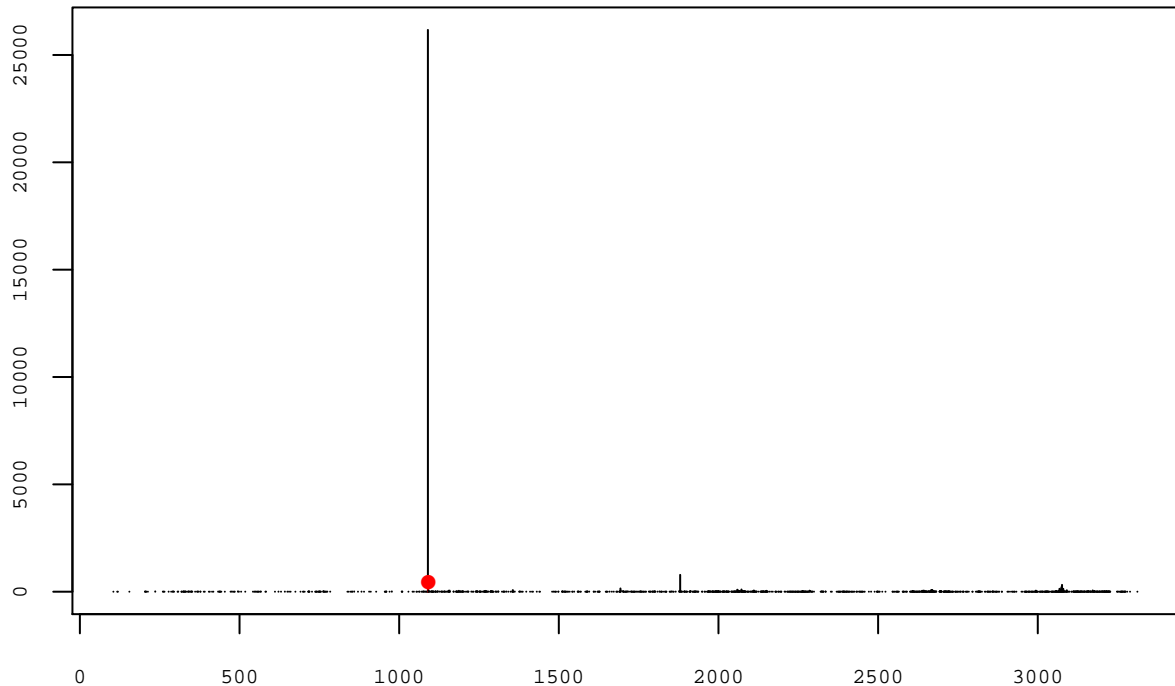


PggD7141\_c1\_g1\_i1

5' ATGGTTGGTATGAAGCCTGGTCCG-GATTCTAT '3

3' CCCTTACTTCGGACCAGGCTC '5

Fragment Abundance



Transcript position

Cleavage site: 1091 Tag abundance: 456 Weighted abundance: 228 Category: 2  
sRNA abundance: 371 Alignment score: 3 MFE ratio: 0.747 p-value: 0.038