

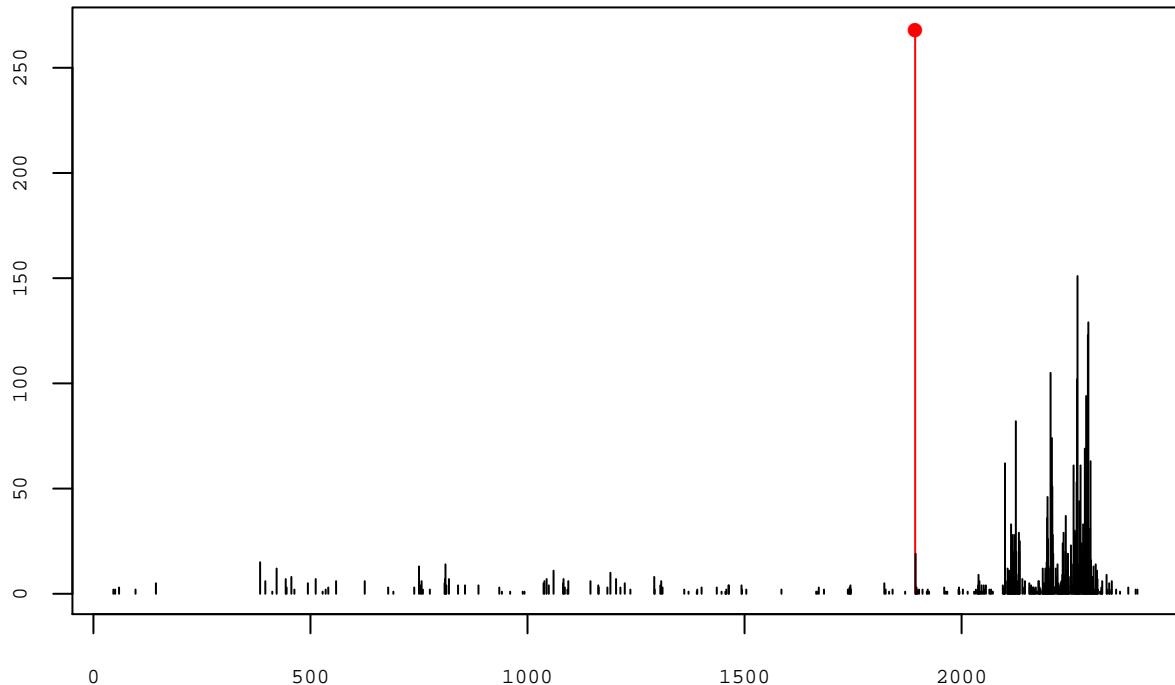
PggD9652\_c11\_g1\_i1

5' TTTAATAGGGGGA-CCCTTCAGTCCAATTCACC '3

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3' TCCCTCGAGGGAAGTCAGGTT '5

Fragment Abundance



Transcript position

Cleavage site: 1893 Tag abundance: 268 Weighted abundance: 67 Category: 0  
sRNA abundance: 8354 Alignment score: 2.5 MFE ratio: 0.794 p-value: 0.003