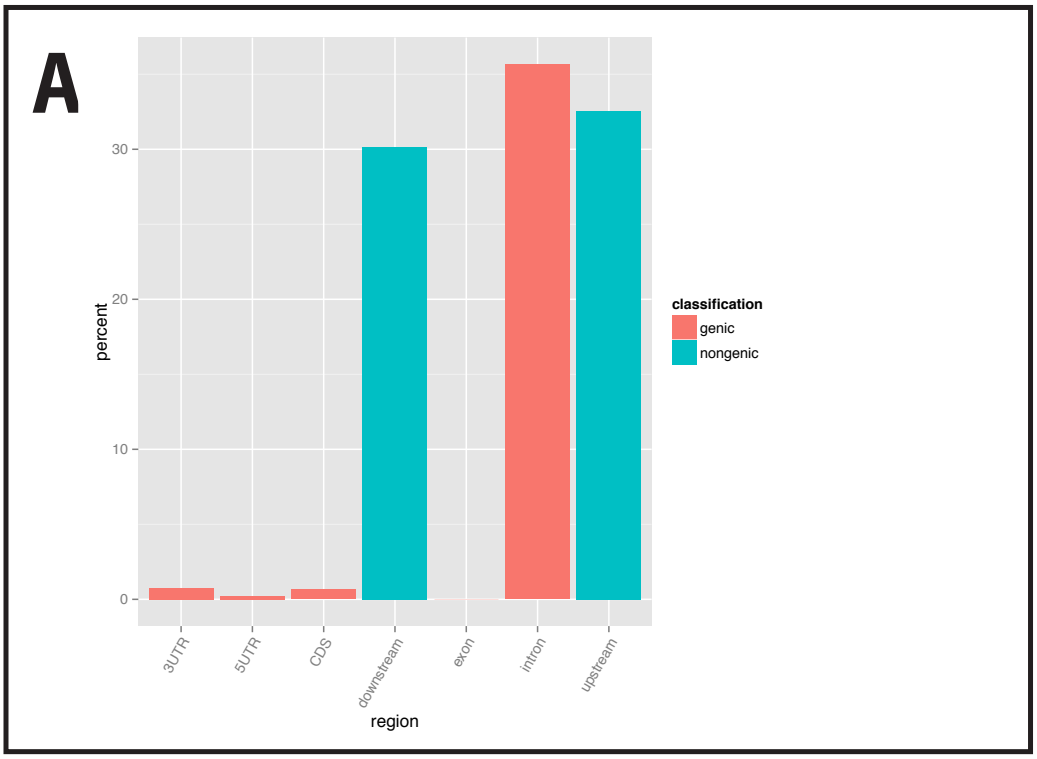
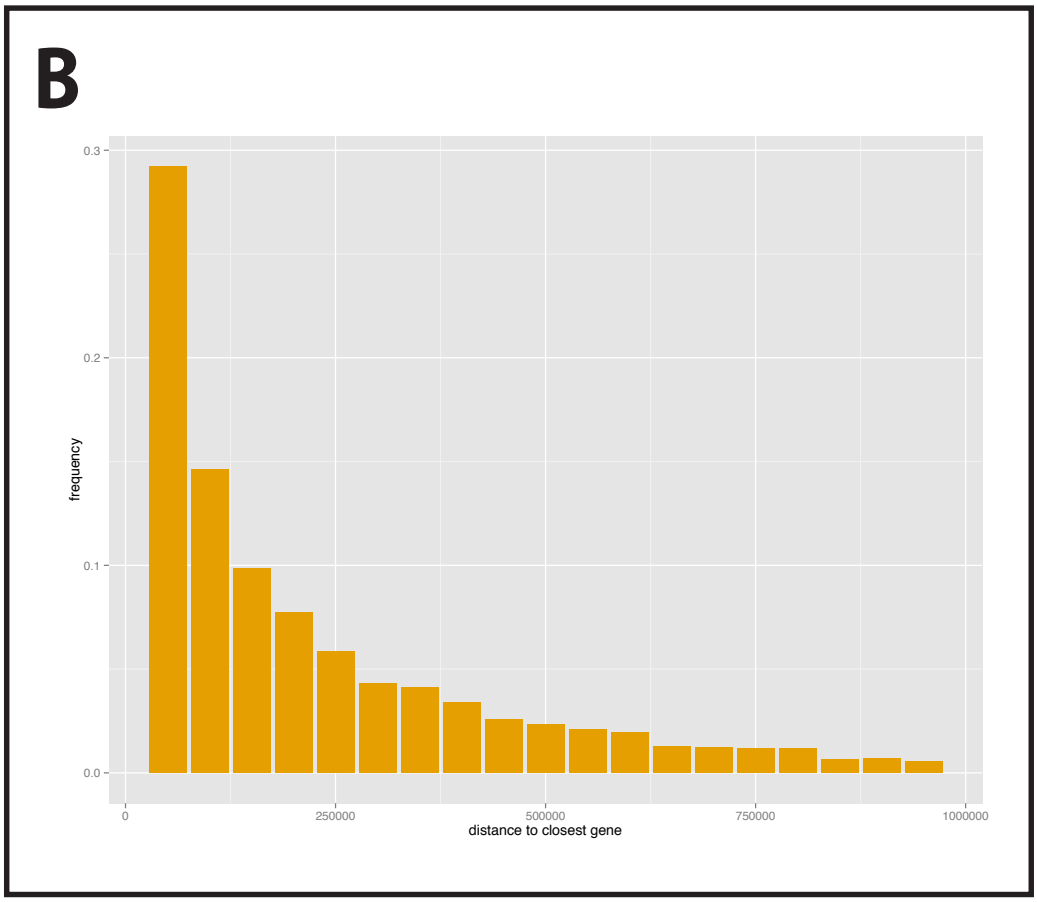
Figure S3


A
percent
30
20
10
0
3UTR
5UTR
CDS
downstream
exon
intron
upstream
region
classification
genic
nongenic


B
frequency
0.3
0.2
0.1
0.0
0
250000
500000
750000
1000000
distance to closest gene