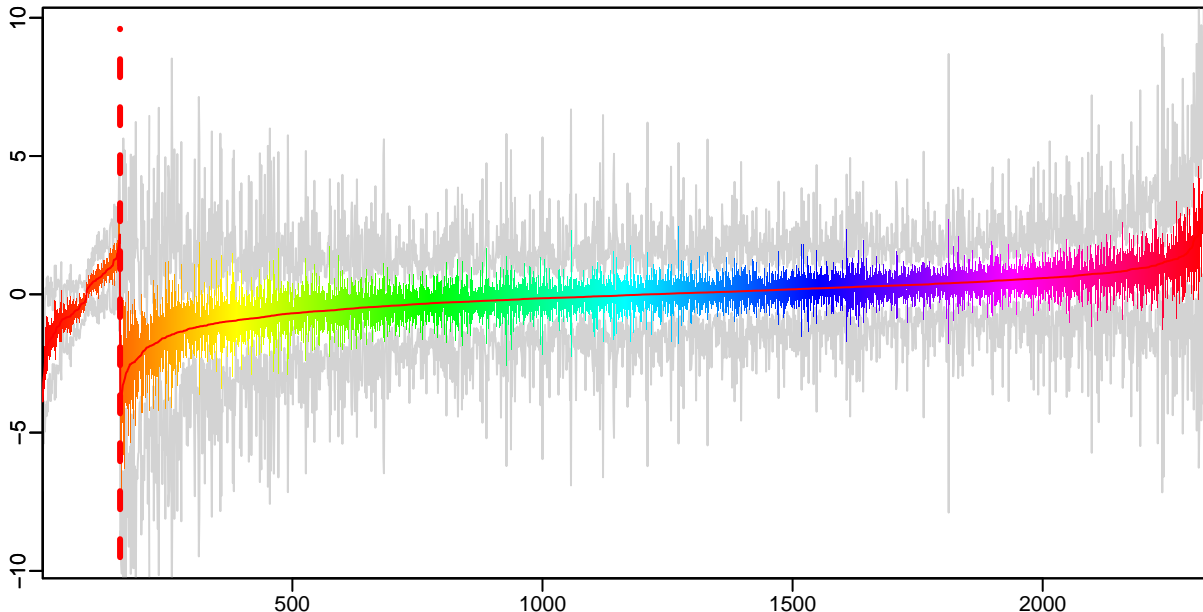
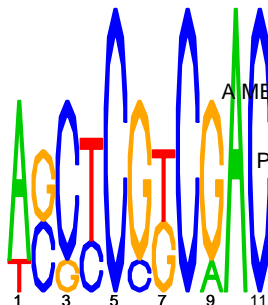
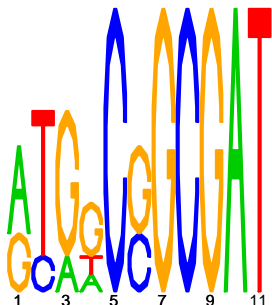


Cluster: 0015 mtu; resid: 0.46; r/c: 8/155

13 Oct 25 14:25:18 iter=2000
cMonkey Version 4.9.8 mtu



upstream meme PSSM #1; e=2.5e+00 upstream meme PSSM #2; e=2.9e+03



log10(P) upstream meme

log10(P.clust)=-4.10; 8 seqs; 6 uniq

AL METHYLASE HSDM (M PROTEIN) (DNA METHYLTRANSFERASE): RV2756C

PECIFICITY DETERMINANT (FRAGMENT) HSDS_1 (S PROTEIN): RV2755C

-4.68 hypothetical protein: RV0434

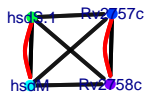
-4.42 hypothetical protein: RV2758C

-4.42 hypothetical protein: RV2757C

-4.22 POSSIBLE MUGONOLACTONE ISOMERASE: RV0316

PROBABLE MEMBRANE-ASSOCIATED PHOSPHOLIPASE C 2 PLCB: RV2350C

AL MEMBRANE PROTEIN KGTP (DICARBOXYLATE TRANSPORTER): RV3476C



kgpP

plcB

Rv0434

— string
— operons

-200 -100 -1