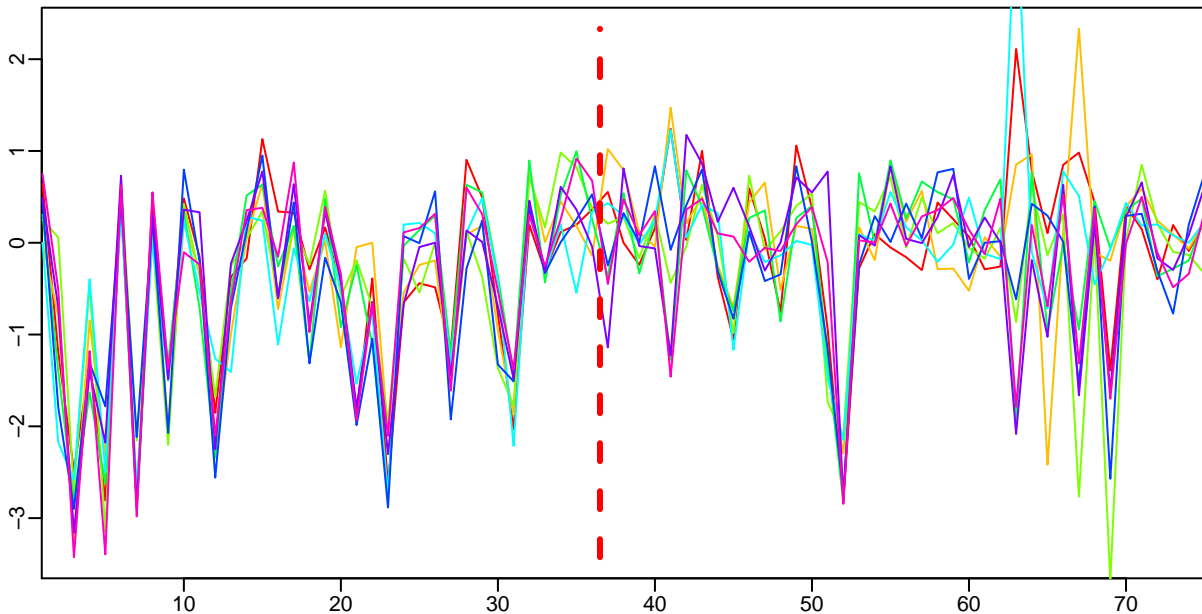
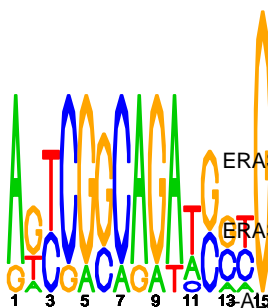
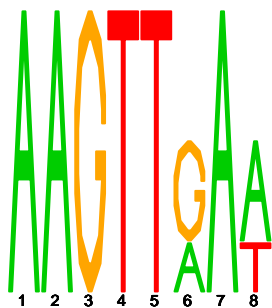


Cluster: 0105 mtu; resid: 0.27; r/c: 8/36

11 Apr 27 11:27:08 iter=2000  
cMonkey Version 4.8.3 mtu



upstream meme PSSM #1;  $e=5.1e+00$  upstream meme PSSM #2;  $e=7.8e+03$



log10(P) upstream meme

log10(P.clust)=-3.36; 8 seqs; 7 uniq

-4.29 30S ribosomal protein S3; Rv0707

ERASE EMB7 (ARABINOSYLINDOLYLACETYLINOSITOL SYNTHASE); Rv3795

ERASE EMB7 (ARABINOSYLINDOLYLACETYLINOSITOL SYNTHASE); Rv3794

ALV-D-ALV CARBOXYPEPTIDASE (D-AMINO ACID HYDROLASE); Rv2911

embA -3.44 hypothetical protein; Rv0433

Rv2953 PROBABLY CONSERVED INTEGRAL MEMBRANE PROTEIN; Rv2120c

Rv2120c -1.79 hypothetical protein; Rv2953

Rv0433 Rv3922c -1.74 hypothetical protein; Rv3922c

— string  
— operons

-200 -100 -1