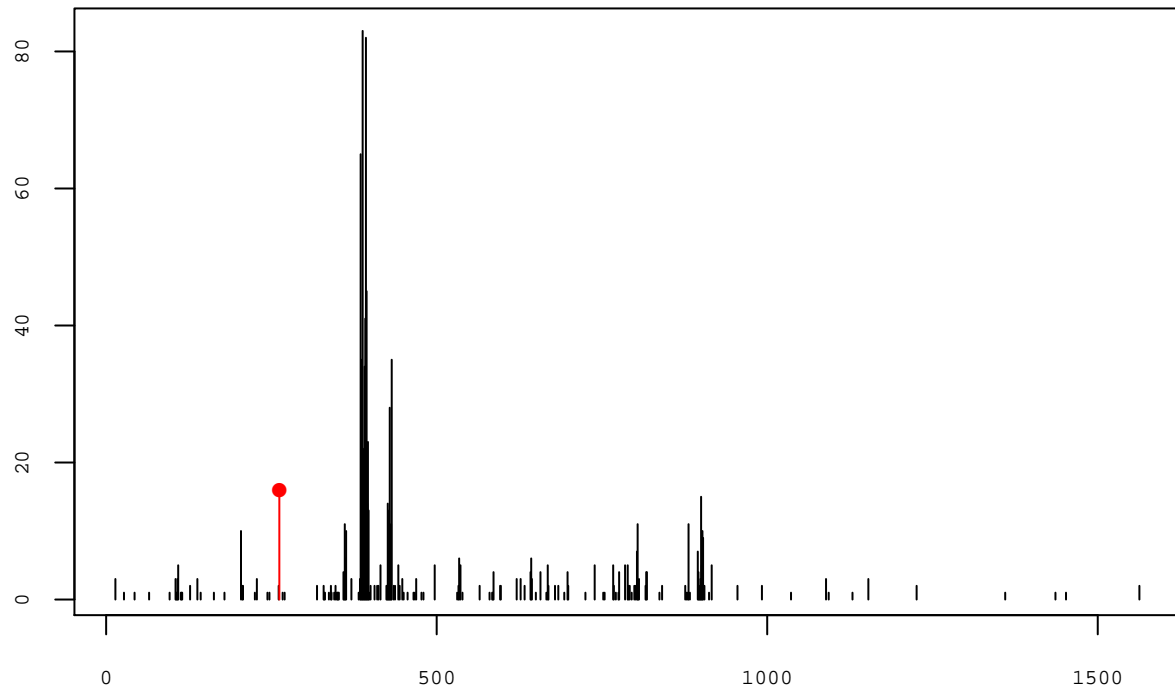


PggD12654\_c81\_g1\_i1

5' TCTTACGGCTGATACATTACTGATAAGATTA '3  
|||||  
3' CGACTATGTAAATG-CTATA '5

Fragment Abundance



Transcript position

Cleavage site: 262 Tag abundance: 16 Weighted abundance: 16 Category: 2  
sRNA abundance: 9 Alignment score: 3 MFE ratio: 0.881 p-value: 0.014