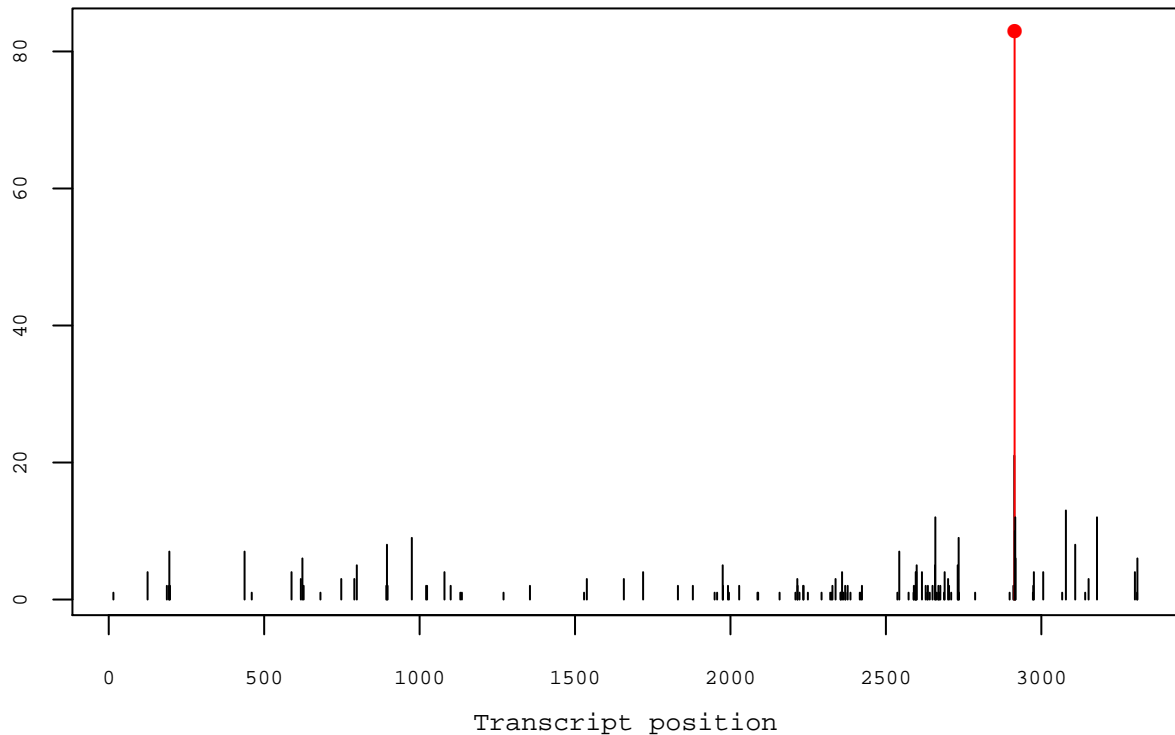


PggD2924_c32_g1_i1

5' AAATTT-GGGTGTAAATATGTAAATATATAAG '3
||| o |||||
3' AAAGTCCACAATTTATACATTTAT '5

Fragment Abundance



Cleavage site: 2914 Tag abundance: 83 Weighted abundance: 0.619 Category: 0
sRNA abundance: 5 Alignment score: 1.5 MFE ratio: 0.807 p-value: 0.025