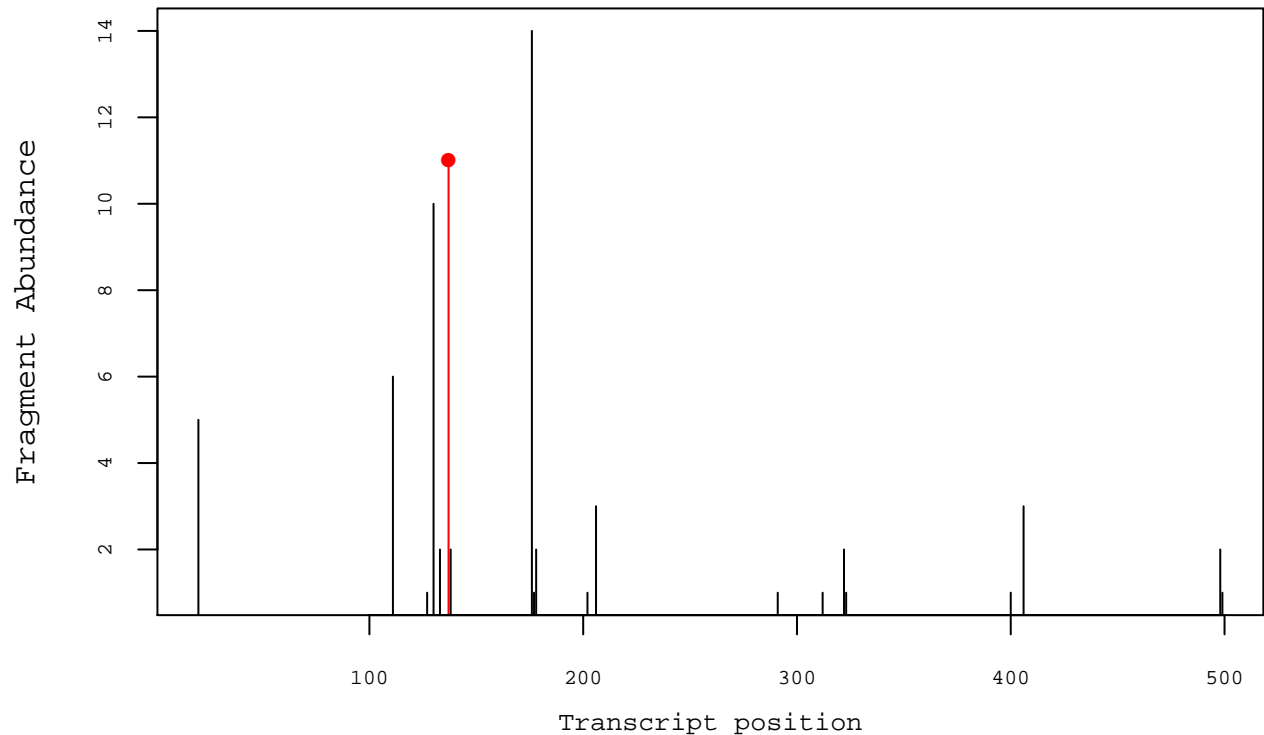


PggD2931\_c59\_g1\_i1

5' TAGATCACCGTGATCTACGGTCAGGATCGGAC 3'  
||||| |||||||||  
3' CTAGTGTCAGTATGATGCCAGTCCT 5'



Cleavage site: 137    Tag abundance: 11    Weighted abundance: 5    Category: 2  
sRNA abundance: 7    Alignment score: 1    MFE ratio: 0.897    p-value: 0.015