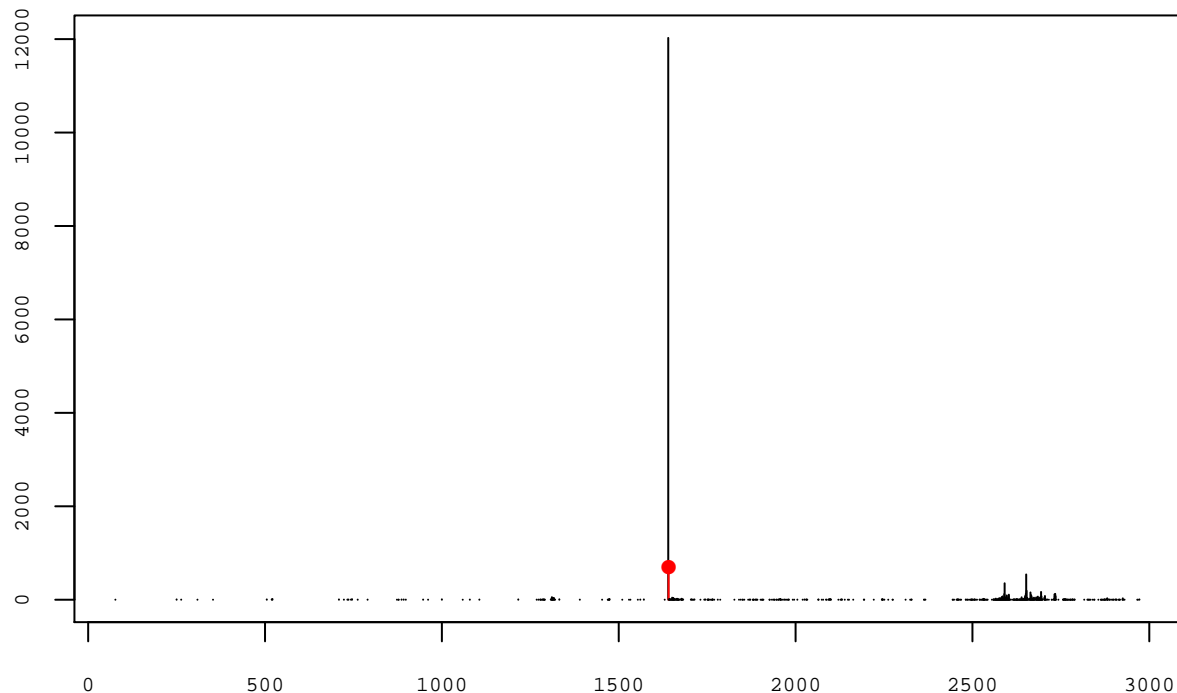


PggD1242\_c13\_g1\_i2

5' CAGCTAGCATAACAGGGAGCCAGGCAAATTCTA '3  
|||||  
3' CCGTATGTCCCTCGGTCCGTA '5

Fragment Abundance



Transcript position

Cleavage site: 1641	Tag abundance: 691	Weighted abundance: 345.5	Category: 2
sRNA abundance: 22	Alignment score: 2	MFE ratio: 0.942	p-value: 0.025