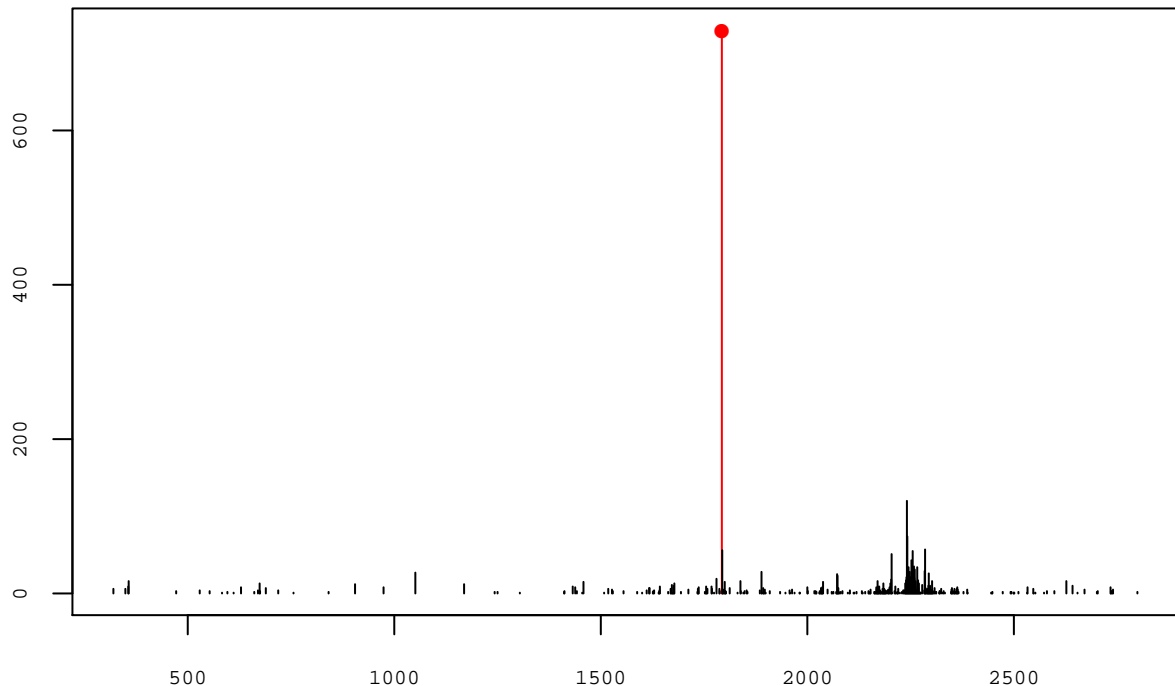


PggD2930_c0_g1_i1

5' ACACAGTGGGGAACCCTTCAGTCCAGTAACCC 3'
|| |||o| |||||o
3' TC-CCTCGAGGGAAGTCAGGTT 5'

Fragment Abundance



Transcript position

Cleavage site: 1793 Tag abundance: 729 Weighted abundance: 182.25 Category: 0
sRNA abundance: 8354 Alignment score: 4 MFE ratio: 0.704 p-value: 0.008