

**A**

Bias direction

DE

Non-DE

Proportion of genes with different stop codon positions

0.6

0.4

0.2

0.0

Autosome+PAR

Young strata

Genomic compartment

Old strata

**B**

Mean indel number

6

4

2

0

Autosome+PAR

Young strata

Genomic compartment

Old strata

