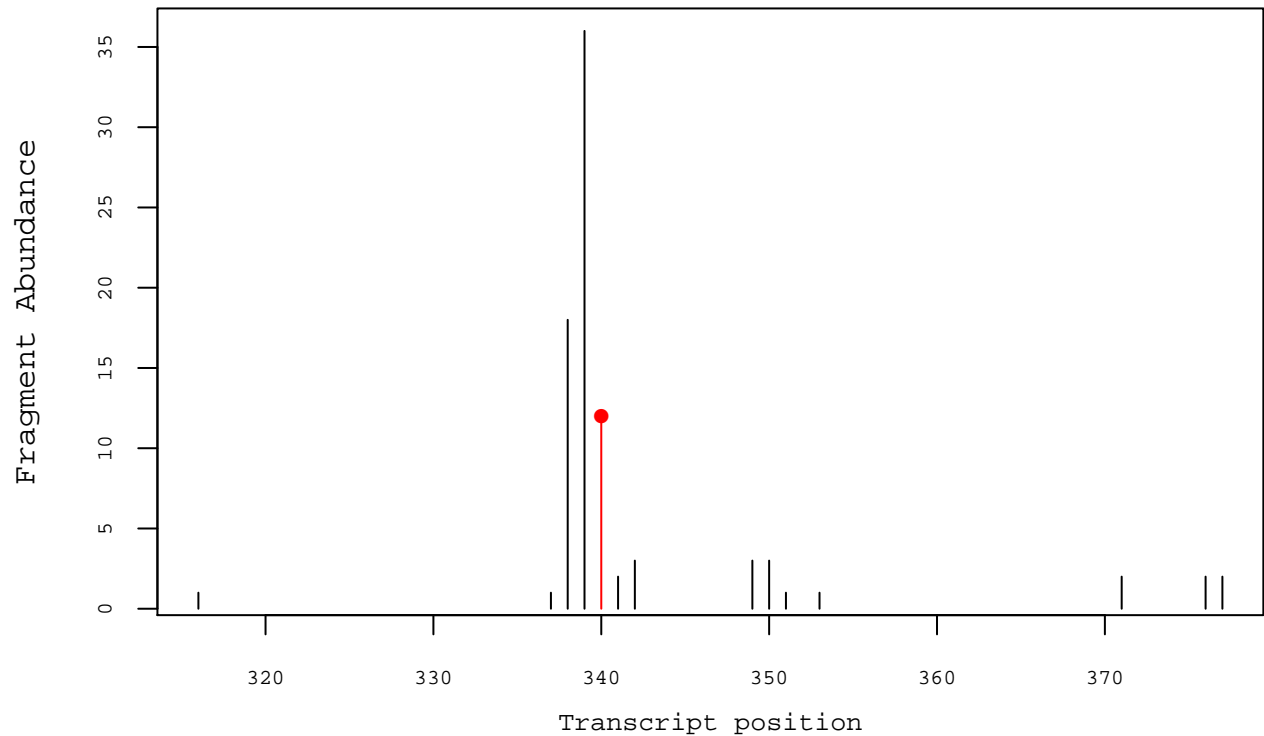


PggD8965_c122_g1_i1

5' CAGCTCGTGTTCGCTCGTTAAGAAGCTCGTG '3
||||||| | |||||
3' CGAGCACAAGCCAAGCAATTCTTC '5



Cleavage site: 340	Tag abundance: 12	Weighted abundance: 0.983	Category: 2
sRNA abundance: 9	Alignment score: 3	MFE ratio: 0.752	p-value: 0.047