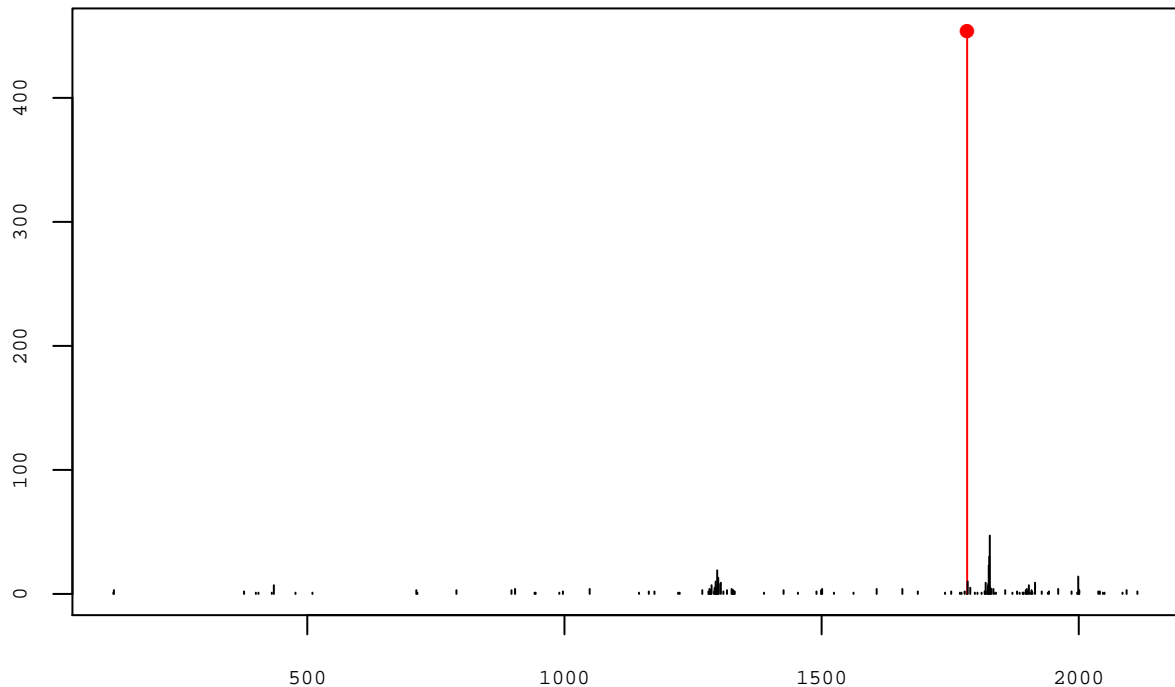


PggD620_c8_g1_i1

5' CGCTGGAGCAGCATCATCAGGATTCTCTTATC 3'
○ |||||
3' TACGTCGTAGTAGTCCTAAGG 5'

Fragment Abundance



Transcript position

Cleavage site: 1783 Tag abundance: 454 Weighted abundance: 454 Category: 0
sRNA abundance: 760 Alignment score: 2 MFE ratio: 0.924 p-value: 0.007