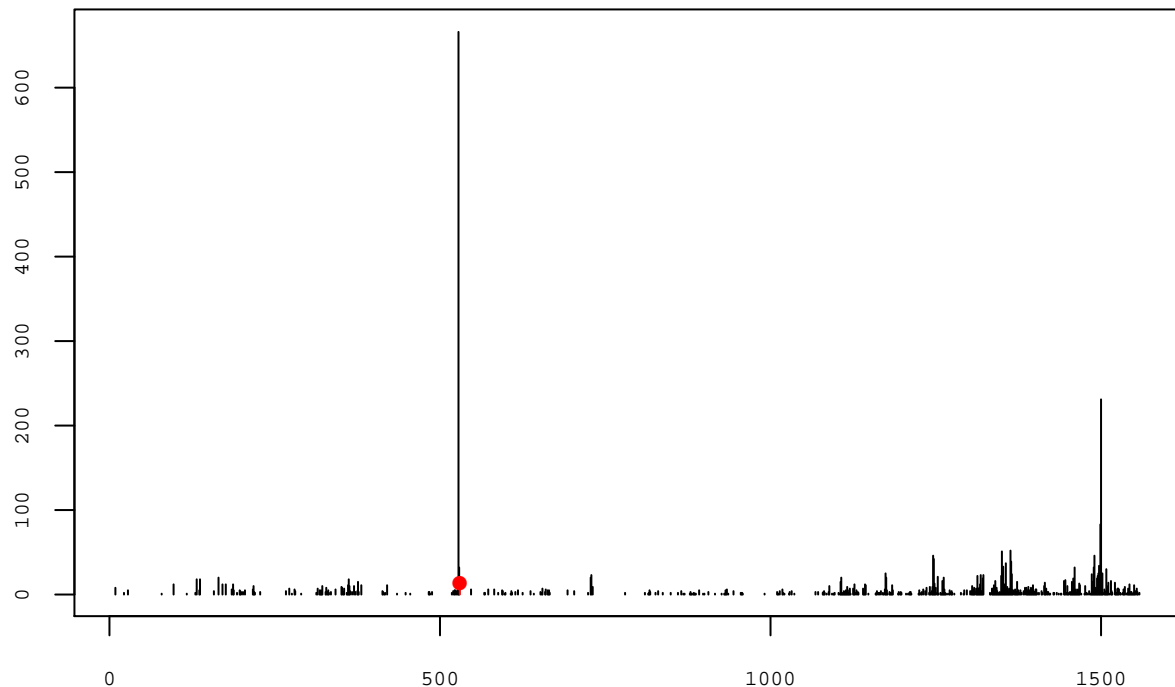


PggD12227\_c16\_g1\_i1

5' AGAAAAAGCTCCATTCAATCCAAAGCGCGCTT '3  
||||| |||||  
3' CGAGGGAAGTTAGGTTTCT '5

Fragment Abundance



Transcript position

Cleavage site: 530 Tag abundance: 14 Weighted abundance: 14 Category: 2  
sRNA abundance: 5 Alignment score: 2 MFE ratio: 0.761 p-value: 0.048