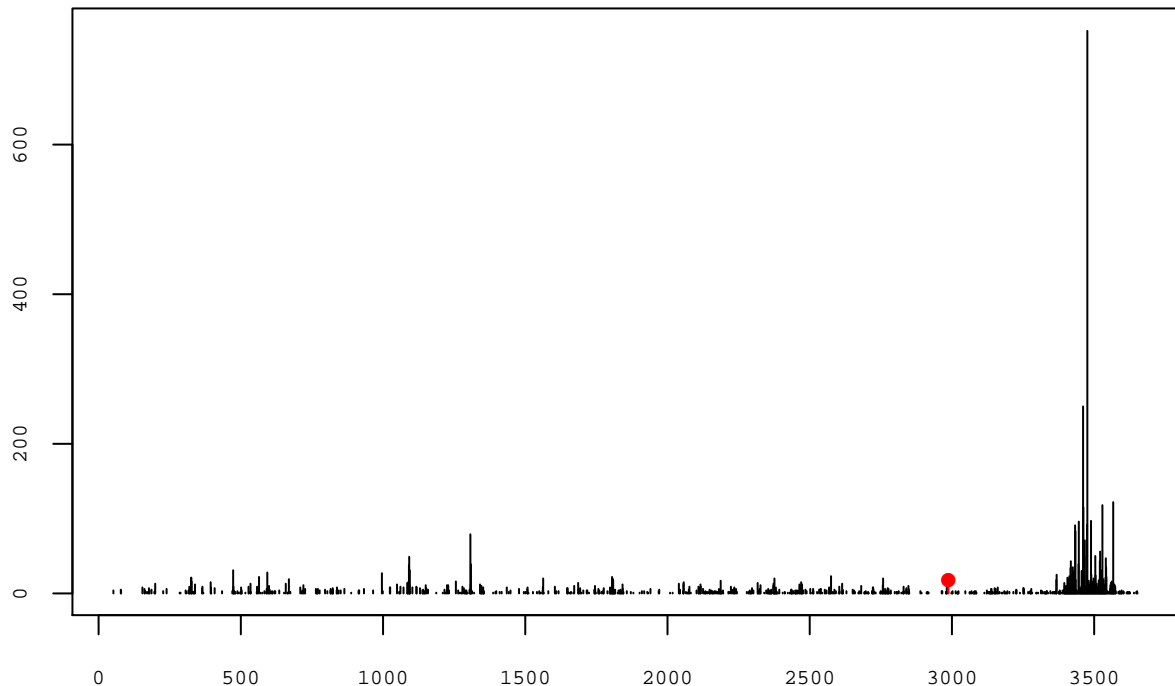


PggD11872\_c0\_g1\_i2

5' CCTCTTAGATCAGGCTGGCAGCTT-GTATTGT '3  
|o||||| |||||o  
3' AGTCTAGTACGACCGTCGAAGT '5

Fragment Abundance



Transcript position

Cleavage site: 2988 Tag abundance: 18 Weighted abundance: 18 Category: 2  
sRNA abundance: 442 Alignment score: 3 MFE ratio: 0.753 p-value: 0.035