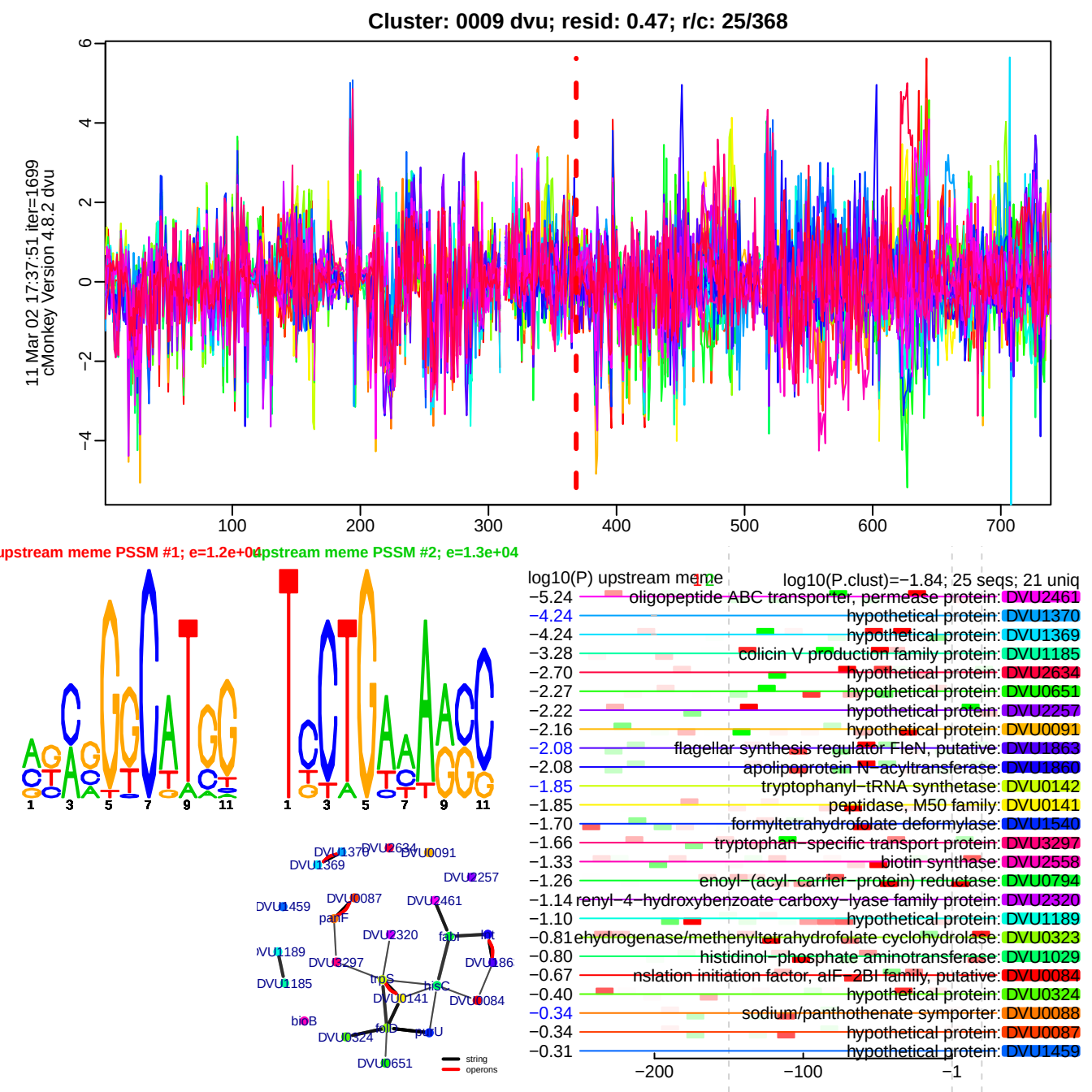
Cluster: 0009 dvu; resid: 0.47; r/c: 25/368
11 Mar 02 17:37:51 iter=1699
cMonkey Version 4.8.2 dvu
upstream meme PSSM #1; e=1.2e+04
upstream meme PSSM #2; e=1.3e+04
log10(P) upstream meme
log10(P.clust)=−1.84; 25 seqs; 21 uniq
−5.24 oligopeptide ABC transporter, permease protein: DVU2461
−4.24 hypothetical protein: DVU1370
−4.24 hypothetical protein: DVU1369
−3.28 colicin V production family protein: DVU1185
−2.70 hypothetical protein: DVU2634
−2.27 hypothetical protein: DVU0651
−2.22 hypothetical protein: DVU2257
−2.16 hypothetical protein: DVU0091
−2.08 flagellar synthesis regulator FleN, putative: DVU1863
−2.08 apolipoprotein N−acyltransferase: DVU1860
−1.85 tryptophanyl−tRNA synthetase: DVU0142
−1.85 peptidase, M50 family: DVU0141
−1.70 formyltetrahydrofolate deformylase: DVU1540
−1.66 tryptophan−specific transport protein: DVU3297
−1.33 biotin synthase: DVU2558
−1.26 enoyl−(acyl−carrier−protein) reductase: DVU0794
−1.14 renyl−4−hydroxybenzoate carboxy−lyase family protein: DVU2320
−1.10 hypothetical protein: DVU1189
−0.81 ehydrogenase/methenyltetrahydrofolate cyclohydrolase: DVU0323
−0.80 histidinol−phosphate aminotransferase: DVU1029
−0.67 nslation initiation factor, aIF−2Bl family, putative: DVU0084
−0.40 hypothetical protein: DVU0324
−0.34 sodium/panthothenate symporter: DVU0088
−0.34 hypothetical protein: DVU0087
−0.31 hypothetical protein: DVU1459
DVU1370
DVU2634
DVU0091
DVU1369
DVU2257
DVU0087
DVU2461
DVU1459
panF
DVU2320
fabI
DVU1189
DVU3297
DVU1186
DVU1185
trpS
hisC
DVU0141
DVU0084
bioB
DVU0324
purU
DVU0651
string
operons