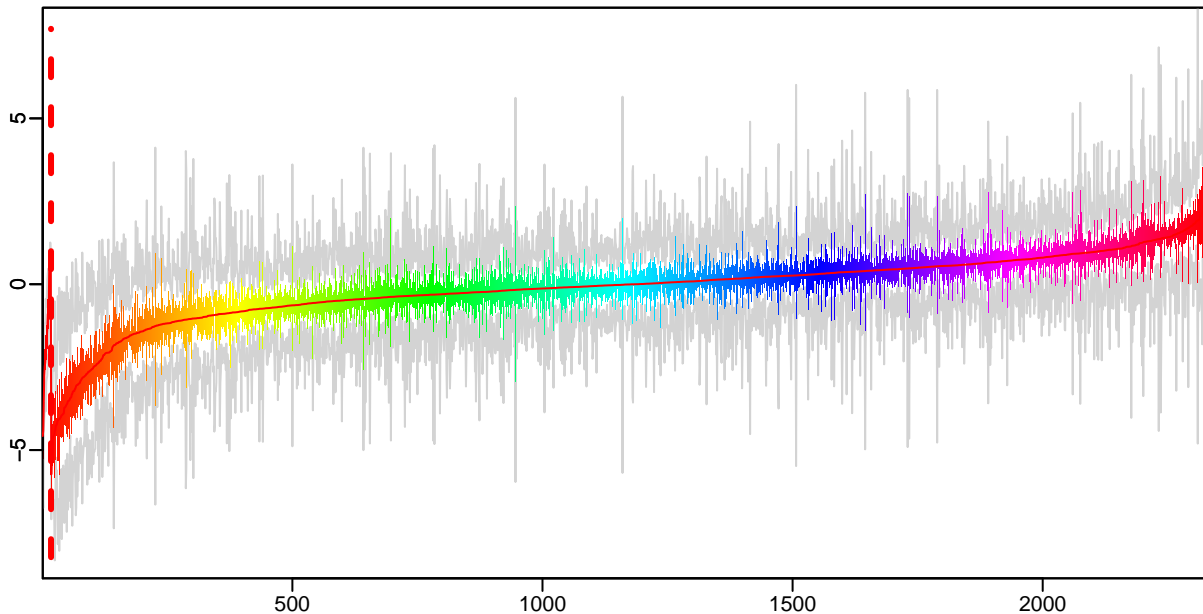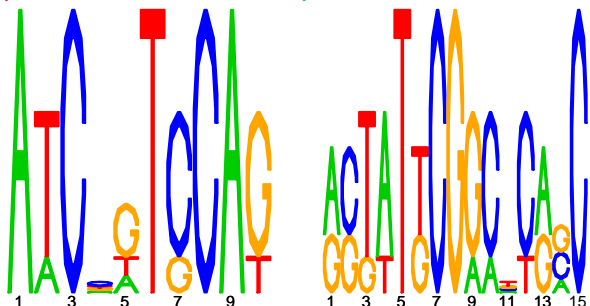# Cluster: 0354 mtu; resid: 0.44; r/c: 13/17

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

upstream meme PSSM #1; e=580 **upstream meme PSSM #2; e=4.5e+03**

log10(P) upstream meme   log10(P.clust)=−4.95; 13 seqs; 5 uniq

−5.62	30S ribosomal protein S19: RV0705
−5.62	50S ribosomal protein L2: RV0704
−5.62	50S ribosomal protein L4: RV0702
−5.62	50S ribosomal protein L3: RV0701
−5.62	30S ribosomal protein S10: RV0700
−5.22	DNA−directed RNA polymerase subunit beta': RV0668
−5.22	DNA−directed RNA polymerase subunit beta: RV0667
−4.85	phenylalanyl−tRNA synthetase subunit alpha: RV1649
−4.24	elongation factor Tu: RV0685
−4.24	elongation factor G: RV0684
−4.24	30S ribosomal protein S7: RV0683
−4.24	30S ribosomal protein S12: RV0682
−3.97	POSSIBLE BETA−1,3−GLUCANASE PRECURSOR: RV0315