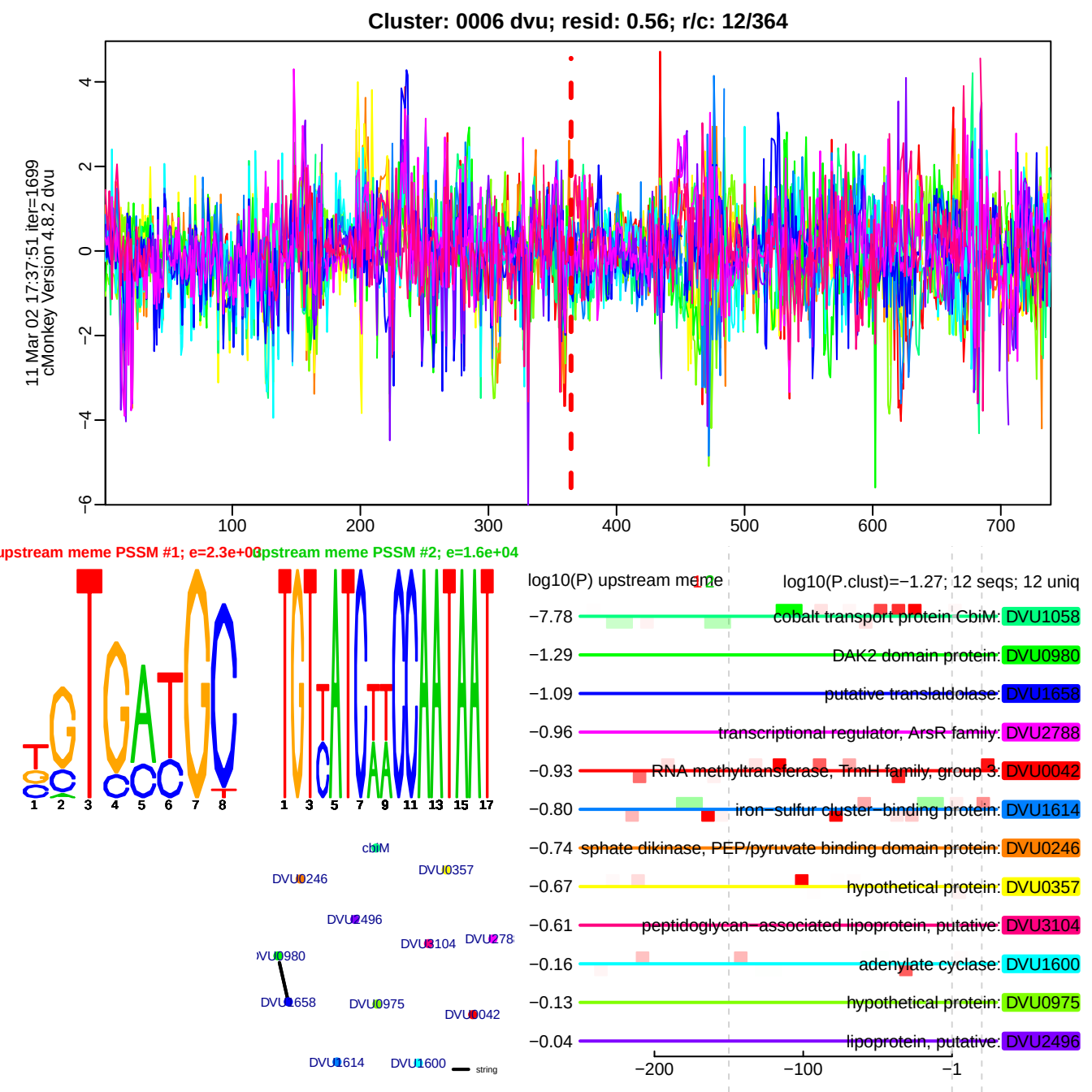
Cluster: 0006 dvu; resid: 0.56; r/c: 12/364
11 Mar 02 17:37:51 iter=1699
cMonkey Version 4.8.2 dvu
upstream meme PSSM #1; e=2.3e+03
upstream meme PSSM #2; e=1.6e+04
log10(P) upstream meme
log10(P.clust)=−1.27; 12 seqs; 12 uniq
−7.78 cobalt transport protein CbiM: DVU1058
−1.29 DAK2 domain protein: DVU0980
−1.09 putative translaldolase: DVU1658
−0.96 transcriptional regulator, ArsR family: DVU2788
−0.93 RNA methyltransferase, TrmH family, group 3: DVU0042
−0.80 iron−sulfur cluster−binding protein: DVU1614
−0.74 sphate dikinase, PEP/pyruvate binding domain protein: DVU0246
−0.67 hypothetical protein: DVU0357
−0.61 peptidoglycan−associated lipoprotein, putative: DVU3104
−0.16 adenylate cyclase: DVU1600
−0.13 hypothetical protein: DVU0975
−0.04 lipoprotein, putative: DVU2496
string