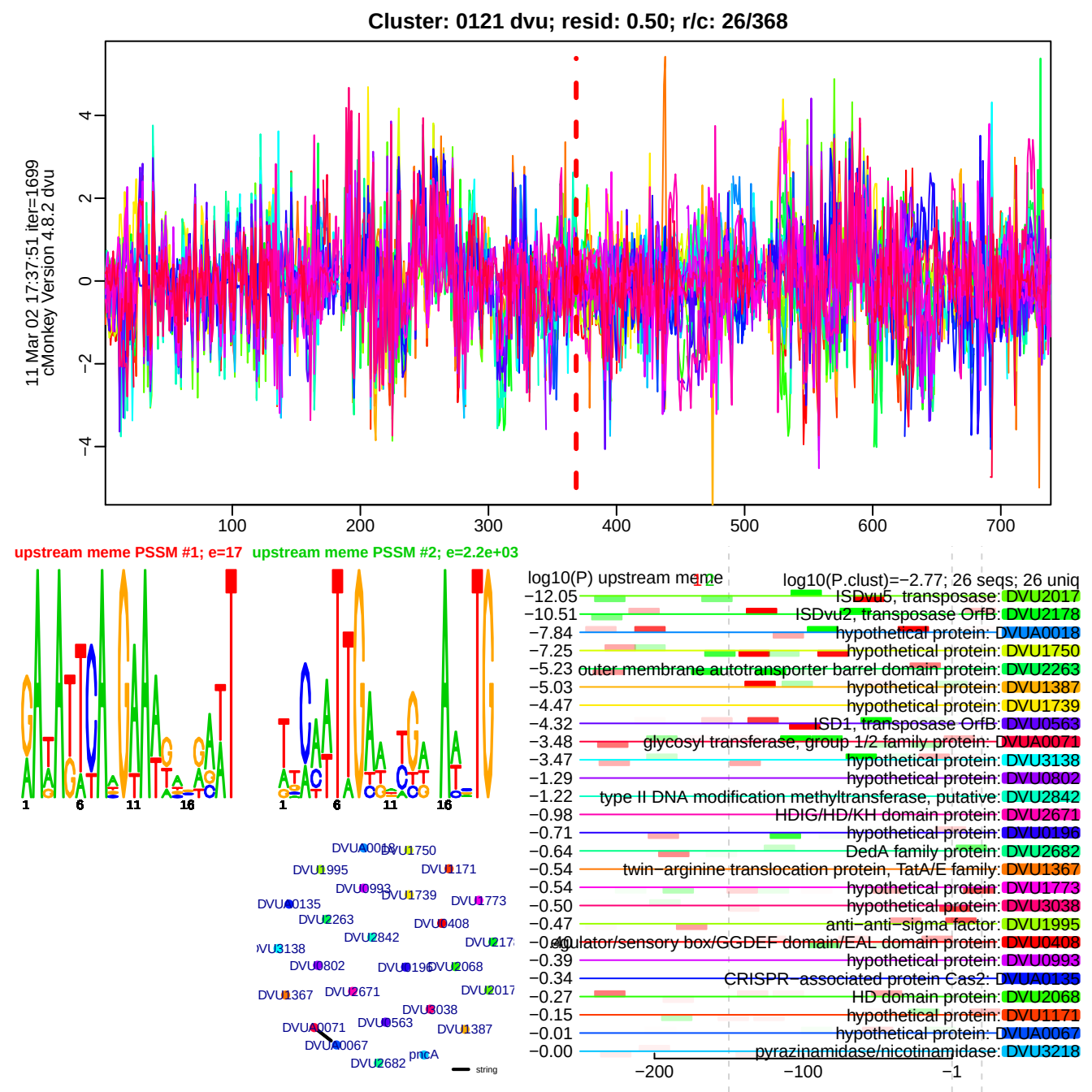
Cluster: 0121 dvu; resid: 0.50; r/c: 26/368
11 Mar 02 17:37:51 iter=1699
cMonkey Version 4.8.2 dvu
upstream meme PSSM #1; e=17
upstream meme PSSM #2; e=2.2e+03
log10(P) upstream meme
log10(P.clust)=−2.77; 26 seqs; 26 uniq
−12.05 ISDvu5, transposase: DVU2017
−10.51 ISDvu2, transposase OrfB: DVU2178
−7.84 hypothetical protein: DVUA0018
−7.25 hypothetical protein: DVU1750
−5.23 outer membrane autotransporter barrel domain protein: DVU2263
−5.03 hypothetical protein: DVU1387
−4.47 hypothetical protein: DVU1739
−4.32 ISD1, transposase OrfB: DVU0563
−3.48 glycosyl transferase, group 1/2 family protein: DVUA0071
−3.47 hypothetical protein: DVU3138
−1.29 hypothetical protein: DVU0802
−1.22 type II DNA modification methyltransferase, putative: DVU2842
−0.98 HDIG/HD/KH domain protein: DVU2671
−0.71 hypothetical protein: DVU0196
−0.64 DedA family protein: DVU2682
−0.54 twin−arginine translocation protein, TatA/E family: DVU1367
−0.54 hypothetical protein: DVU1773
−0.50 hypothetical protein: DVU3038
−0.47 anti−anti−sigma factor: DVU1995
−0.40 regulator/sensory box/GGDEF domain/EAL domain protein: DVU0408
−0.39 hypothetical protein: DVU0993
−0.34 CRISPR−associated protein Cas2: DVUA0135
−0.27 HD domain protein: DVU2068
−0.15 hypothetical protein: DVU1171
−0.01 hypothetical protein: DVUA0067
−0.00 pyrazinamidase/nicotinamidase: DVU3218
DVUA0018
DVU1750
DVU1995
DVU1171
DVU0993
DVU1739
DVU1773
DVUA0135
DVU2263
DVU0408
DVU2842
DVU3138
DVU0802
DVU0196
DVU2068
DVU1367
DVU2671
DVU2017
DVU3038
DVUA0071
DVU0563
DVU1387
DVUA0067
DVU2682
pncA
string