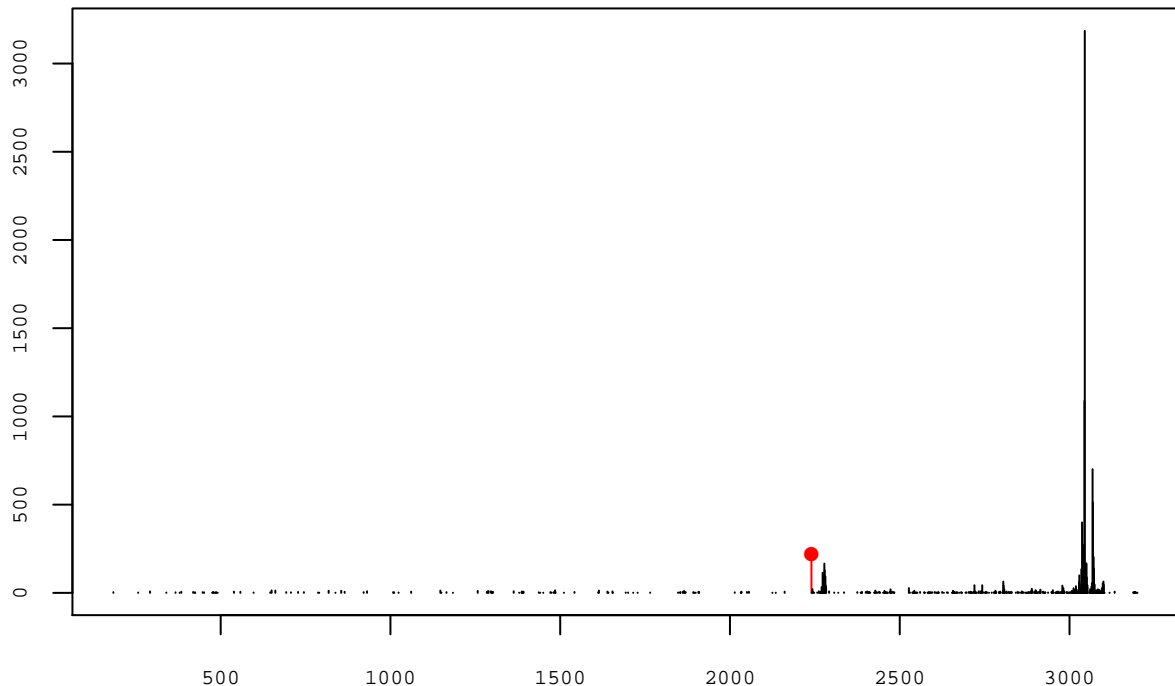


PggD11481_c9_g1_i1

5' GTTGTGGCATAACAGGGACCCAGGCATAGTCA '3
|||||
3' ACCGTATGTCCTCGGTCCGT '5

Fragment Abundance



Transcript position

Cleavage site: 2240 Tag abundance: 220 Weighted abundance: 110 Category: 2
sRNA abundance: 102 Alignment score: 2 MFE ratio: 0.852 p-value: 0.048