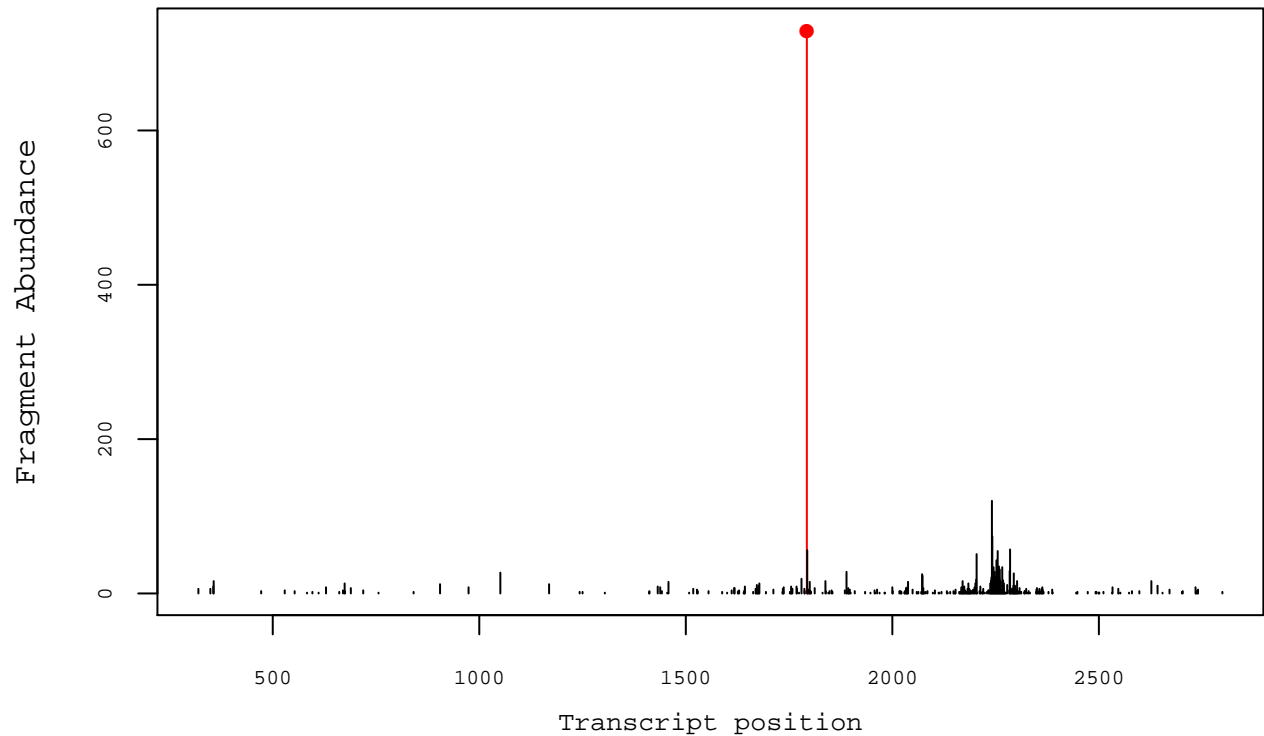


PggD2930_c0_g1_i1

5' ACACAGTGGGGAACCTTCAGTCCAGTAACCC '3

| | ○ | ||||| ||||| ||||| ○

3' C-CCTCGAGGGAAGTCAGGTT 5'



Cleavage site: 1793 Tag abundance: 729 Weighted abundance: 182.25 Category: 0
sRNA abundance: 1130 Alignment score: 4 MFE ratio: 0.726 p-value: 0.008