical protein: DVU1072
−4.59 hypothetical protein: DVU1980
−4.02 thioredoxin, putative: DVU0378
−3.89 glycerate dehydrogenase: DVU1412
−3.77 hypothetical protein: DVU2112
−3.72 lipoprotein, putative: DVU3352
−3.72 adenosylmethionine:tRNA ribosyltransferase−isomerase: DVU3351
−2.87 pyridine nucleotide−disulfide oxidoreductase: DVU3212
−2.63 lipoprotein, putative: DVU0761
−2.59 HMGL−like domain protein: DVU0398
−2.18 6−phosphogluconolactonase: DVU2313
−1.34 rhodanese−like domain protein: DVU1368
−1.11 saccharopine dehydrogenase: DVU0418
−0.77 hypothetical protein: DVU0274
−0.66 GDP−fucose synthetase: DVU0090
−0.66 fructose−1,6−bisphosphate aldolase, class II: DVU2143
−0.59 lipoprotein, putative: DVU0290
−0.55 phenylacetate−coenzyme A ligase: DVU2735
−0.41 enolase: DVU0322
−0.40 ribosome recycling factor: DVU0870
−0.39 hypothetical protein: DVU2420
−0.28 flagellin, putative: DVU2082
−0.28 hypothetical protein: DVU2769
−0.22 hypothetical protein: DVU2460
−0.18 hypothetical protein: DVU1886
−0.16 hypothetical protein: DVU2108
−0.16 hypothetical protein: DVU2109
−0.15 dihydrodipicolinate synthase: DVU1868
−0.15 histidinol phosphatase, putative: DVU2490
−0.07 putative glutamate synthase subunit beta: DVU1613
−200
−100
−1
DVU1886
DVU2769
DVU2460
paaK−3
DVU2108
DVU2420
DVU0378
DVU2109
DVU1613
DVU0290
DVU2490
DVU0398
DVU2112
DVU0274
DVU1368
DVU3212
DVU0761
DVU2082
lysS
DVU1412
DVU2203
wcaG
queA
dapA
DVU3352
string
operons