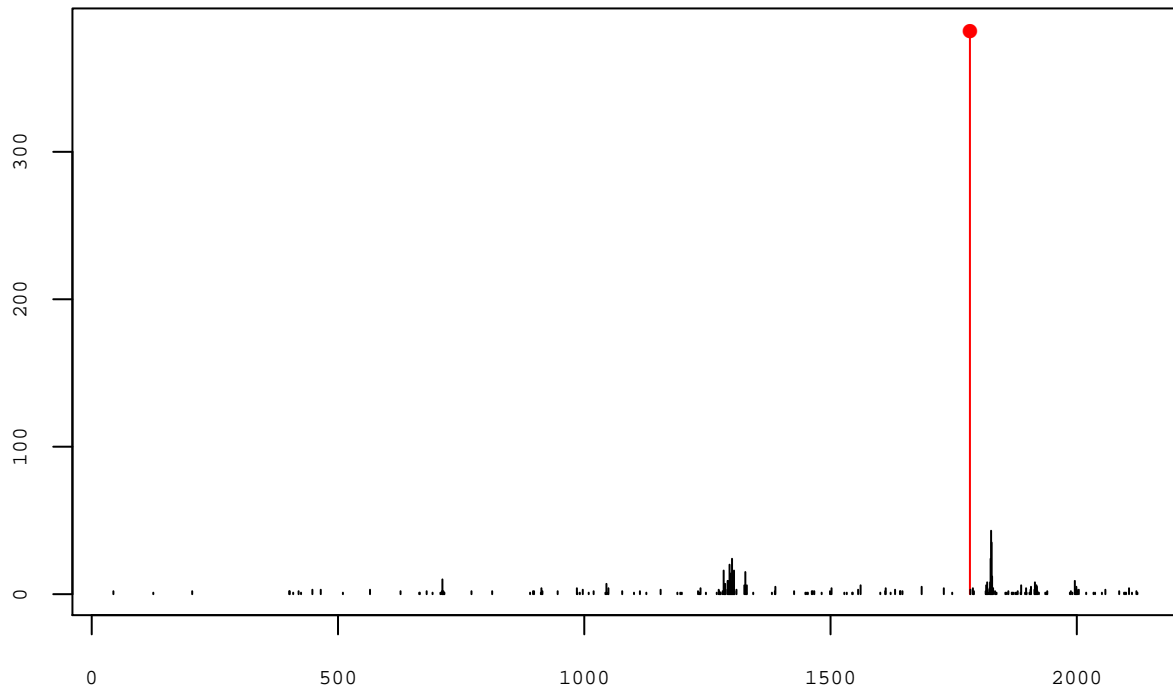


PggD620_c8_g1_i1

5' CGCTGGAGCAGCATCATCAGGATTCTTTATC '3
○ |||||
3' TACGTCGTAGTAGTCCTAAGG '5

Fragment Abundance



Transcript position

Cleavage site: 1783 Tag abundance: 382 Weighted abundance: 382 Category: 0
sRNA abundance: 48 Alignment score: 2 MFE ratio: 0.924 p-value: 0.007