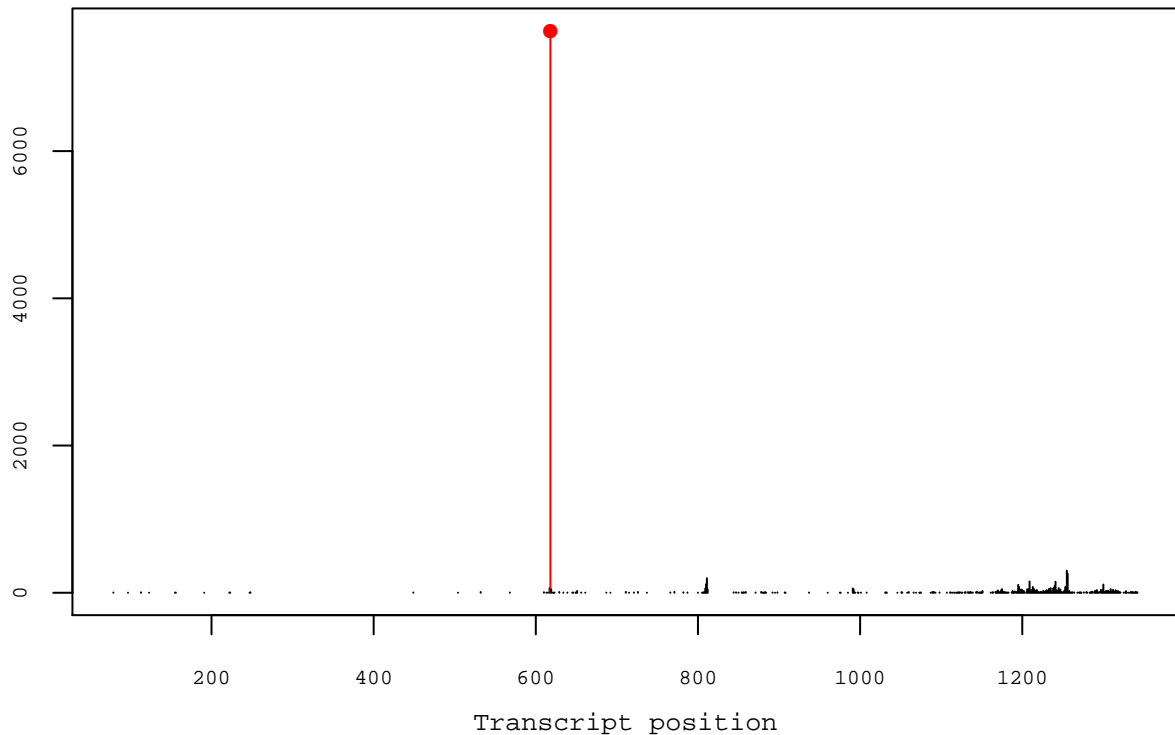


PggD6412_c14_g2_i1

5' GGAAATGGAGCTCCCTTCAAACCAAAGCCATC '3
|o|||||||||||||
3' ATCTCGAGGGAAGTTAGGTTT '5

Fragment Abundance



Cleavage site: 618 Tag abundance: 7633 Weighted abundance: 7633 Category: 0
sRNA abundance: 106556 Alignment score: 2.5 MFE ratio: 0.863 p-value: 0.001