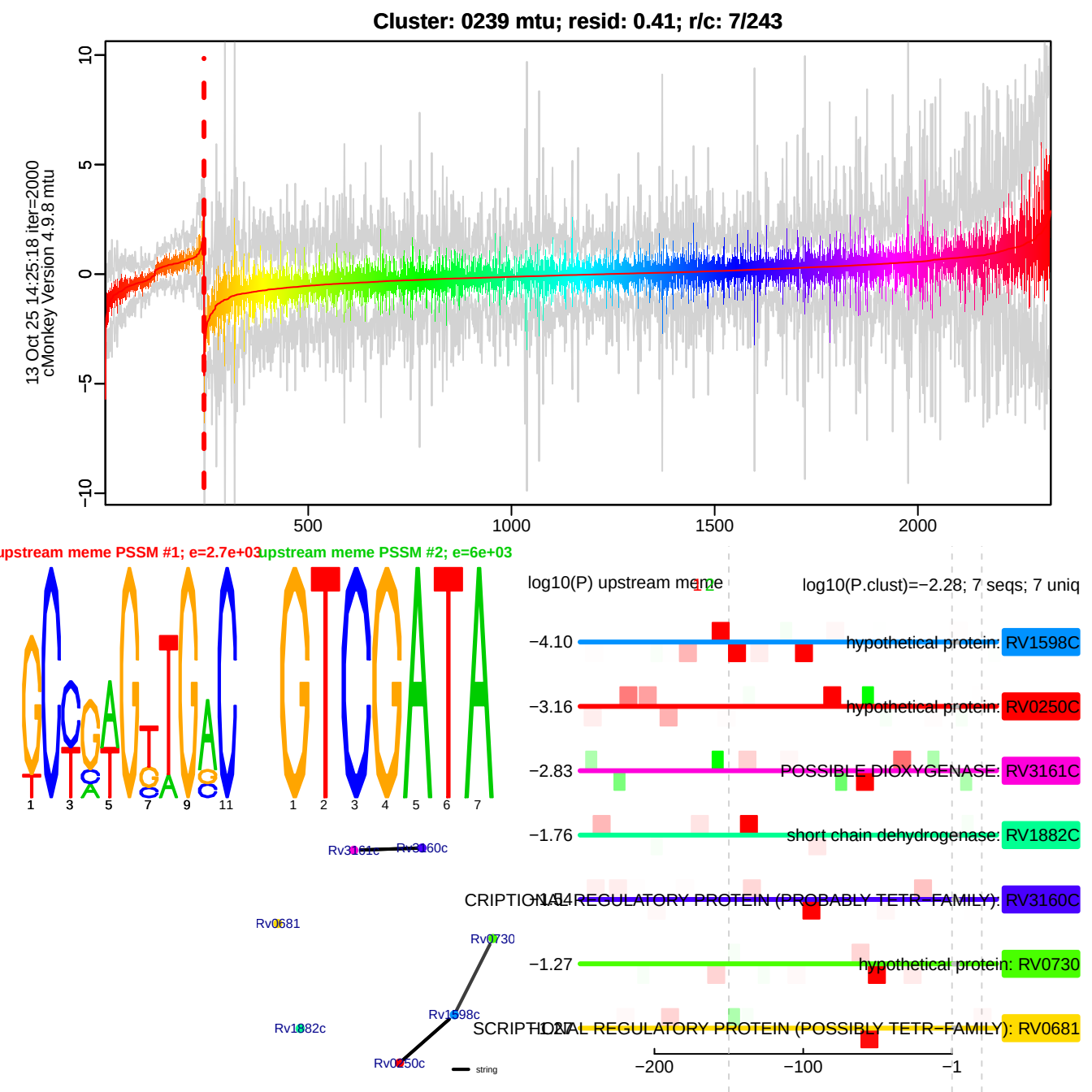
Cluster: 0239 mtu; resid: 0.41; r/c: 7/243
13 Oct 25 14:25:18 iter=2000
cMonkey Version 4.9.8 mtu
upstream meme PSSM #1; e=2.7e+03
upstream meme PSSM #2; e=6e+03
log10(P) upstream meme
log10(P.clust)=−2.28; 7 seqs; 7 uniq
−4.10 hypothetical protein: RV1598C
−3.16 hypothetical protein: RV0250C
−2.83 POSSIBLE DIOXYGENASE: RV3161C
−1.76 short chain dehydrogenase: RV1882C
−1.54 CRIPTIONAL REGULATORY PROTEIN (PROBABLY TETR−FAMILY): RV3160C
−1.27 hypothetical protein: RV0730
−1.27 SCRIPTIONAL REGULATORY PROTEIN (POSSIBLY TETR−FAMILY): RV0681
−200
−100
−1
Rv0681
Rv0730
Rv1598c
Rv1882c
Rv0250c
string