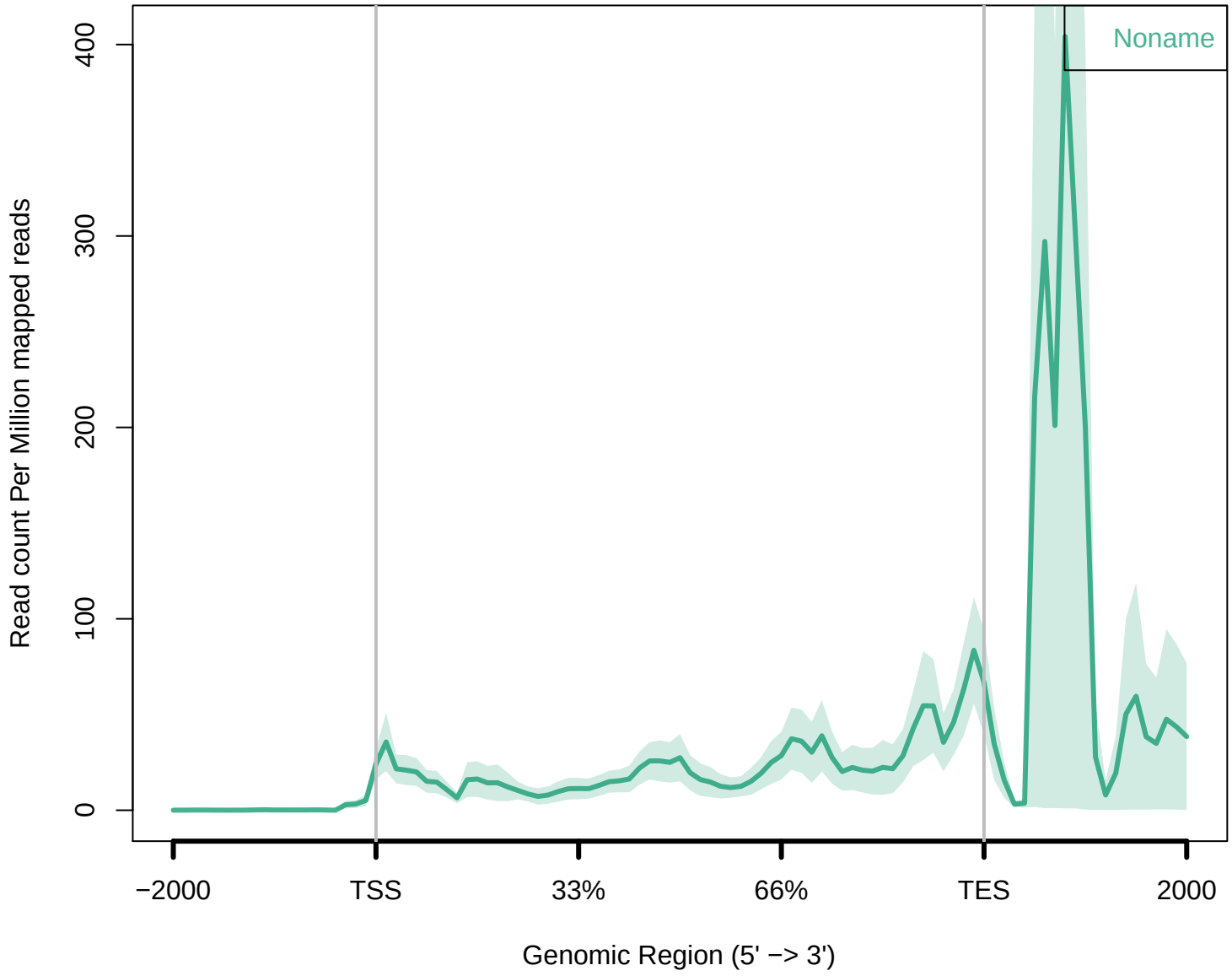
Noname
Read count Per Million mapped reads
0
100
200
300
400
−2000
TSS
33%
66%
TES
2000
Genomic Region (5' −> 3')