

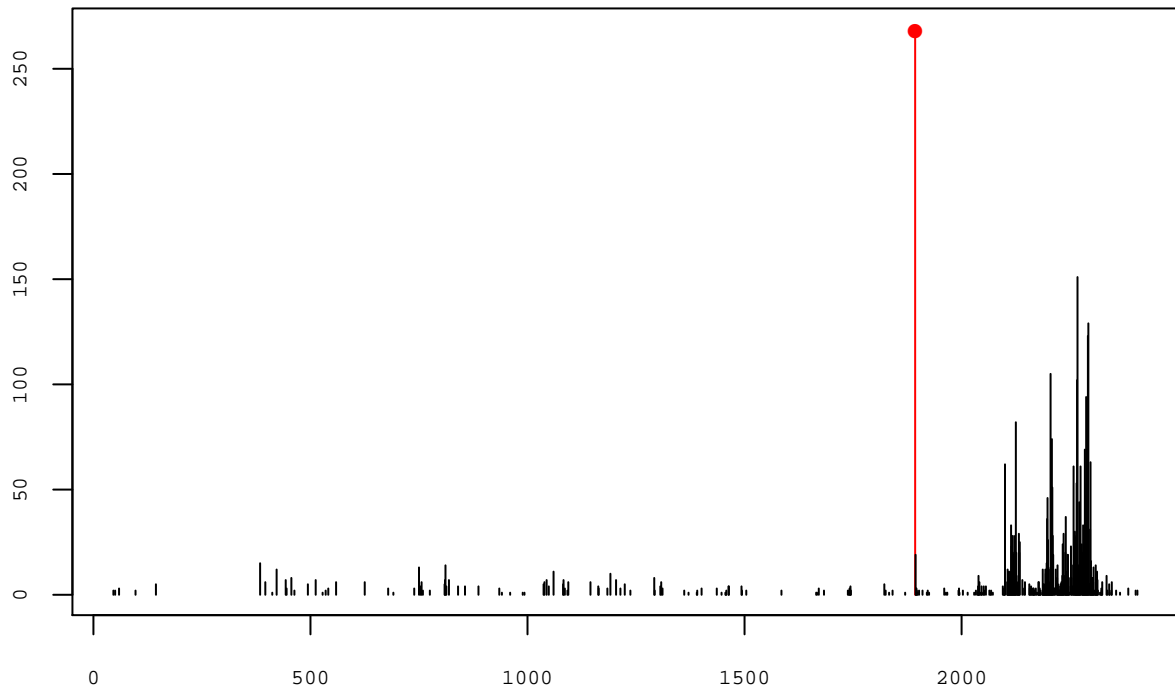
PggD9652_c11_g1_i1

5' TTTAATAGGGGGA-CCCTTCAGTCCAATTCACC '3

|||o| |||||

3' CCCTCGAGGGAAGTCAGGTT '5

Fragment Abundance



Transcript position

Cleavage site: 1893 Tag abundance: 268 Weighted abundance: 67 Category: 0
sRNA abundance: 1130 Alignment score: 2.5 MFE ratio: 0.786 p-value: 0.003