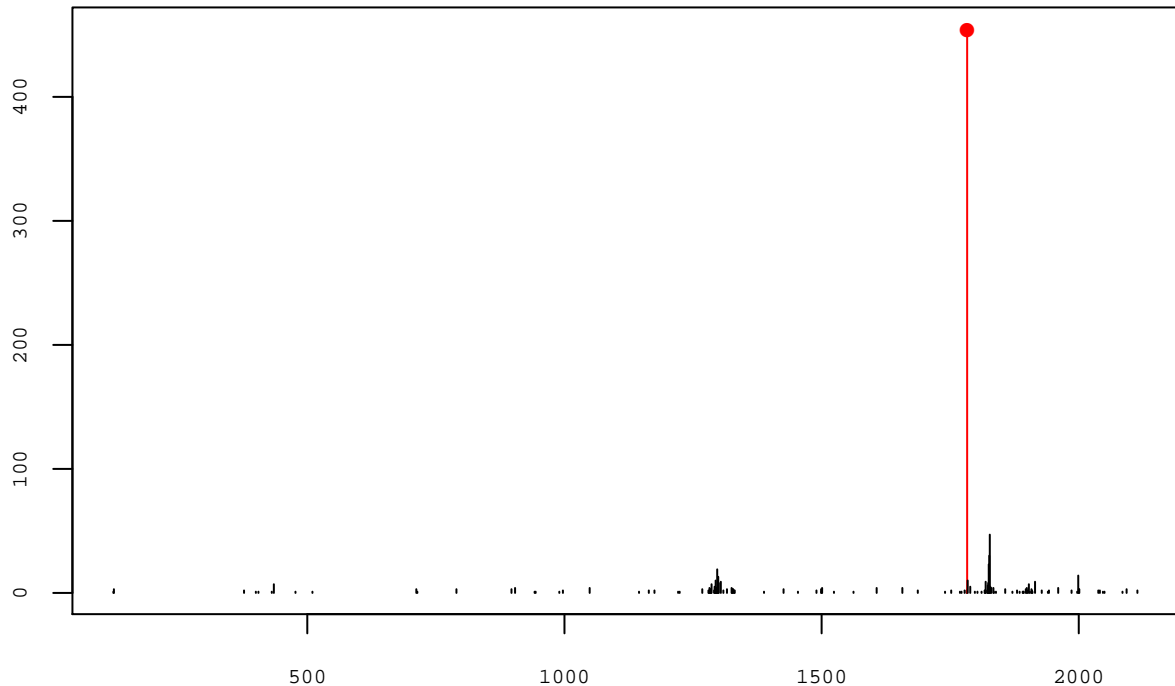


PggD620\_c8\_g1\_i1

5' CGCTGGAGCAGCATCATCAGGATTCTTTATC 3'  
○ ||||| ||||| ○ |||||  
3' TACGTCGTAGTAGTTCTAAGA 5'

Fragment Abundance



Transcript position

Cleavage site: 1783 Tag abundance: 454 Weighted abundance: 454 Category: 0  
sRNA abundance: 108 Alignment score: 2.5 MFE ratio: 0.956 p-value: 0.005