

Single nucleotide frequencies in the various datasets containing 10kb long sequences (total of 214 sequences)

Nucleotide	upstream miRNA sequences	Upstream protein coding sequences (UPS)		Repeat-masked control sequences (RGS)		Unmasked control sequences (GS)	
				Standard		Standard	
		Average	deviation	Average	deviation	Average	deviation
A	0.26	0.27	0.03	0.30	0.03	0.29	0.03
T	0.28	0.27	0.03	0.30	0.03	0.29	0.03
C	0.23	0.22	0.02	0.20	0.02	0.20	0.02
G	0.23	0.22	0.02	0.20	0.02	0.20	0.02