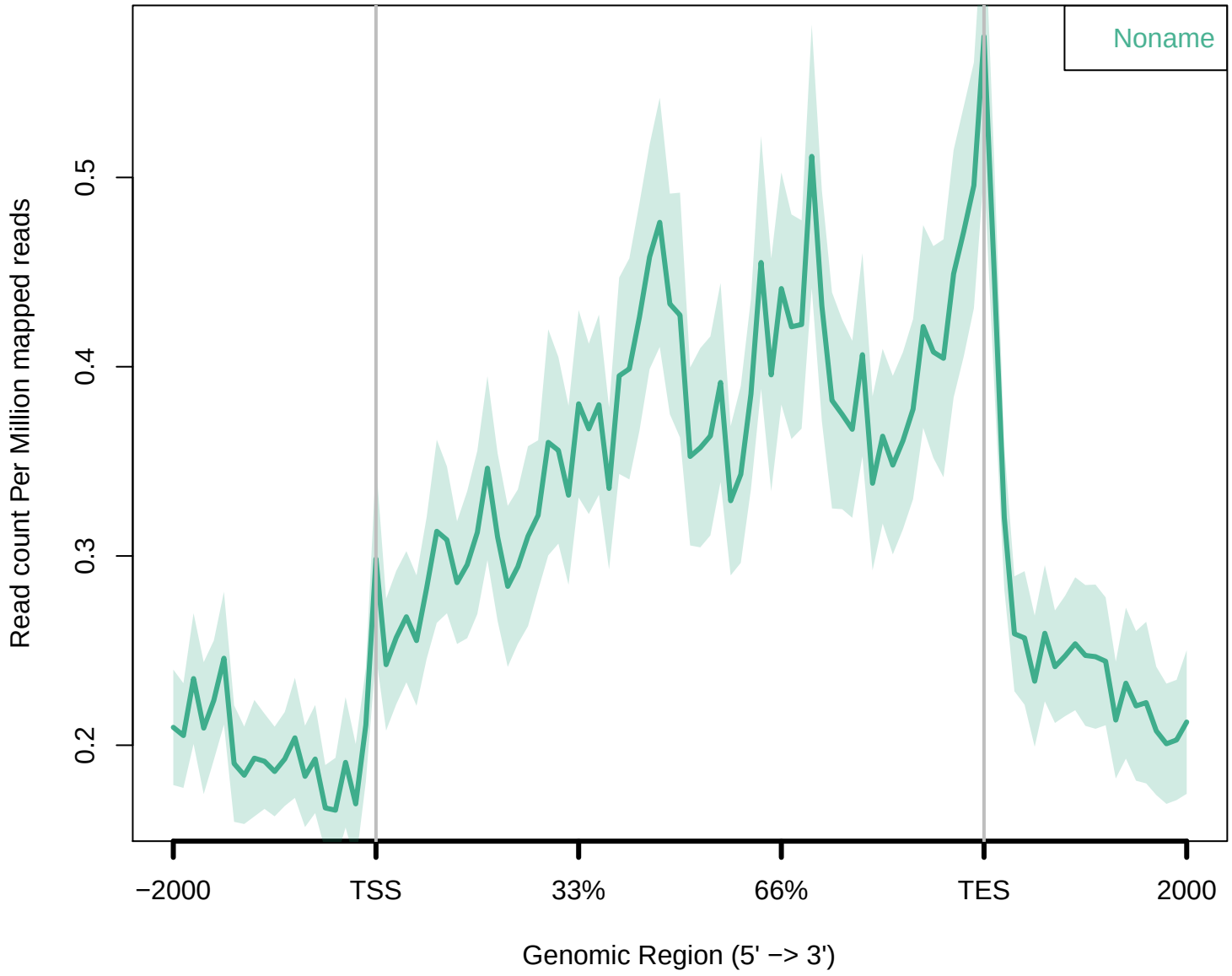
Noname
Read count Per Million mapped reads
0.2
0.3
0.4
0.5
−2000
TSS
33%
66%
TES
2000
Genomic Region (5' −> 3')