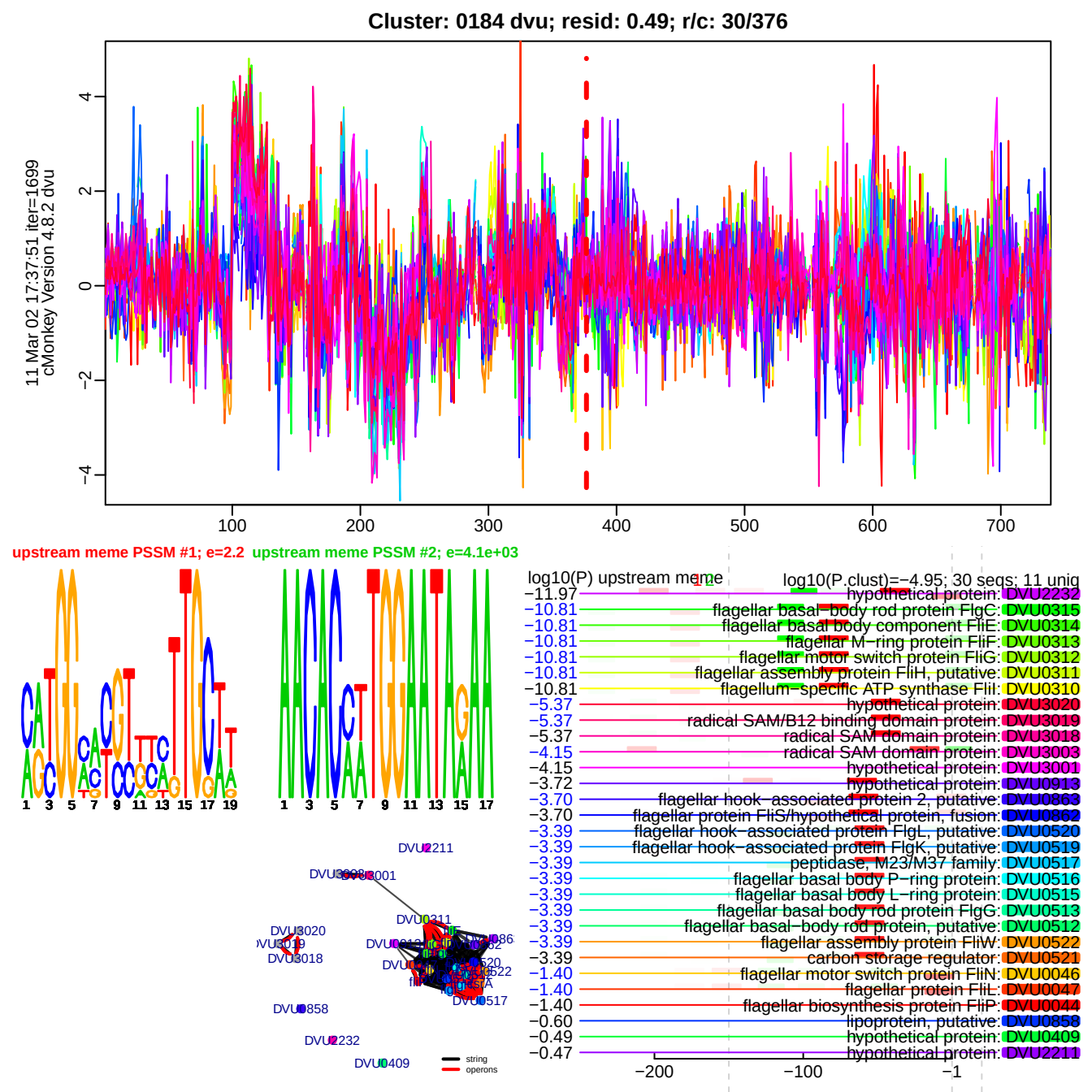
Cluster: 0184 dvu; resid: 0.49; r/c: 30/376
11 Mar 02 17:37:51 iter=1699
cMonkey Version 4.8.2 dvu
upstream meme PSSM #1; e=2.2
upstream meme PSSM #2; e=4.1e+03
log10(P) upstream meme
log10(P.clust)=−4.95; 30 seqs; 11 uniq
−11.97 hypothetical protein: DVU2232
−10.81 flagellar basal−body rod protein FlgC: DVU0315
−10.81 flagellar basal body component FliE: DVU0314
−10.81 flagellar M−ring protein FliF: DVU0313
−10.81 flagellar motor switch protein FliG: DVU0312
−10.81 flagellar assembly protein FliH, putative: DVU0311
−10.81 flagellum−specific ATP synthase FliI: DVU0310
−5.37 hypothetical protein: DVU3020
−5.37 radical SAM/B12 binding domain protein: DVU3019
−5.37 radical SAM domain protein: DVU3018
−4.15 radical SAM domain protein: DVU3003
−4.15 hypothetical protein: DVU3001
−3.72 hypothetical protein: DVU0913
−3.70 flagellar hook−associated protein 2, putative: DVU0863
−3.70 flagellar protein FliS/hypothetical protein, fusion: DVU0862
−3.39 flagellar hook−associated protein FlgL, putative: DVU0520
−3.39 flagellar hook−associated protein FlgK, putative: DVU0519
−3.39 peptidase, M23/M37 family: DVU0517
−3.39 flagellar basal body P−ring protein: DVU0516
−3.39 flagellar basal body L−ring protein: DVU0515
−3.39 flagellar basal body rod protein FlgG: DVU0513
−3.39 flagellar basal−body rod protein, putative: DVU0512
−3.39 flagellar assembly protein FliW: DVU0522
−3.39 carbon storage regulator: DVU0521
−1.40 flagellar motor switch protein FliN: DVU0046
−1.40 flagellar protein FliL: DVU0047
−1.40 flagellar biosynthesis protein FliP: DVU0044
−0.60 lipoprotein, putative: DVU0858
−0.49 hypothetical protein: DVU0409
−0.47 hypothetical protein: DVU2211
DVU2211
DVU3020
DVU3019
DVU3018
DVU0311
DVU0517
DVU0858
DVU2232
DVU0409
string
operons