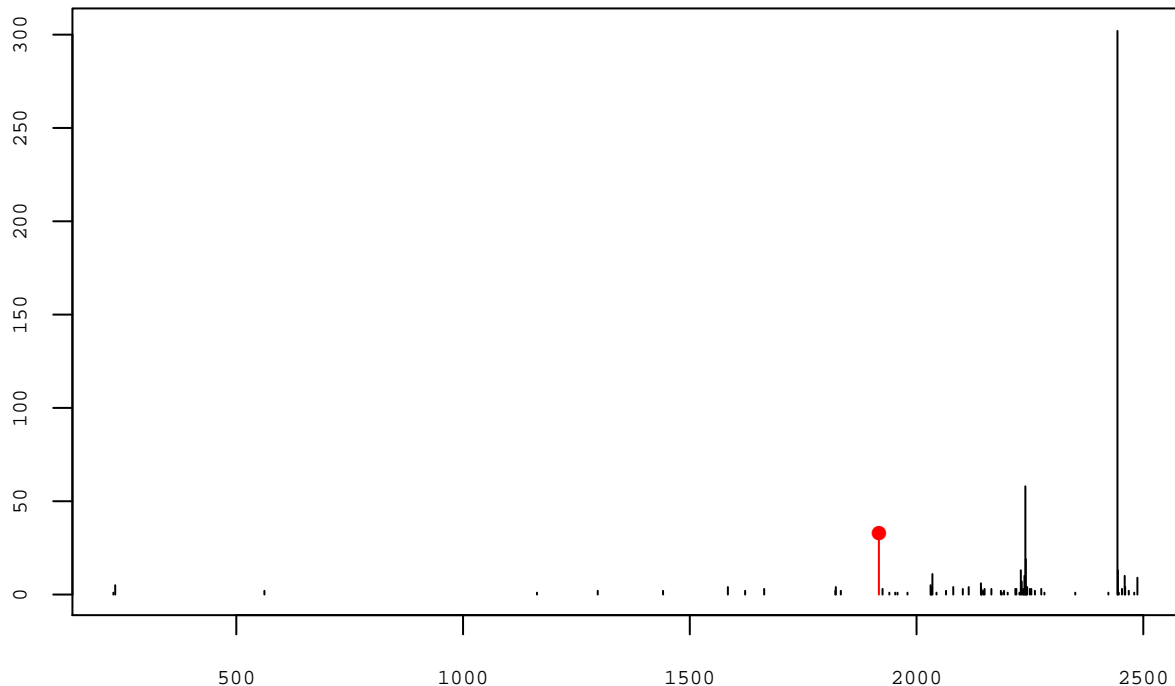


PggD12654\_c9\_g1\_i1

5' CTCTTCTGCAGCATCATCAGGATTCTCAGCCT '3  
|||||o|||||  
3' TACGTCGTAGTAGTTCTAAGA '5

Fragment Abundance



Transcript position

Cleavage site: 1917 Tag abundance: 33 Weighted abundance: 11 Category: 2  
sRNA abundance: 108 Alignment score: 2 MFE ratio: 0.983 p-value: 0.018