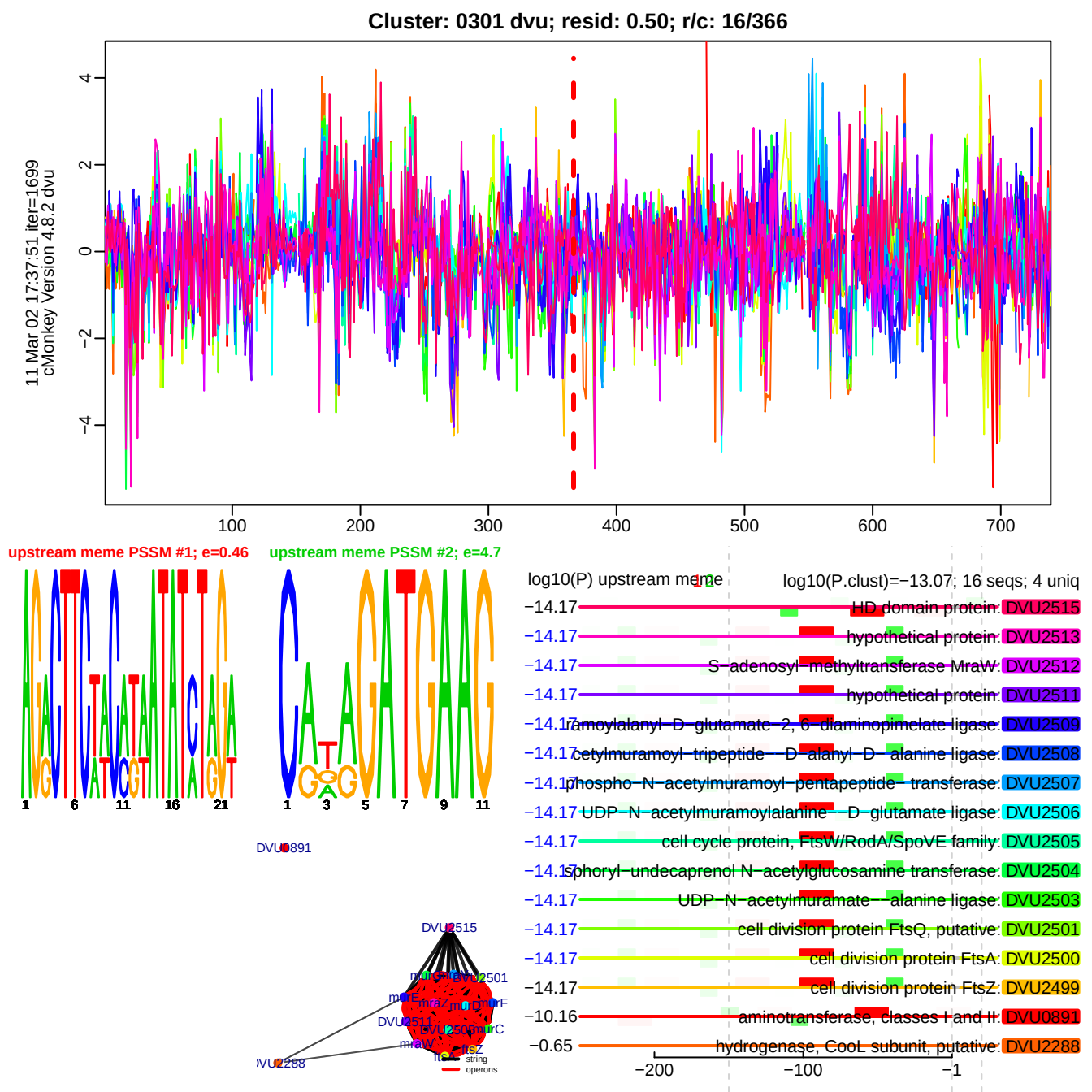
Cluster: 0301 dvu; resid: 0.50; r/c: 16/366
11 Mar 02 17:37:51 iter=1699
cMonkey Version 4.8.2 dvu
upstream meme PSSM #1; e=0.46
upstream meme PSSM #2; e=4.7
log10(P) upstream meme
log10(P.clust)=−13.07; 16 seqs; 4 uniq
−14.17 HD domain protein: DVU2515
−14.17 hypothetical protein: DVU2513
−14.17 S−adenosyl−methyltransferase MraW: DVU2512
−14.17 hypothetical protein: DVU2511
−14.17 amoylalanyl−D−glutamate−2, 6−diaminopimelate ligase: DVU2509
−14.17 cetylmuramoyl−tripeptide−−D−alanyl−D−alanine ligase: DVU2508
−14.17 phospho−N−acetylmuramoyl−pentapeptide−transferase: DVU2507
−14.17 UDP−N−acetylmuramoylalanine−−D−glutamate ligase: DVU2506
−14.17 cell cycle protein, FtsW/RodA/SpoVE family: DVU2505
−14.17 phoryl−undecaprenol N−acetylglucosamine transferase: DVU2504
−14.17 UDP−N−acetylmuramate−−alanine ligase: DVU2503
−14.17 cell division protein FtsQ, putative: DVU2501
−14.17 cell division protein FtsA: DVU2500
−14.17 cell division protein FtsZ: DVU2499
−10.16 aminotransferase, classes I and II: DVU0891
−0.65 hydrogenase, CooL subunit, putative: DVU2288
DVU0891
DVU2515
DVU2501
DVU2511
DVU2508
DVU2288
string
operons