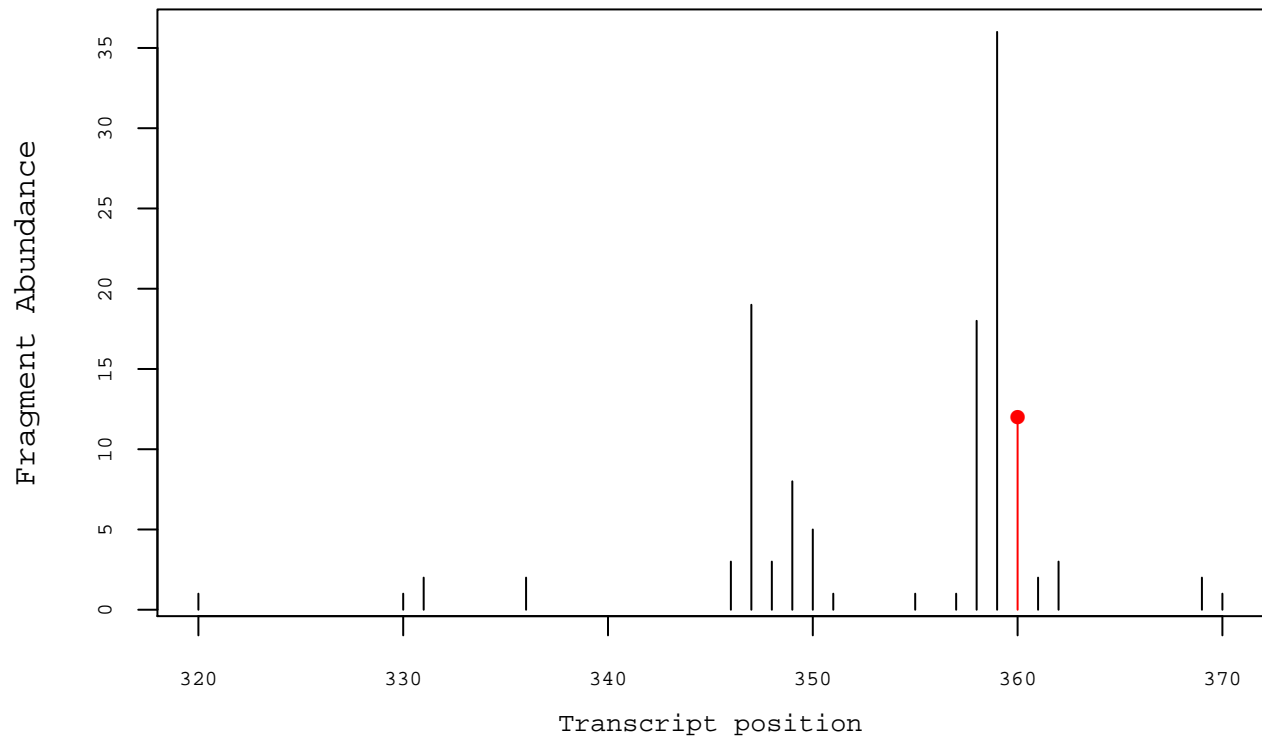


PggD15562\_c117\_g1\_i1

5' CAGCTCGTGTTCGGCTCGTTAAGAAGCTCGTG 3'  
 |||||  
 3' CGAGCACAAAGCCGAGCAATTTGTT 5'



Cleavage site: 360 Tag abundance: 12 Weighted abundance: 0.983 Category: 2  
sRNA abundance: 9 Alignment score: 3.5 MFE ratio: 0.888 p-value: 0.04