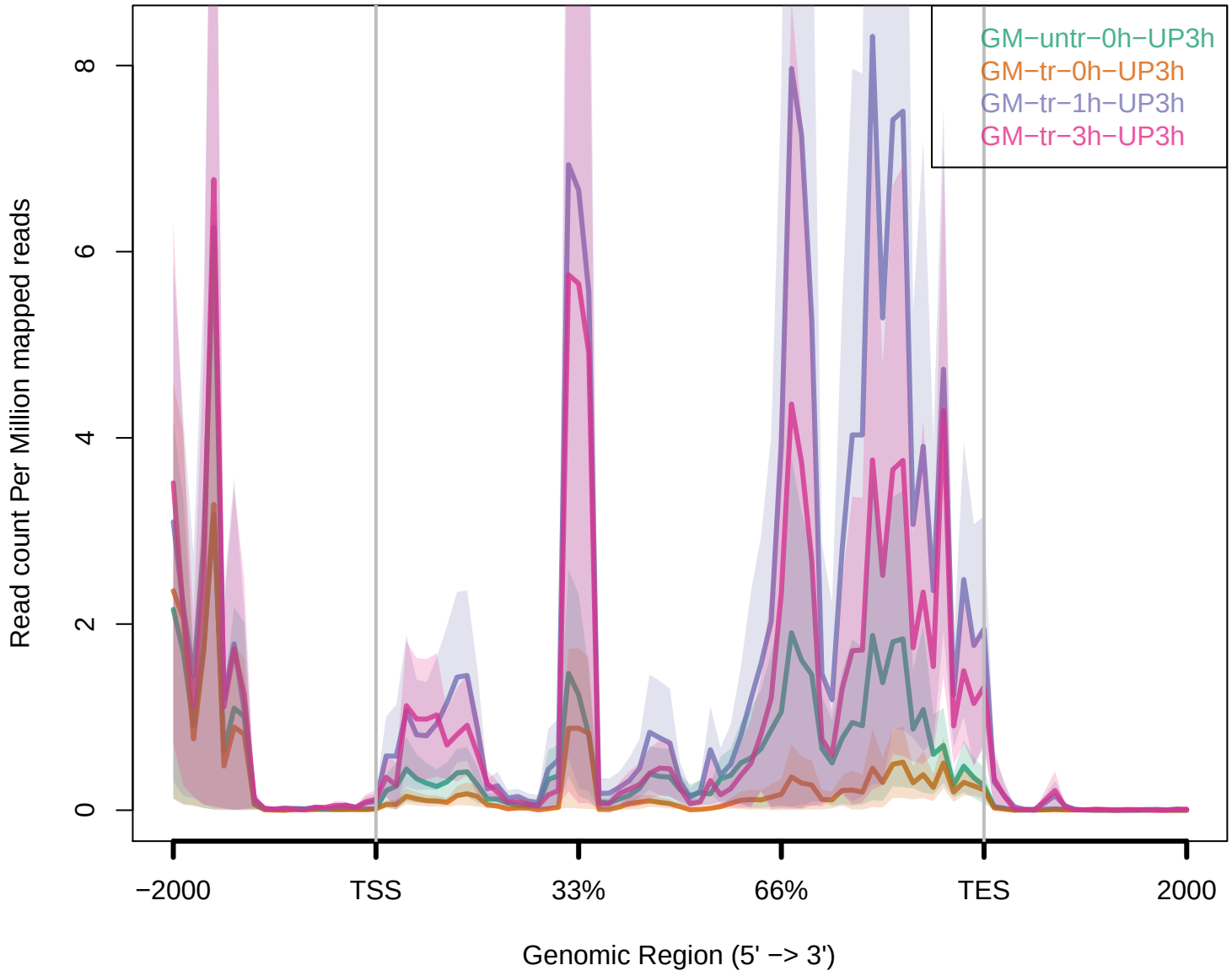
GM−untr−0h−UP3h
GM−tr−0h−UP3h
GM−tr−1h−UP3h
GM−tr−3h−UP3h
Read count Per Million mapped reads
0
2
4
6
8
−2000
TSS
33%
66%
TES
2000
Genomic Region (5' −> 3')