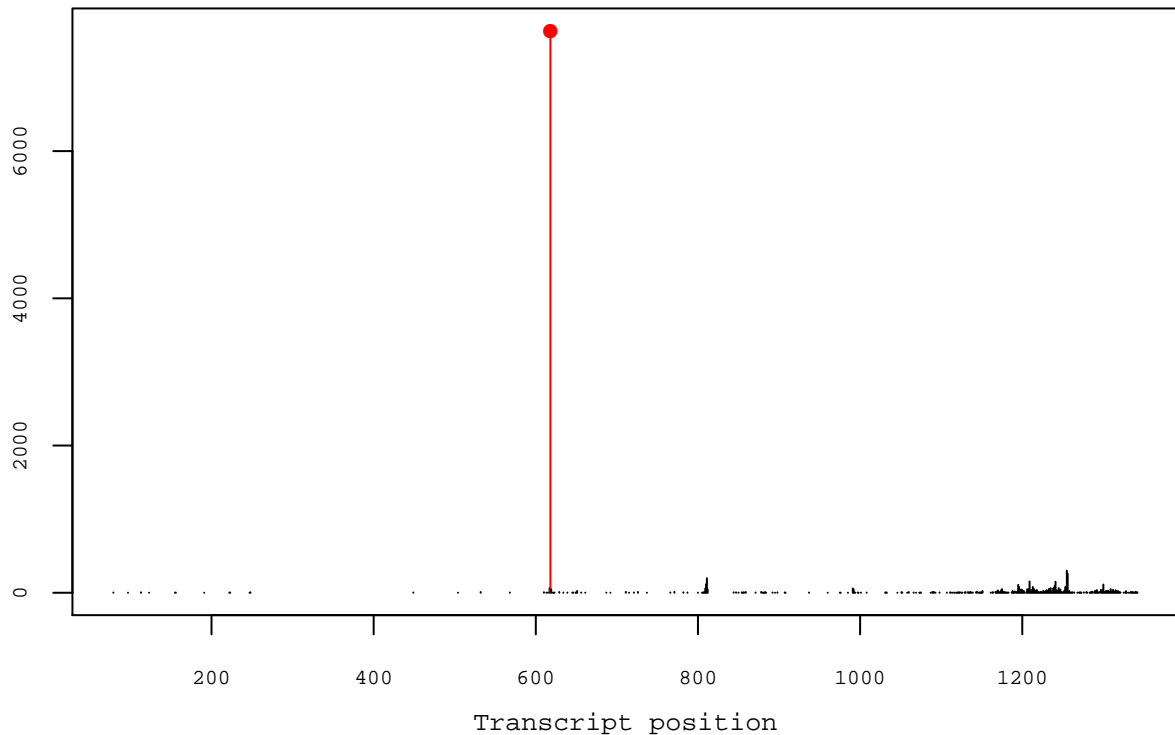


PggD6412\_c14\_g2\_i1

5' GGAAATGGAGCTCCCTTCAAACCAAAGCCATC '3  
o |||||  
3' TCTCGAGGGAAGTTAGGTTT '5

Fragment Abundance



Cleavage site: 618 Tag abundance: 7633 Weighted abundance: 7633 Category: 0  
sRNA abundance: 26628 Alignment score: 2.5 MFE ratio: 0.866 p-value: 0.001