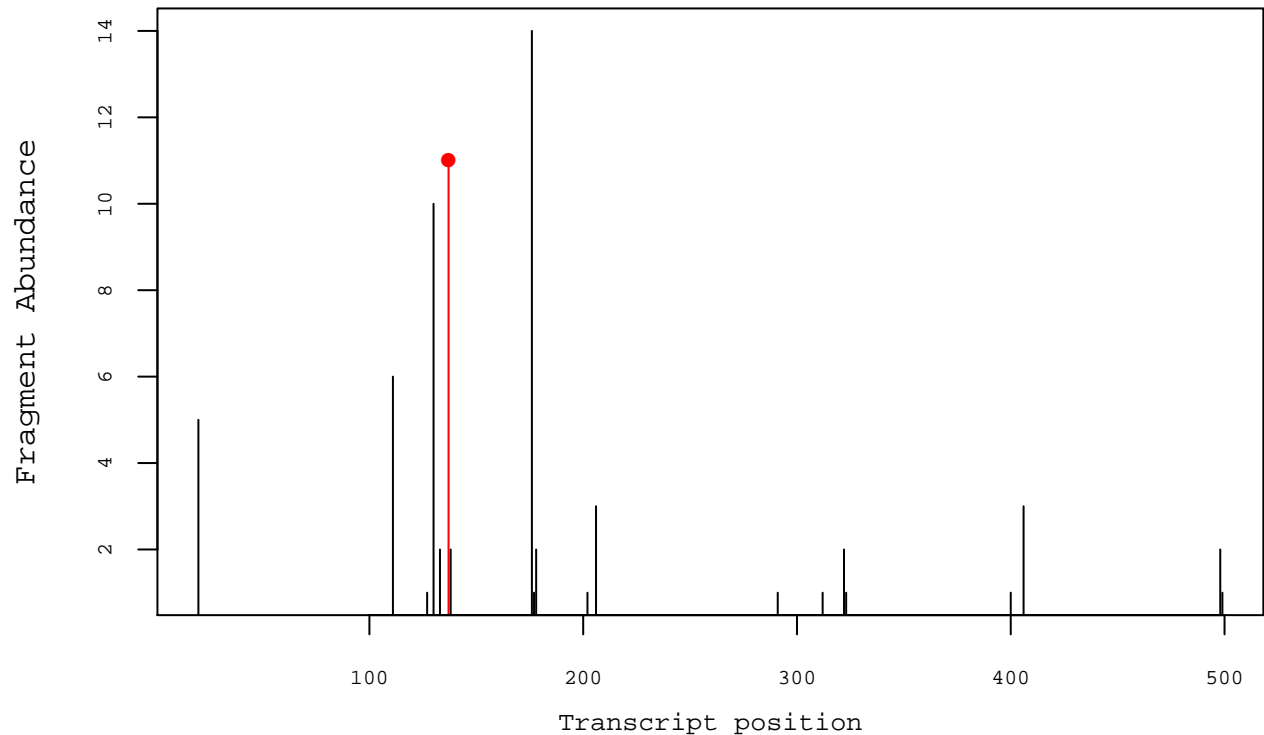


PggD2931\_c59\_g1\_i1

5' TAGATCACCGTGATCTACGGTCAGGATCGGAC '3  
 ||||| |||||  
 3' CTAGTGCCACTAGATGCCAGTACT '5



Cleavage site: 137    Tag abundance: 11    Weighted abundance: 5    Category: 2  
sRNA abundance: 9    Alignment score: 3    MFE ratio: 0.765    p-value: 0.015