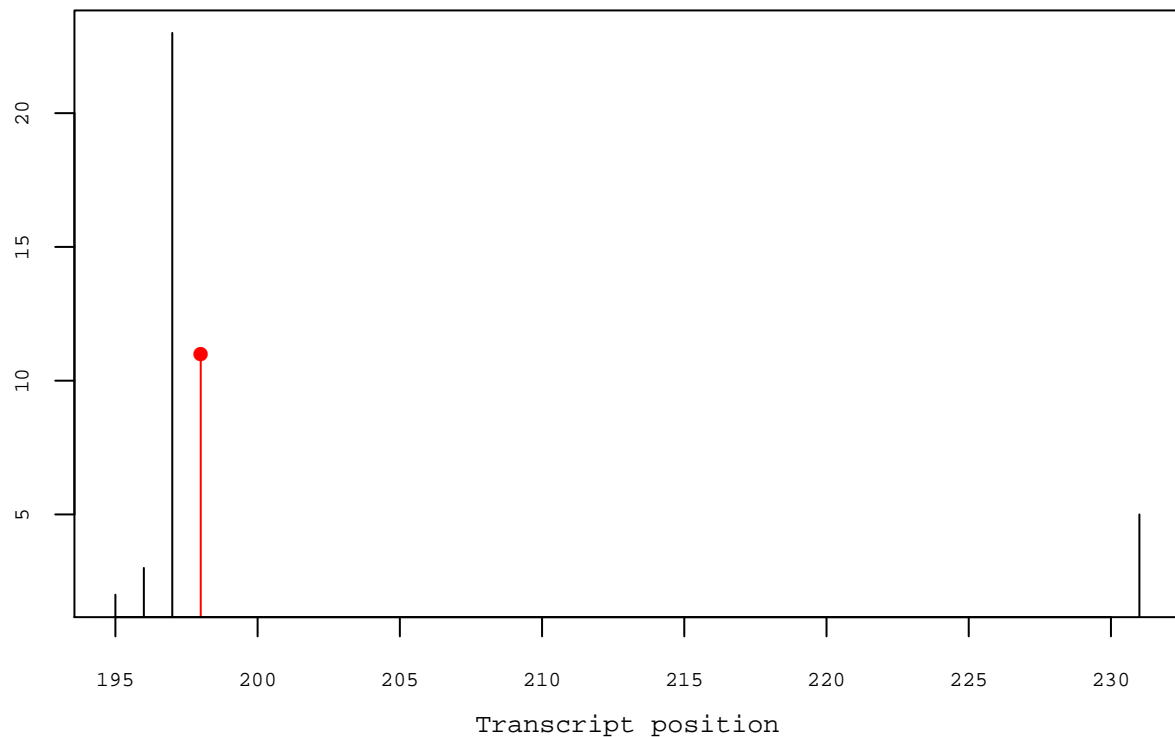


PggD7237\_c562\_g1\_i1

5' TGTACCCTCCAAATAACCTAATAT-ATATGCAA '3  
||||| |||||  
3' ATGGGAGGCTTATTGGATTATACA '5

Fragment Abundance



Cleavage site: 198 Tag abundance: 11 Weighted abundance: 5.5 Category: 2  
sRNA abundance: 21 Alignment score: 3 MFE ratio: 0.754 p-value: 0.005