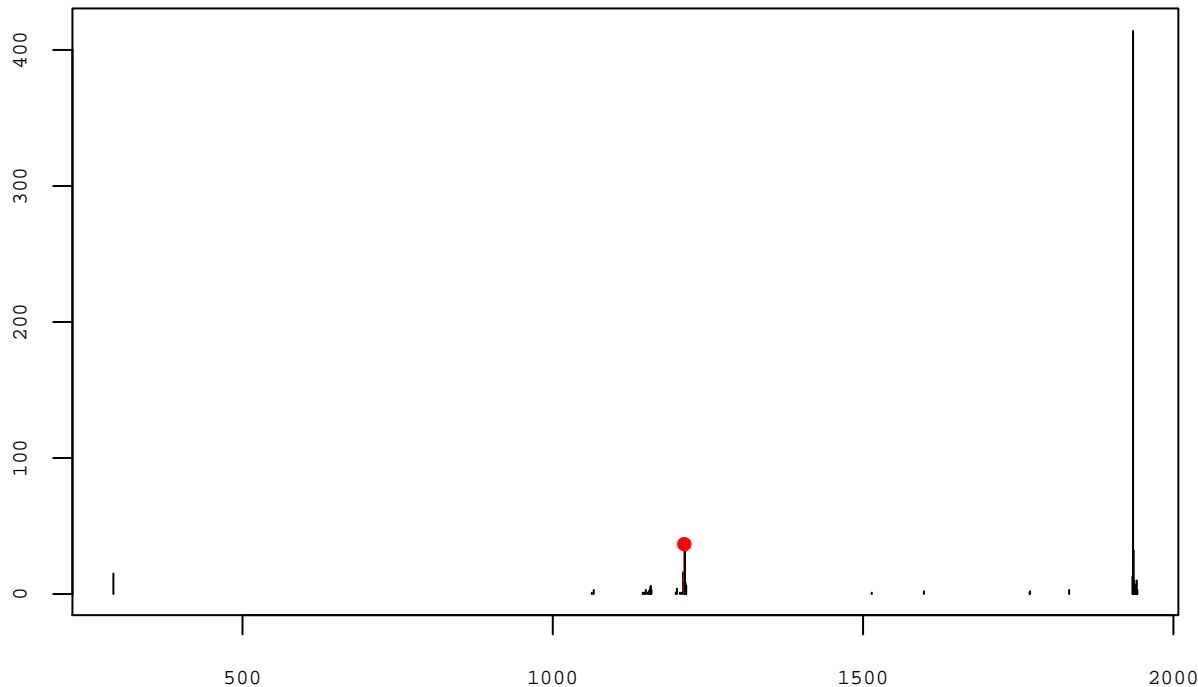


PggD15095\_c245\_g1\_i1

5' CACGTAGAGCTACGTCAGGAGTCCGGACAGAA '3  
|||||  
3' ACATCTCGATGCAGTCCTCAGGCG '5

Fragment Abundance



Transcript position

Cleavage site: 1212	Tag abundance: 37	Weighted abundance: 18.333	Category: 2
sRNA abundance: 27	Alignment score: 2	MFE ratio: 0.957	p-value: 0.003