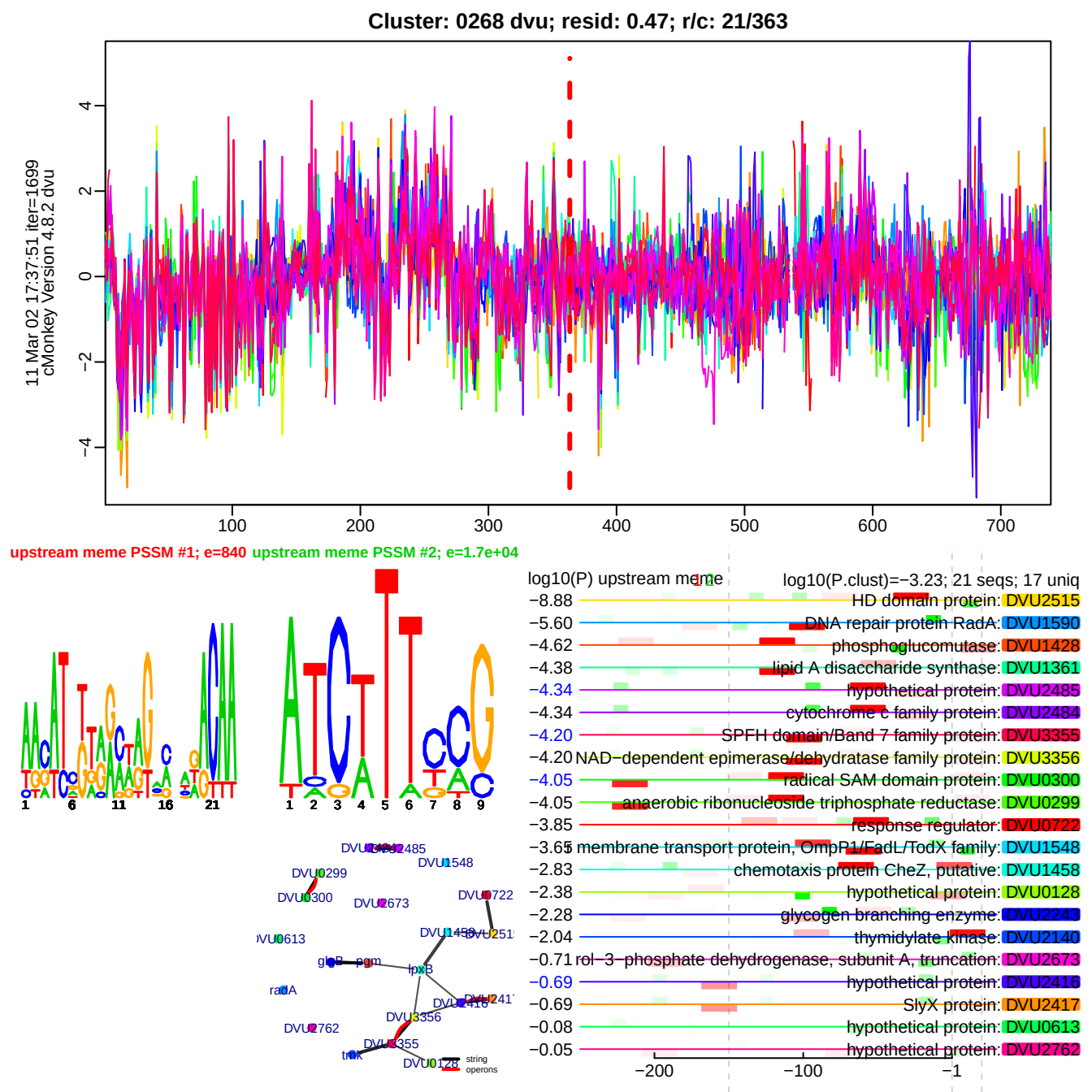
Cluster: 0268 dvu; resid: 0.47; r/c: 21/363
11 Mar 02 17:37:51 iter=1699
cMonkey Version 4.8.2 dvu
upstream meme PSSM #1; e=840
upstream meme PSSM #2; e=1.7e+04
log10(P) upstream meme
log10(P.clust)=−3.23; 21 seqs; 17 uniq
−8.88 HD domain protein: DVU2515
−5.60 DNA repair protein RadA: DVU1590
−4.62 phosphoglucomutase: DVU1428
−4.38 lipid A disaccharide synthase: DVU1361
−4.34 hypothetical protein: DVU2485
−4.34 cytochrome c family protein: DVU2484
−4.20 SPFH domain/Band 7 family protein: DVU3355
−4.20 NAD−dependent epimerase/dehydratase family protein: DVU3356
−4.05 radical SAM domain protein: DVU0300
−4.05 anaerobic ribonucleoside triphosphate reductase: DVU0299
−3.85 response regulator: DVU0722
−3.65 membrane transport protein, OmpP1/FadL/TodX family: DVU1548
−2.83 chemotaxis protein CheZ, putative: DVU1458
−2.38 hypothetical protein: DVU0128
−2.28 glycogen branching enzyme: DVU2243
−2.04 thymidylate kinase: DVU2140
−0.71 rol−3−phosphate dehydrogenase, subunit A, truncation: DVU2673
−0.69 hypothetical protein: DVU2416
−0.69 SlyX protein: DVU2417
−0.08 hypothetical protein: DVU0613
−0.05 hypothetical protein: DVU2762
DVU2485
DVU1548
DVU0299
DVU0300
DVU2673
DVU0722
DVU0613
DVU1458
DVU2510
glgB
pgm
lpxB
radA
DVU2416
DVU2417
DVU3356
DVU2762
DVU3355
tmk
DVU0128
string
operons