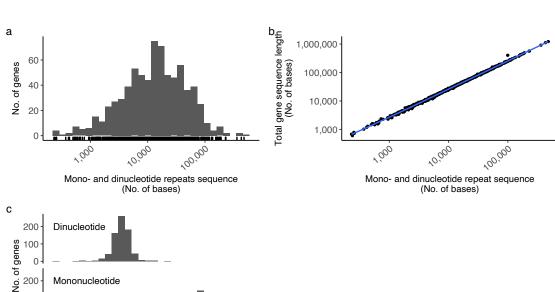
100

0.10



Ratio of bases in repeats (No. of bases in mono- and dinucleotide repeats / total gene length)

0.15

0.20

0.25