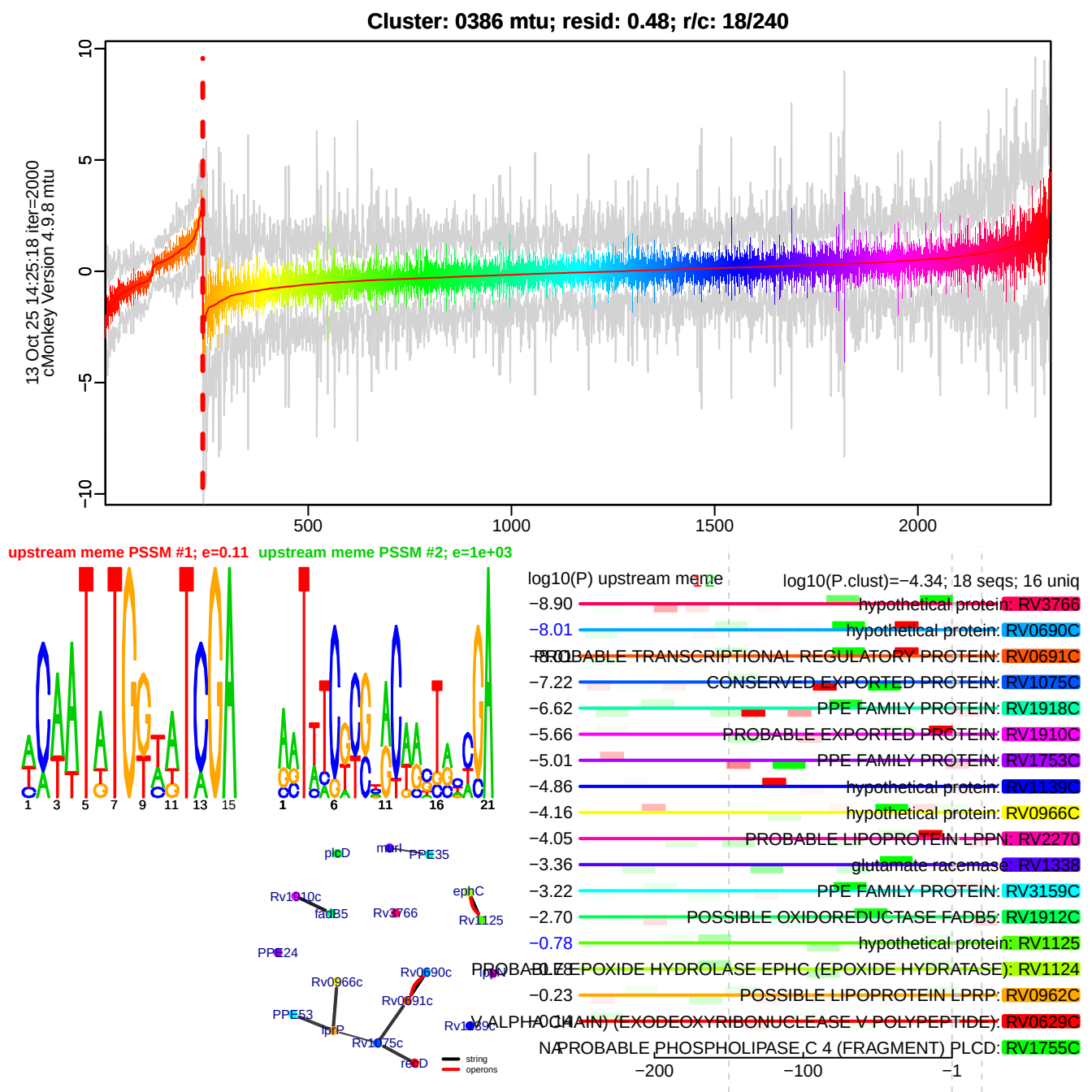
Cluster: 0386 mtu; resid: 0.48; r/c: 18/240
13 Oct 25 14:25:18 iter=2000
cMonkey Version 4.9.8 mtu
upstream meme PSSM #1; e=0.11
upstream meme PSSM #2; e=1e+03
log10(P) upstream meme
log10(P.clust)=−4.34; 18 seqs; 16 uniq
−8.90 hypothetical protein: RV3766
−8.01 hypothetical protein: RV0690C
PROBABLE TRANSCRIPTIONAL REGULATORY PROTEIN: RV0691C
−7.22 CONSERVED EXPORTED PROTEIN: RV1075C
−6.62 PPE FAMILY PROTEIN: RV1918C
−5.66 PROBABLE EXPORTED PROTEIN: RV1910C
−5.01 PPE FAMILY PROTEIN: RV1753C
−4.86 hypothetical protein: RV1139C
−4.16 hypothetical protein: RV0966C
−4.05 PROBABLE LIPOPROTEIN LPPN: RV2270
−3.36 glutamate racemase: RV1338
−3.22 PPE FAMILY PROTEIN: RV3159C
−2.70 POSSIBLE OXIDOREDUCTASE FADB5: RV1912C
−0.78 hypothetical protein: RV1125
PROBABLE EPOXIDE HYDROLASE EPHC (EPOXIDE HYDRATASE): RV1124
−0.23 POSSIBLE LIPOPROTEIN LPRP: RV0962C
V ALPHA CHAIN) (EXODEOXYRIBONUCLEASE V POLYPEPTIDE): RV0629C
NA PROBABLE PHOSPHOLIPASE C 4 (FRAGMENT) PLCD: RV1755C
plcD
mprA
PPE35
Rv1910c
fadB5
Rv3766
ephC
Rv1125
PPE24
Rv0690c
Rv0966c
Rv0691c
PPE53
lprP
Rv1075c
recD
string
operons