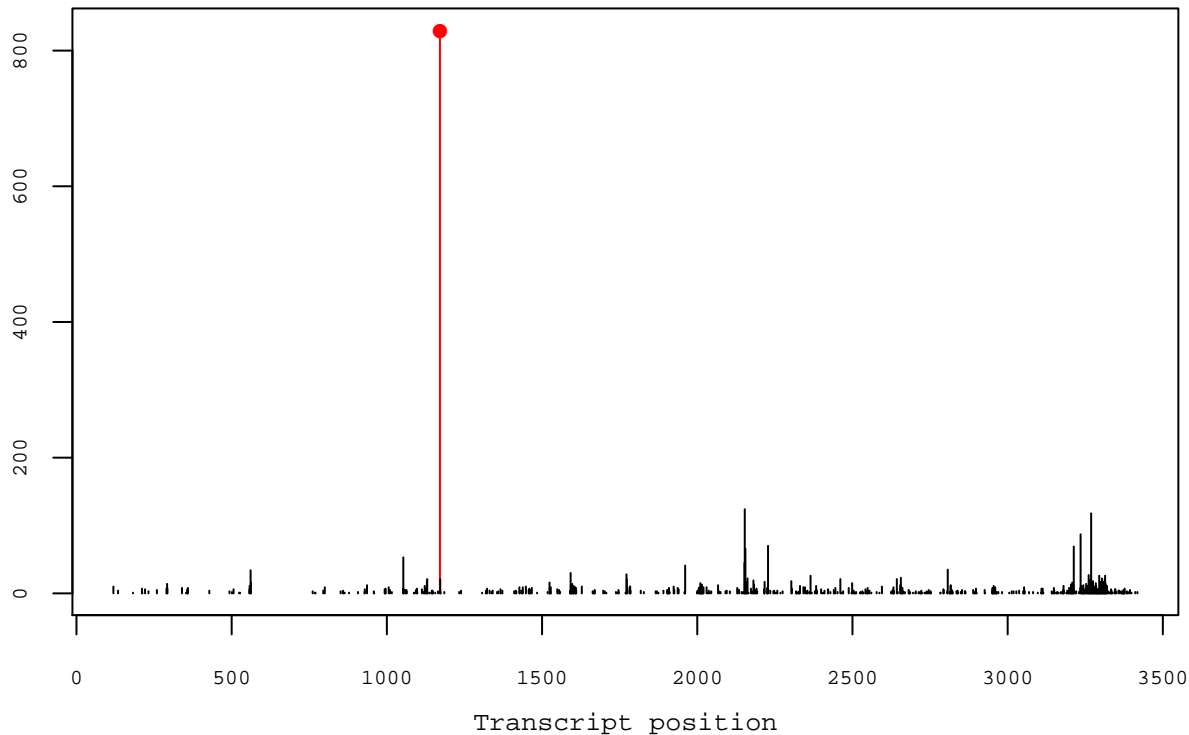


PggD9924_c6_g1_i1

5' GCTTGTGGGAATGAAGCCTGGTCCGGATTCCA '3
○ |||||
3' TCCCCTTACTTCGGACCAGGCT '5

Fragment Abundance



Cleavage site: 1171 Tag abundance: 829 Weighted abundance: 829 Category: 0
sRNA abundance: 696 Alignment score: 3 MFE ratio: 0.829 p-value: 0