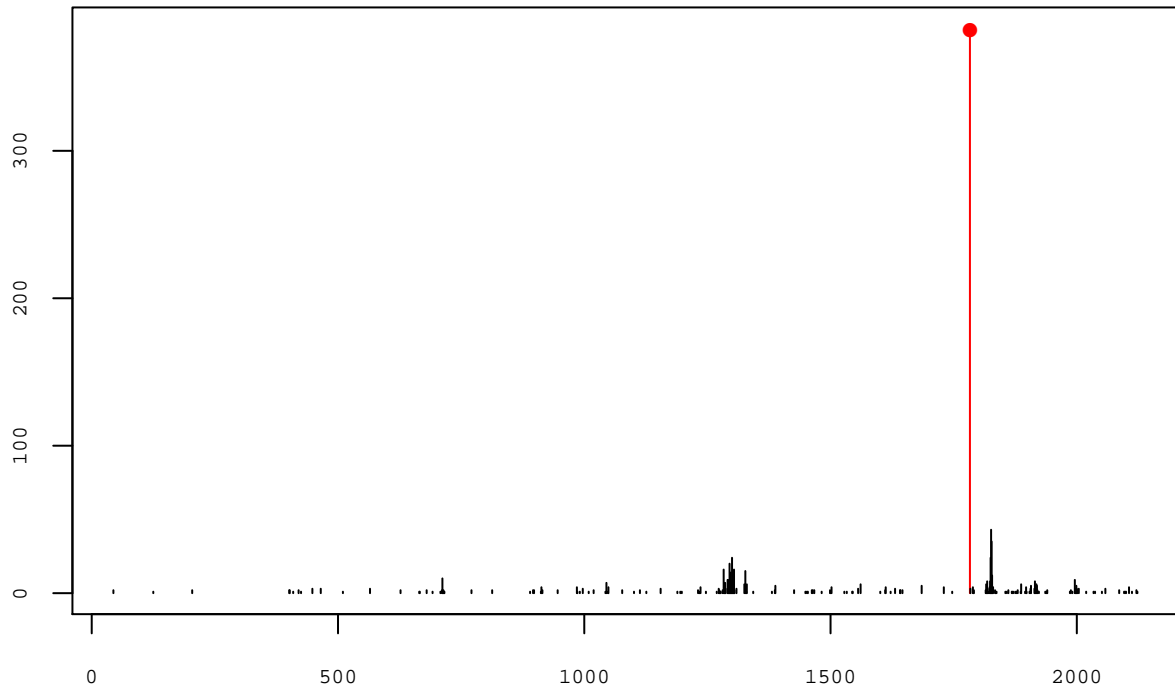


PggD620\_c8\_g1\_i1

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

Fragment Abundance



Transcript position

Cleavage site: 1783 Tag abundance: 382 Weighted abundance: 382 Category: 0  
sRNA abundance: 37 Alignment score: 2.5 MFE ratio: 0.956 p-value: 0.005