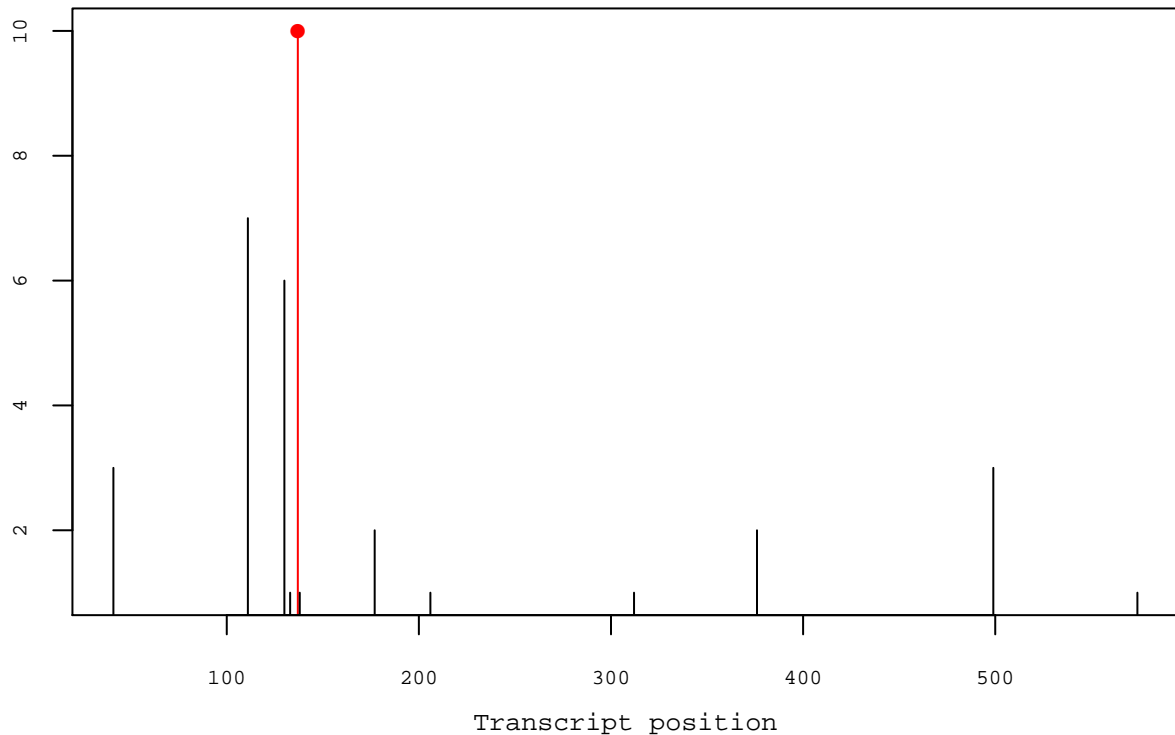


PggD2931\_c59\_g1\_i1

5' TAGATCACCGTGATCTACGGTCAGGATCGGAC '3  
||| ||| o ||| ||| ||| ||| o |  
3' CTAGTGCCATTAGATGCCAGTCTT '5

Fragment Abundance



Cleavage site: 137 Tag abundance: 10 Weighted abundance: 4.833 Category: 0  
sRNA abundance: 11 Alignment score: 2.5 MFE ratio: 0.843 p-value: 0.003