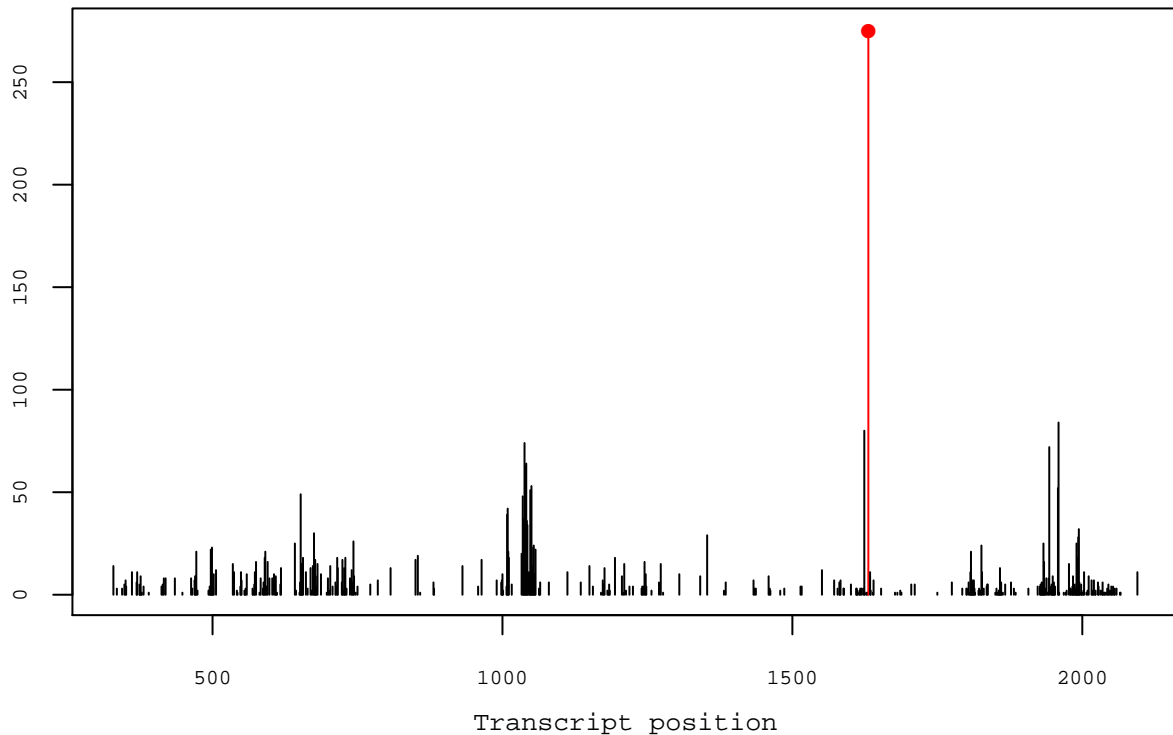


PggD1342\_c43\_g1\_i1

5' GACTCCGACTGTGCTCTCTCTCTTCTGTCATC 3'  
| | | | | o | | | | | | | | | | | | | |  
3' TCCGACATGAGAGAGAGAAGA 5'

Fragment Abundance



Cleavage site: 1631 Tag abundance: 275 Weighted abundance: 67 Category: 0  
sRNA abundance: 37 Alignment score: 2.5 MFE ratio: 0.816 p-value: 0.006