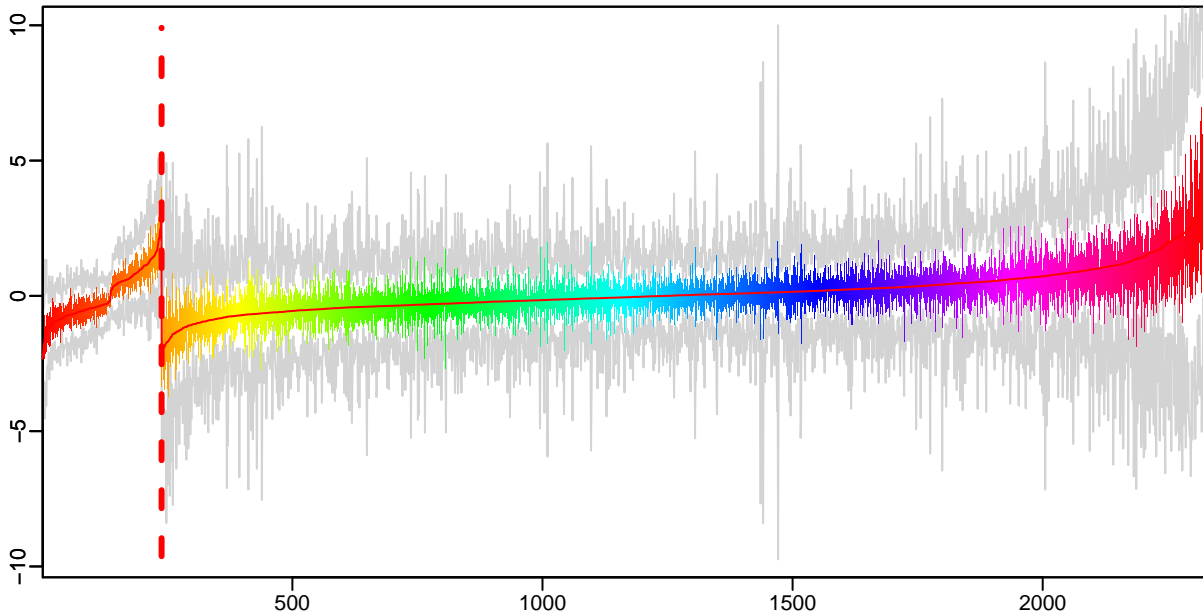


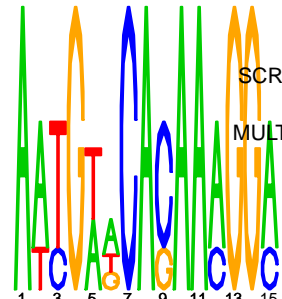
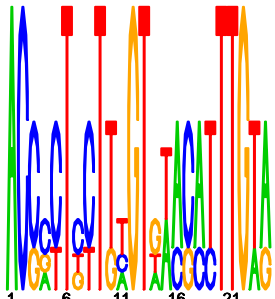
Cluster: 0385 mtu; resid: 0.48; r/c: 10/238

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



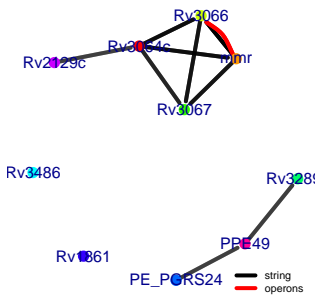
upstream meme PSSM #1; e=0.63

upstream meme PSSM #2; e=9.2



log10(P) upstream meme      log10(P.clust)=-7.42; 10 seqs; 9 uniq

- DESCRIPTION: TRANSCRIPTIONAL REGULATORY PROTEIN (PROBABLY DEOR-FAMILY): RV3066
- MULTIDRUG TRANSPORT INTEGRAL MEMBRANE PROTEIN MMR: RV3065
- PROBABLE CONSERVED INTEGRAL MEMBRANE PROTEIN: RV3064C
- 10.92 PPE FAMILY PROTEIN: RV3125C
- 6.37 PROBABLE CONSERVED TRANSMEMBRANE PROTEIN: RV1861
- 3.67 PE\_PGRS FAMILY PROTEIN: RV1325C
- 0.46 POSSIBLE TRANSMEMBRANE PROTEIN: RV3289C
- 0.21 short chain dehydrogenase: RV2129C
- 0.19 hypothetical protein: RV3486
- 0.04 hypothetical protein: RV3067



-200      -100      -1