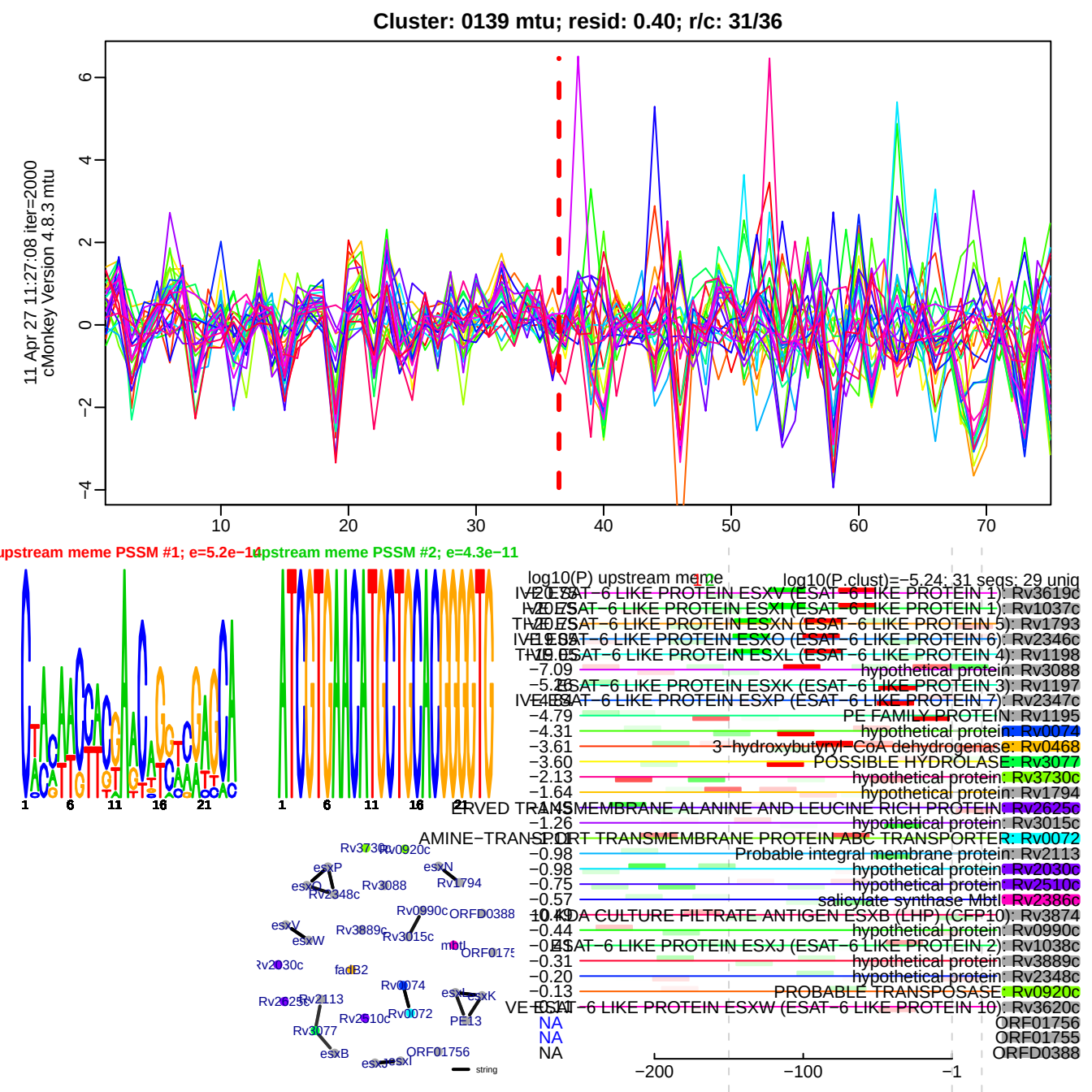
Cluster: 0139 mtu; resid: 0.40; r/c: 31/36
11 Apr 27 11:27:08 iter=2000
cMonkey Version 4.8.3 mtu
upstream meme PSSM #1; e=5.2e−14
upstream meme PSSM #2; e=4.3e−11
log10(P) upstream meme
log10(P clust)=−5.24: 31 seqs: 29 uniq
−7.09 hypothetical protein: Rv3088
−4.79 PE FAMILY PROTEIN: Rv1195
−4.31 hypothetical protein: Rv0074
−3.61 3−hydroxybutyryl−CoA dehydrogenase: Rv0468
−3.60 POSSIBLE HYDROLASE: Rv3077
−2.13 hypothetical protein: Rv3730c
−1.64 hypothetical protein: Rv1794
−1.26 hypothetical protein: Rv3015c
−0.98 Probable integral membrane protein: Rv2113
−0.98 hypothetical protein: Rv2030c
−0.75 hypothetical protein: Rv2510c
−0.57 salicylate synthase MbtI: Rv2386c
−0.44 hypothetical protein: Rv0990c
−0.31 hypothetical protein: Rv3889c
−0.20 hypothetical protein: Rv2348c
−0.13 PROBABLE TRANSPOSASE: Rv0920c
NA
NA
NA
ORF01756
ORF01755
ORFD0388
−200
−100
−1
string