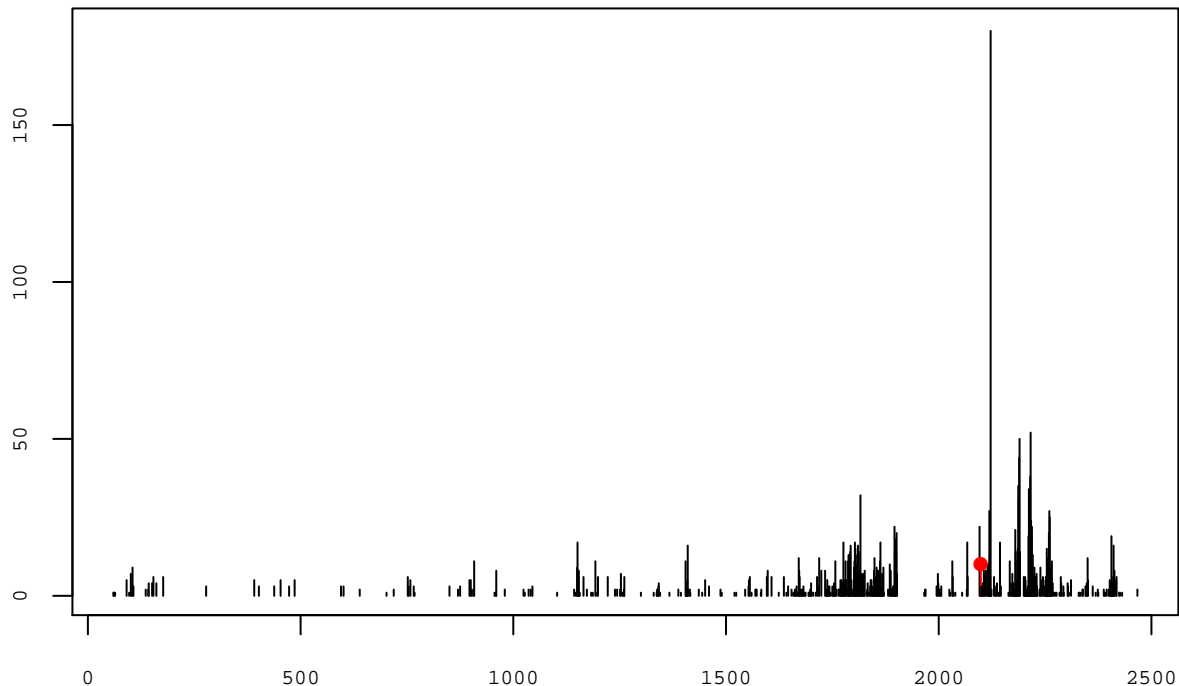


PggD7047\_c0\_g1\_i1

5' GAGGTTACAGGGGGTATAACACGGATATAACT '3  
|o|||||o|||||||o|||  
3' CTAATGTCTCCCATATTGTGTCTA '5

Fragment Abundance



Transcript position

Cleavage site: 2099 Tag abundance: 10 Weighted abundance: 10 Category: 2  
sRNA abundance: 31 Alignment score: 2 MFE ratio: 0.944 p-value: 0.036