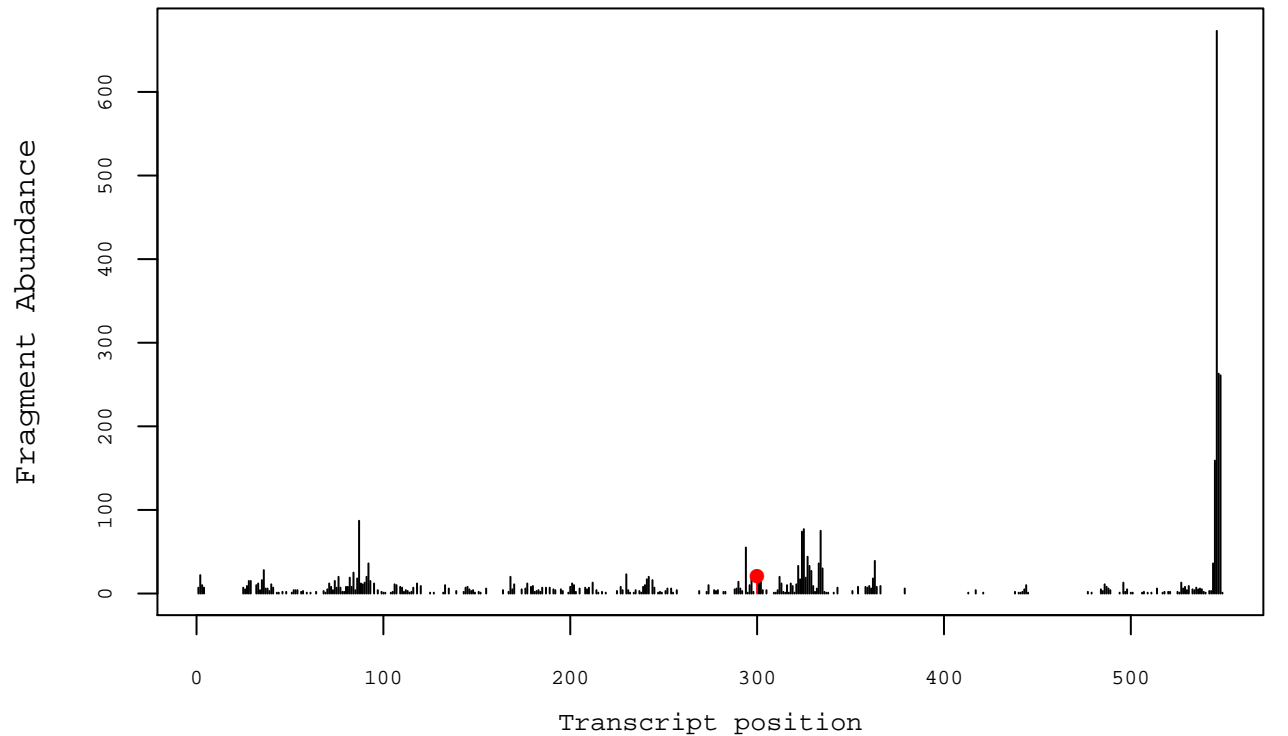


PggD9068_c6_g1_i1

5' GCGGTGTCCGATCCTGACCGTAGATCATGATT '3
 |||||
 3' CCACAGGCTAGGTCTGGCATTTAG '5



Cleavage site: 300	Tag abundance: 21	Weighted abundance: 21	Category: 2
sRNA abundance: 514	Alignment score: 3	MFE ratio: 0.903	p-value: 0.042