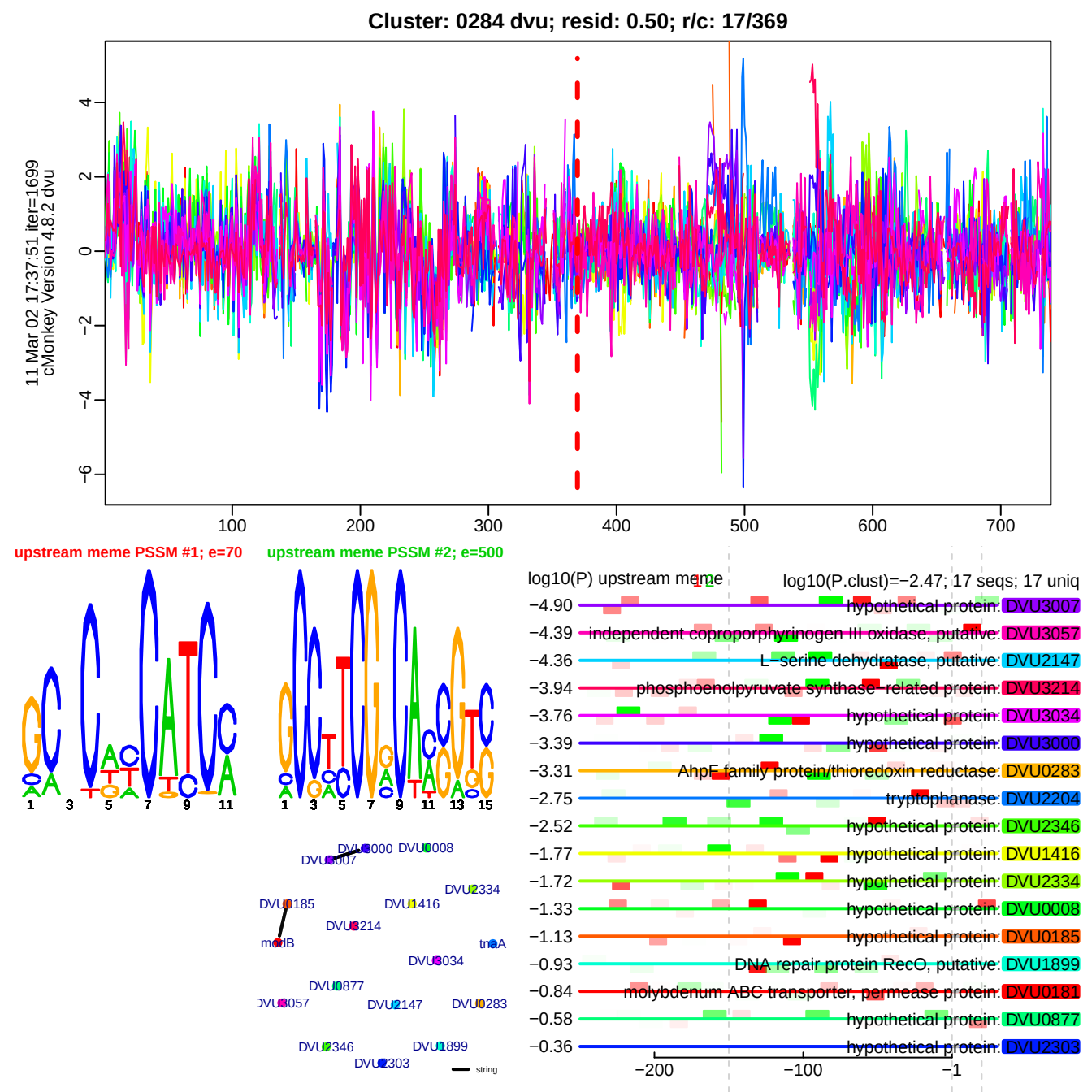
Cluster: 0284 dvu; resid: 0.50; r/c: 17/369
11 Mar 02 17:37:51 iter=1699
cMonkey Version 4.8.2 dvu
upstream meme PSSM #1; e=70
upstream meme PSSM #2; e=500
log10(P) upstream meme
log10(P.clust)=−2.47; 17 seqs; 17 uniq
−4.90 hypothetical protein: DVU3007
−4.39 independent coproporphyrinogen III oxidase, putative: DVU3057
−4.36 L−serine dehydratase, putative: DVU2147
−3.94 phosphoenolpyruvate synthase−related protein: DVU3214
−3.76 hypothetical protein: DVU3034
−3.39 hypothetical protein: DVU3000
−3.31 AhpF family protein/thioredoxin reductase: DVU0283
−2.75 tryptophanase: DVU2204
−2.52 hypothetical protein: DVU2346
−1.77 hypothetical protein: DVU1416
−1.72 hypothetical protein: DVU2334
−1.33 hypothetical protein: DVU0008
−1.13 hypothetical protein: DVU0185
−0.93 DNA repair protein RecO, putative: DVU1899
−0.84 molybdenum ABC transporter, permease protein: DVU0181
−0.58 hypothetical protein: DVU0877
−0.36 hypothetical protein: DVU2303
DVU3000
DVU0008
DVU3007
DVU2334
DVU0185
DVU1416
DVU3214
modB
tnaA
DVU3034
DVU0877
DVU3057
DVU2147
DVU0283
DVU2346
DVU1899
DVU2303
string