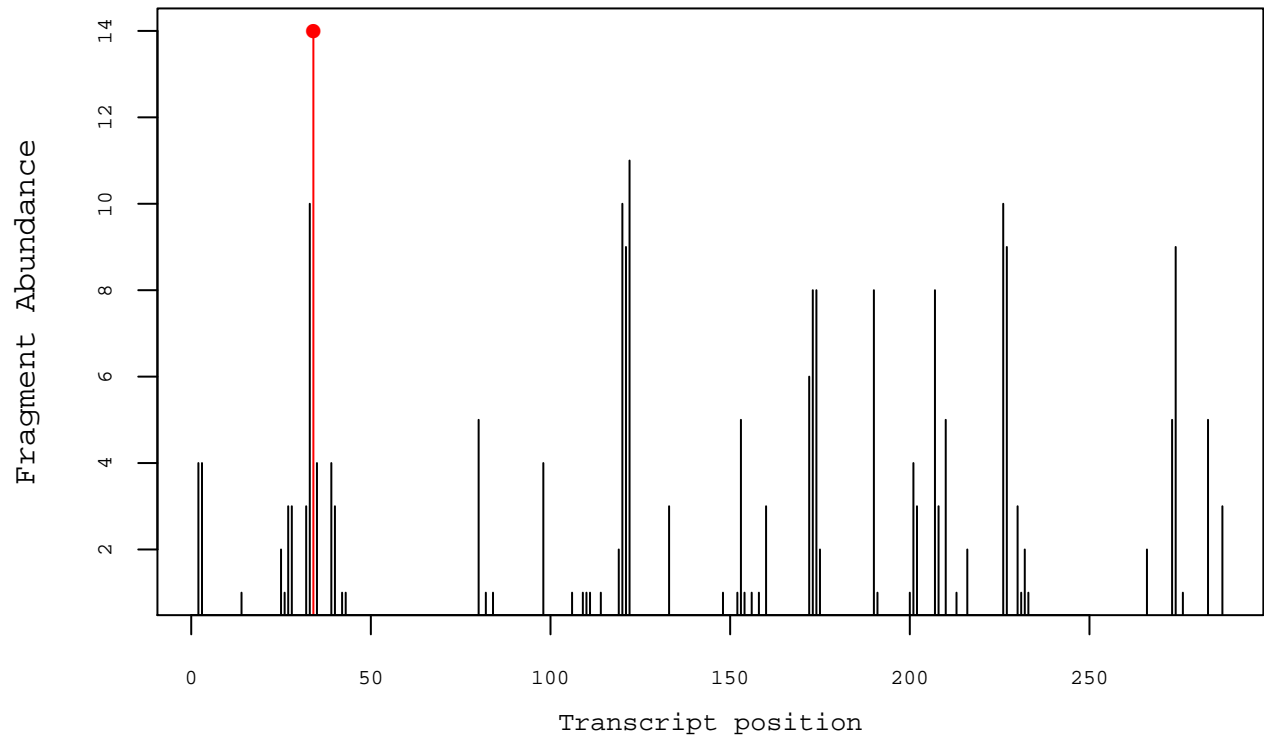


5' TGTCTGATTGTTTGGTTGGCGCTTATGGA AAA 3'
 ||| o ||||| ||| o |||||
 3' AGACCGACAAACCAACTGCGAATA 5'



Cleavage site: 34 Tag abundance: 14 Weighted abundance: 1.75 Category: 0
 sRNA abundance: 22 Alignment score: 2.5 MFE ratio: 0.766 p-value: 0.031