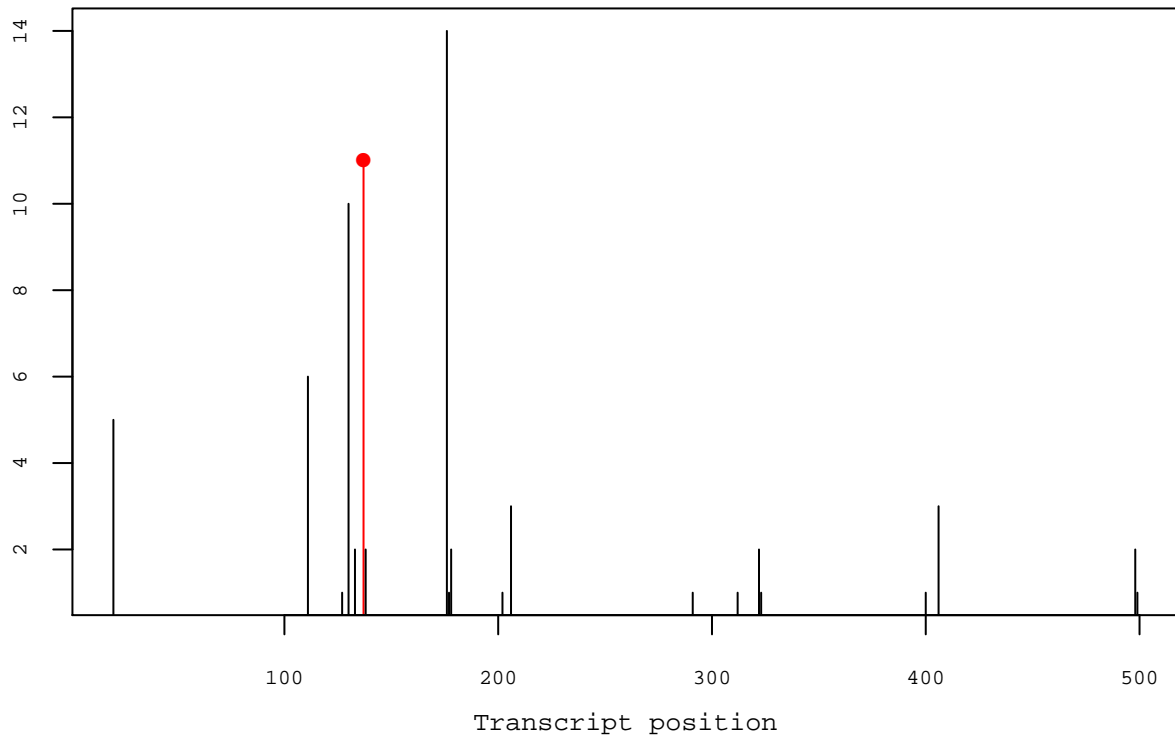


```

5' TAGATCACCGTGATCTACGGTCAGGATCGGAC '3
   ||||| ||||| ||||| ||||| ||||| |||||
3' CTAGTGCCACTAGATGCCAGTCTT '5

```

Fragment Abundance



Cleavage site: 137 Tag abundance: 11 Weighted abundance: 5 Category: 2  
sRNA abundance: 9 Alignment score: 2 MFE ratio: 0.849 p-value: 0.015