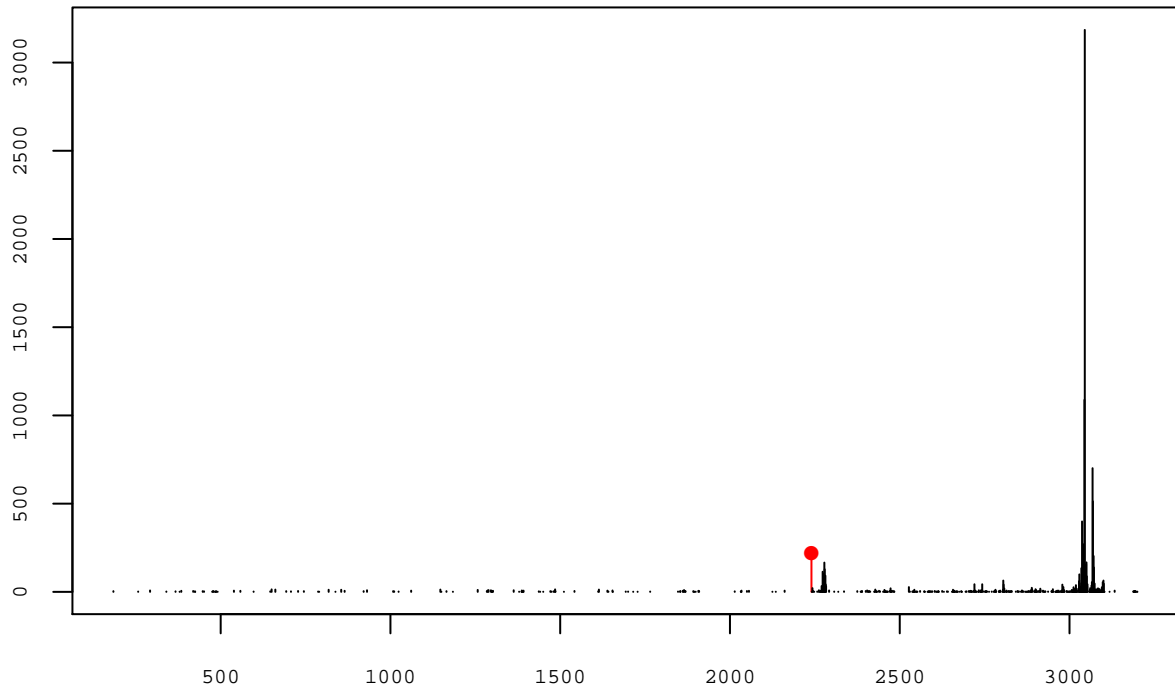


PggD11481\_c9\_g1\_i1

5' GTTGTGGCATAACAGGGACCCAGGCATAGTCA '3  
|||||  
3' CGTATGTCCTCGGTCCGT '5

Fragment Abundance



Transcript position

Cleavage site: 2240    Tag abundance: 220    Weighted abundance: 110    Category: 2  
sRNA abundance: 11    Alignment score: 2    MFE ratio: 0.835    p-value: 0.048