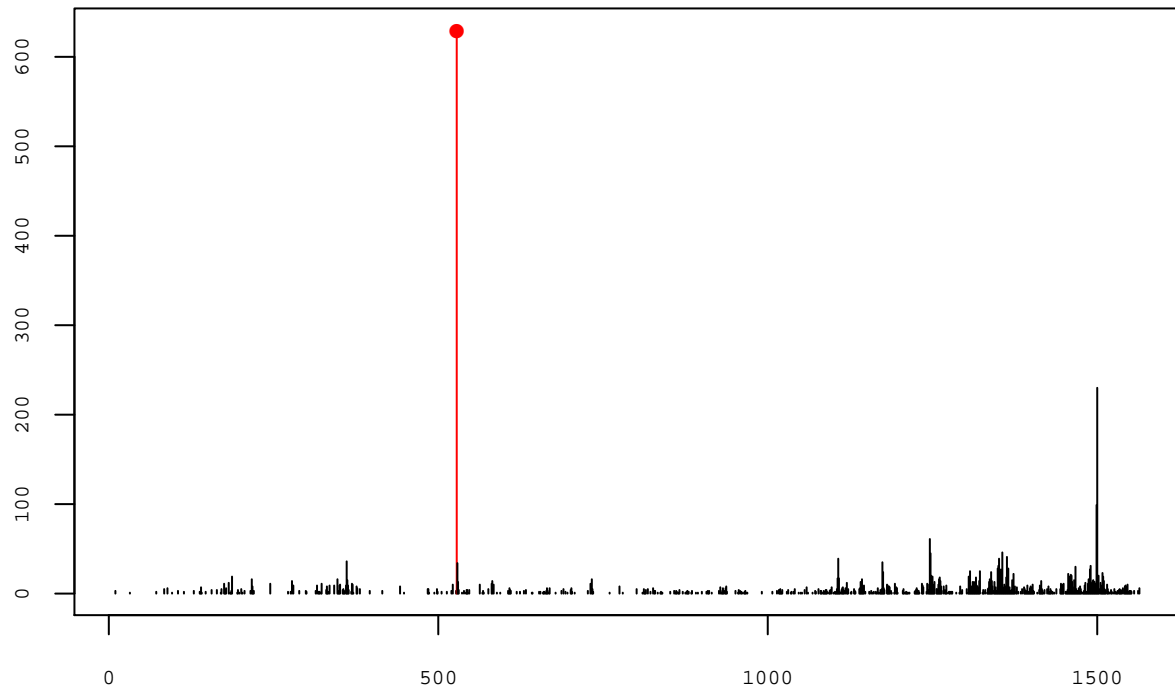


PggD12227\_c16\_g1\_i1

5' GGAGAAAAAGCTCCATTCAATCCAAAGCGCGC '3  
| | | | | | | | | | | | | | | | | |  
3' TCTCGAGGGAAGTTAGGTTT '5

Fragment Abundance



Transcript position

Cleavage site: 528 Tag abundance: 629 Weighted abundance: 629 Category: 0  
sRNA abundance: 26628 Alignment score: 3 MFE ratio: 0.705 p-value: 0.008