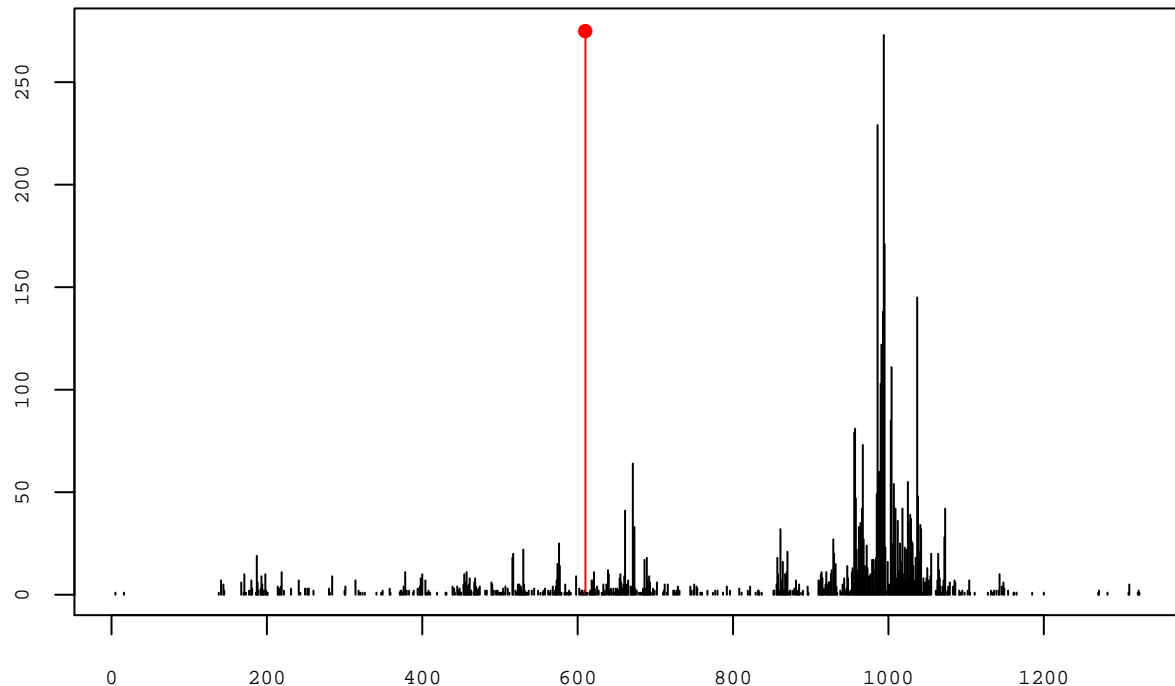


PggD13945\_c35\_g1\_i1

5' AAGGAACTGCGCGGTCTTGAGCAAAACATTGA '3  
|||||  
3' CAACGCGCCAGAACTCGTTTT '5

Fragment Abundance



Transcript position

Cleavage site: 610	Tag abundance: 275	Weighted abundance: 68.75	Category: 0
sRNA abundance: 6	Alignment score: 2	MFE ratio: 0.932	p-value: 0.007