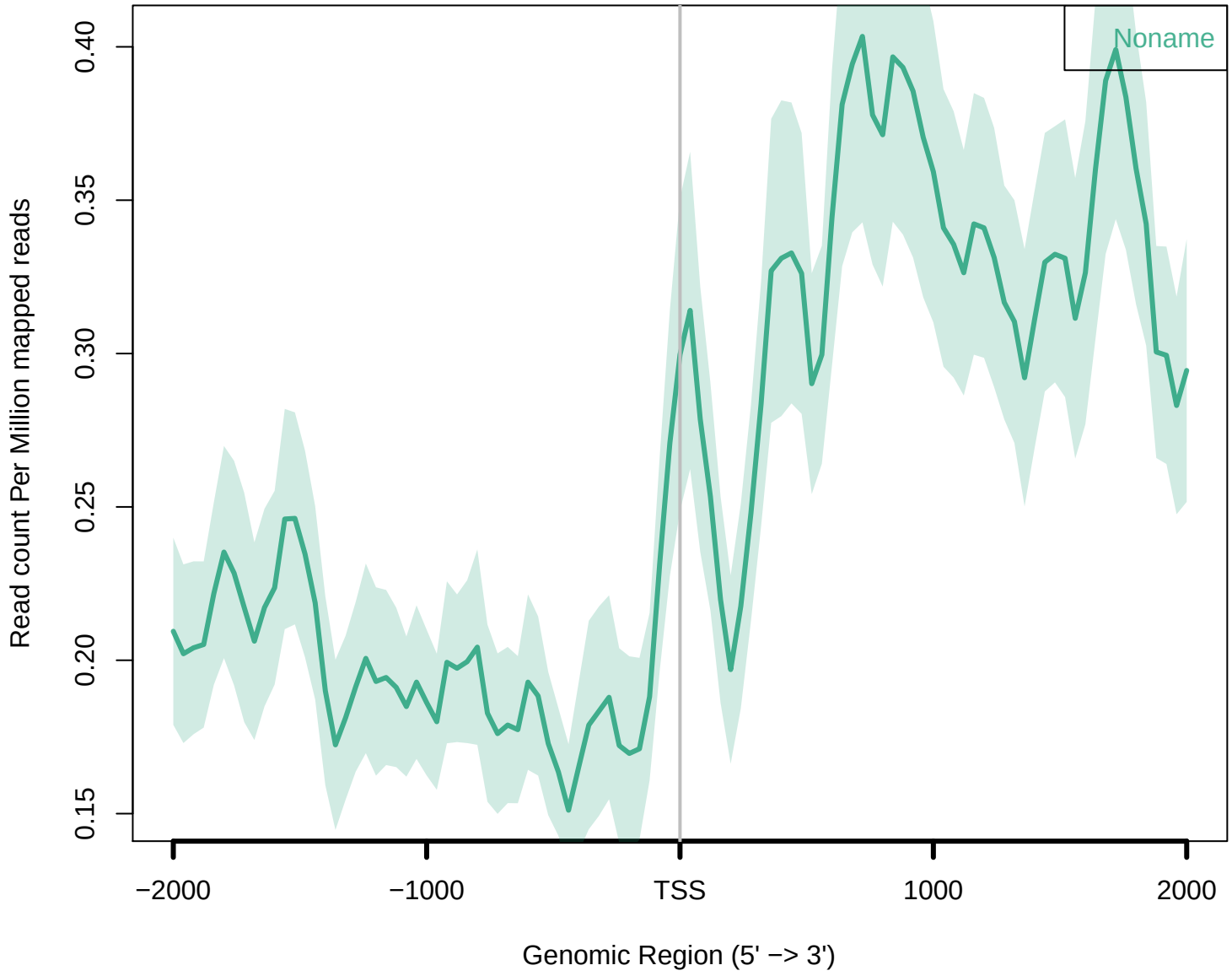
Noname
Read count Per Million mapped reads
0.15
0.20
0.25
0.30
0.35
0.40
−2000
−1000
TSS
1000
2000
Genomic Region (5' −> 3')