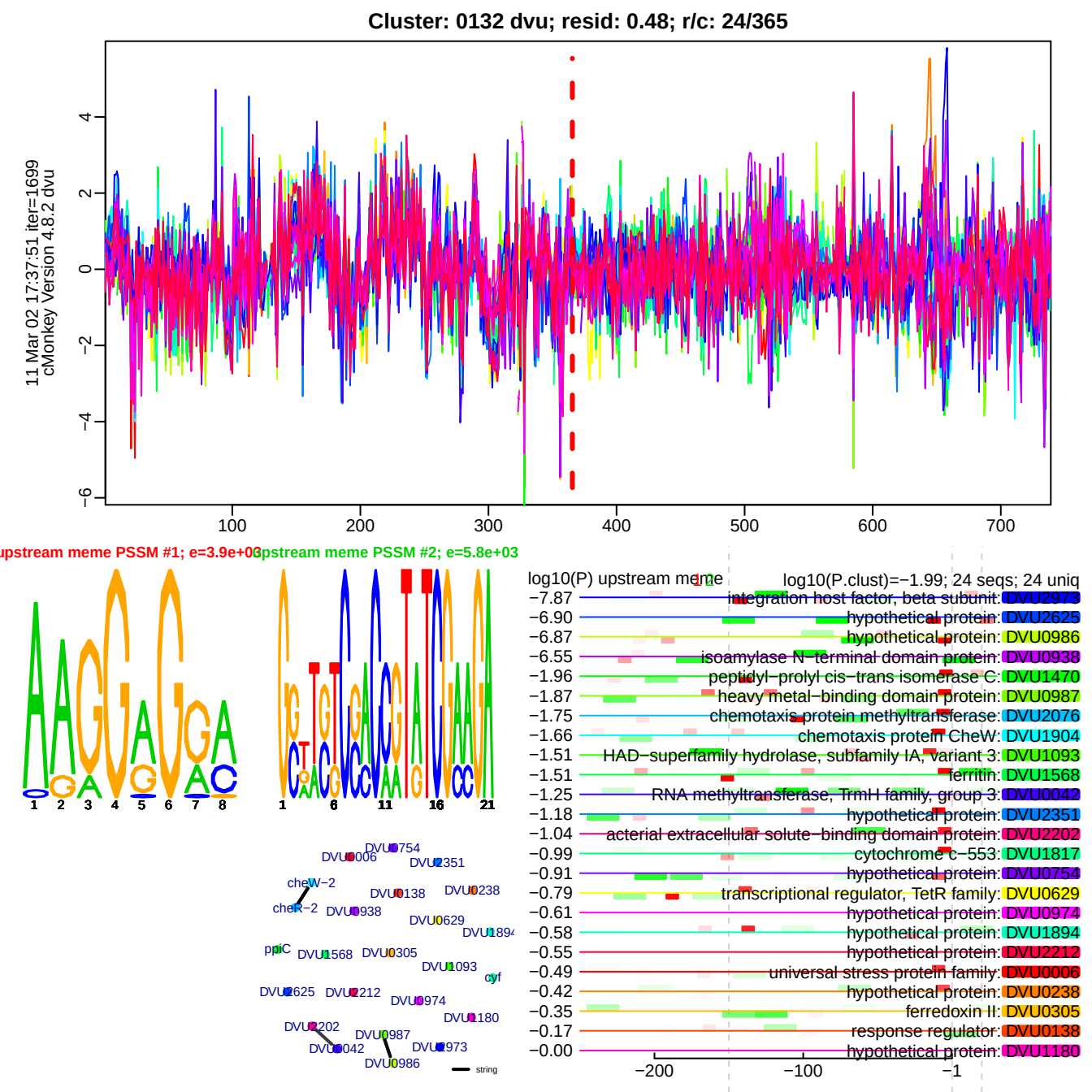
Cluster: 0132 dvu; resid: 0.48; r/c: 24/365
11 Mar 02 17:37:51 iter=1699
cMonkey Version 4.8.2 dvu
upstream meme PSSM #1; e=3.9e+03
upstream meme PSSM #2; e=5.8e+03
log10(P) upstream meme
log10(P.clust)=−1.99; 24 seqs; 24 uniq
−7.87 integration host factor, beta subunit: DVU2973
−6.90 hypothetical protein: DVU2625
−6.87 hypothetical protein: DVU0986
−6.55 isoamylase N−terminal domain protein: DVU0938
−1.96 peptidyl−prolyl cis−trans isomerase C: DVU1470
−1.87 heavy metal−binding domain protein: DVU0987
−1.75 chemotaxis protein methyltransferase: DVU2076
−1.66 chemotaxis protein CheW: DVU1904
−1.51 HAD−superfamily hydrolase, subfamily IA, variant 3: DVU1093
−1.51 ferritin: DVU1568
−1.25 RNA methyltransferase, TrmH family, group 3: DVU0042
−1.18 hypothetical protein: DVU2351
−1.04 acterial extracellular solute−binding domain protein: DVU2202
−0.99 cytochrome c−553: DVU1817
−0.91 hypothetical protein: DVU0754
−0.79 transcriptional regulator, TetR family: DVU0629
−0.61 hypothetical protein: DVU0974
−0.58 hypothetical protein: DVU1894
−0.55 hypothetical protein: DVU2212
−0.49 universal stress protein family: DVU0006
−0.42 hypothetical protein: DVU0238
−0.35 ferredoxin II: DVU0305
−0.17 response regulator: DVU0138
−0.00 hypothetical protein: DVU1180
DVU0754
DVU0006
DVU2351
cheW−2
DVU0138
DVU0238
cheR−2
DVU0938
DVU0629
DVU1894
ppiC
DVU1568
DVU0305
DVU1093
cyf
DVU2625
DVU2212
DVU0974
DVU1180
DVU2202
DVU0987
DVU0042
DVU0973
DVU0986
string