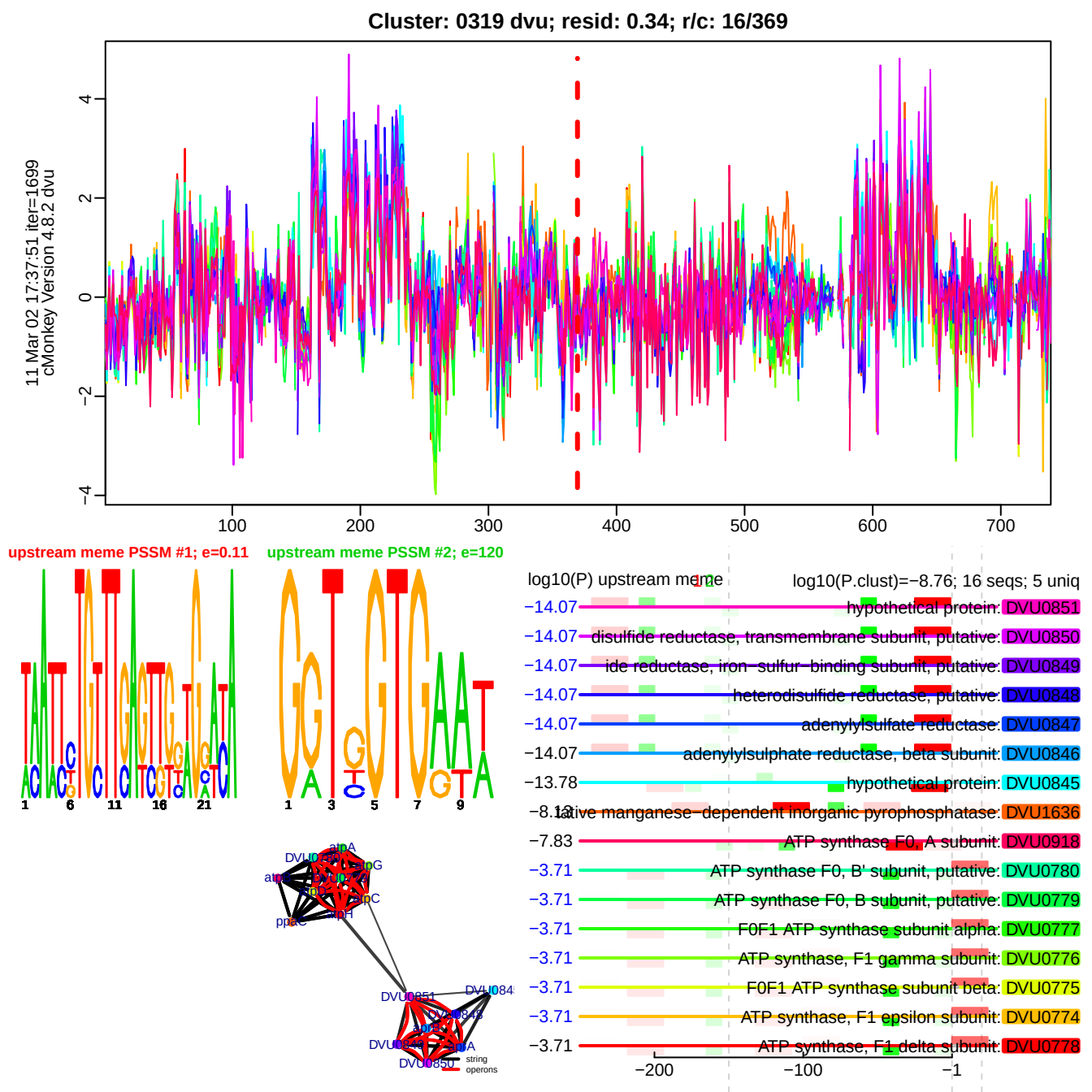
Cluster: 0319 dvu; resid: 0.34; r/c: 16/369
11 Mar 02 17:37:51 iter=1699
cMonkey Version 4.8.2 dvu
upstream meme PSSM #1; e=0.11
upstream meme PSSM #2; e=120
log10(P) upstream meme
log10(P.clust)=−8.76; 16 seqs; 5 uniq
−14.07 hypothetical protein: DVU0851
−14.07 disulfide reductase, transmembrane subunit, putative: DVU0850
−14.07 ide reductase, iron−sulfur−binding subunit, putative: DVU0849
−14.07 heterodisulfide reductase, putative: DVU0848
−14.07 adenylylsulfate reductase: DVU0847
−14.07 adenylylsulphate reductase, beta subunit: DVU0846
−13.78 hypothetical protein: DVU0845
−8.13 ative manganese−dependent inorganic pyrophosphatase: DVU1636
−7.83 ATP synthase F0, A subunit: DVU0918
−3.71 ATP synthase F0, B' subunit, putative: DVU0780
−3.71 ATP synthase F0, B subunit, putative: DVU0779
−3.71 F0F1 ATP synthase subunit alpha: DVU0777
−3.71 ATP synthase, F1 gamma subunit: DVU0776
−3.71 F0F1 ATP synthase subunit beta: DVU0775
−3.71 ATP synthase, F1 epsilon subunit: DVU0774
−3.71 ATP synthase, F1 delta subunit: DVU0778
string
operons