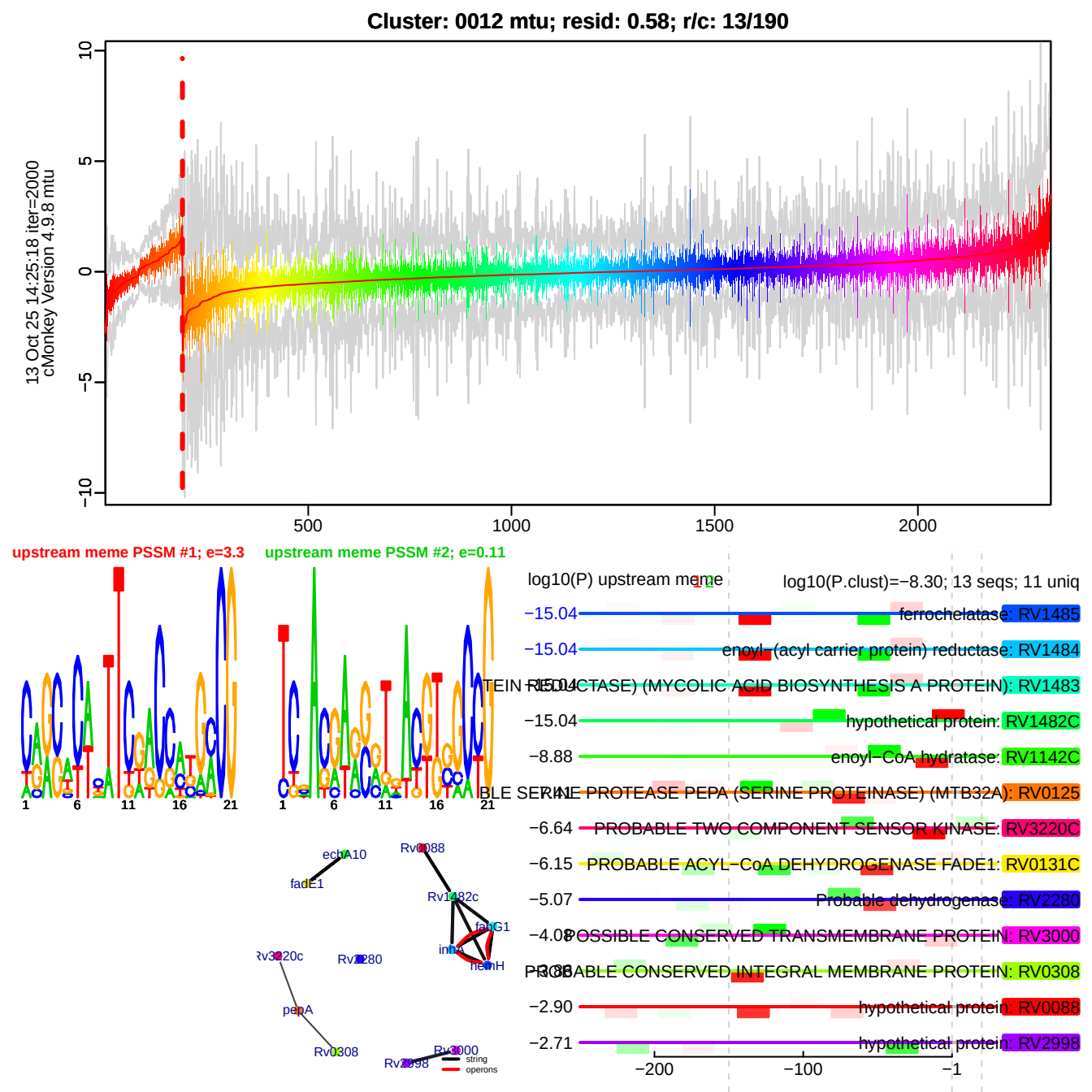
Cluster: 0012 mtu; resid: 0.58; r/c: 13/190
13 Oct 25 14:25:18 iter=2000
cMonkey Version 4.9.8 mtu
upstream meme PSSM #1; e=3.3
upstream meme PSSM #2; e=0.11
log10(P) upstream meme
log10(P.clust)=−8.30; 13 seqs; 11 uniq
−15.04 ferrochelatase: RV1485
−15.04 enoyl (acyl carrier protein) reductase: RV1484
−15.04 TEIN REDUCTASE) (MYCOLIC ACID BIOSYNTHESIS A PROTEIN): RV1483
−15.04 hypothetical protein: RV1482C
−8.88 enoyl−CoA hydratase: RV1142C
−7.41 BLE SERINE PROTEASE PEPA (SERINE PROTEINASE) (MTB32A): RV0125
−6.64 PROBABLE TWO COMPONENT SENSOR KINASE: RV3220C
−6.15 PROBABLE ACYL−CoA DEHYDROGENASE FADE1: RV0131C
−5.07 Probable dehydrogenase: RV2280
−4.08 POSSIBLE CONSERVED TRANSMEMBRANE PROTEIN: RV3000
−3.36 PROBABLE CONSERVED INTEGRAL MEMBRANE PROTEIN: RV0308
−2.90 hypothetical protein: RV0088
−2.71 hypothetical protein: RV2998
echA10
fadE1
Rv0088
Rv1482c
fadG1
inhA
hemH
Rv3220c
Rv2280
pepA
Rv0308
Rv2998
Rv3000
string
operons