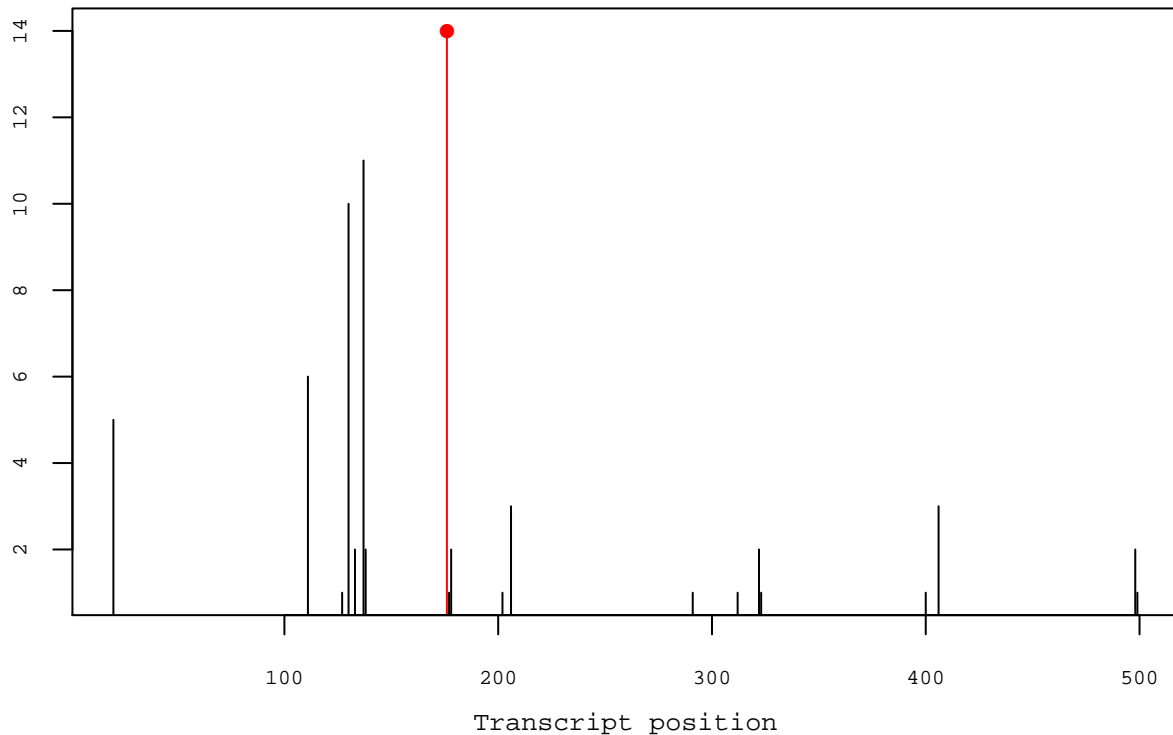


5' TCATCACTGCATGCGGTATATGTATGCAATAA '3  
||| |||||  
3' TAGAGACGTACGCCATATACATAC '5

Fragment Abundance



Cleavage site: 176

sRNA abundance: 21

Tag abundance: 14

Alignment score: 1

Weighted abundance: 4.667

MFE ratio: 0.869

Category: 0

p-value: 0.002