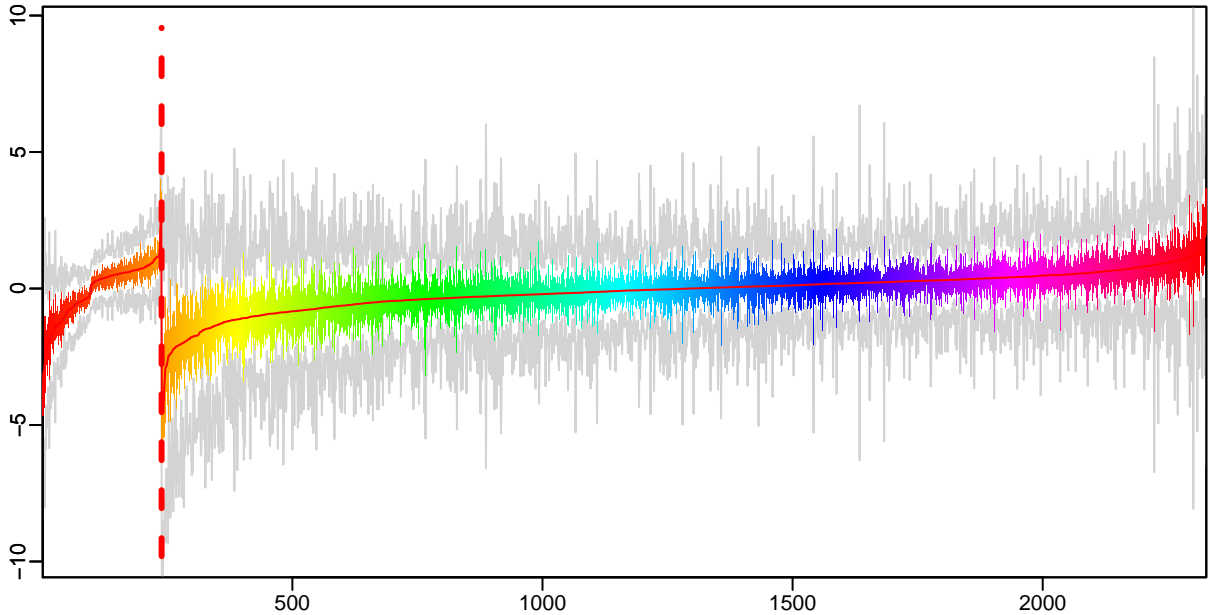
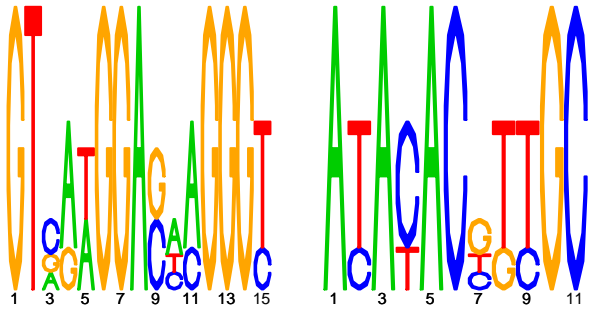


Cluster: 0275 mtu; resid: 0.52; r/c: 13/238

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



upstream meme PSSM #1; e=550 upstream meme PSSM #2; e=2.1e+03



- log10(P) upstream meme log10(P.clust)=-5.23; 13 seqs; 6 uniq
- 7.36 — SUPEROXIDE DISMUTASE [FE] SODA: RV3846
 - 6.43 — BTILIN-LIKE PROTEASE (SUBTILASE-LIKE) (MYGOSIN-2): RV3886C
 - 6.43 — POSSIBLE ESAT-6 LIKE PROTEIN ESXD: RV3891C
 - 6.43 — ESAT-6 LIKE PROTEIN ESXC (ESAT-6 like protein 11): RV3890C
 - 6.43 — hypothetical protein: RV3889C
 - 6.43 — PROBABLE CONSERVED MEMBRANE PROTEIN: RV3888C
 - 6.43 — PROBABLE CONSERVED TRANSMEMBRANE PROTEIN: RV3887C
 - 6.43 — PROBABLE CONSERVED MEMBRANE PROTEIN: RV3885C
 - 6.11 — hypothetical protein: RV3718C
 - 3.35 — Possible ketoacyl reductase: RV1544
 - 3.35 — POSSIBLE FATTY ACYL-CoA REDUCTASE: RV1543
 - 2.44 — PROBABLE CBXX/CFQX FAMILY PROTEIN: RV3884C
 - 2.44 — ASE (ADENYLYL-OR GUANYLYL) (ADENYLATE-OR GUANYLATE): RV2435C

