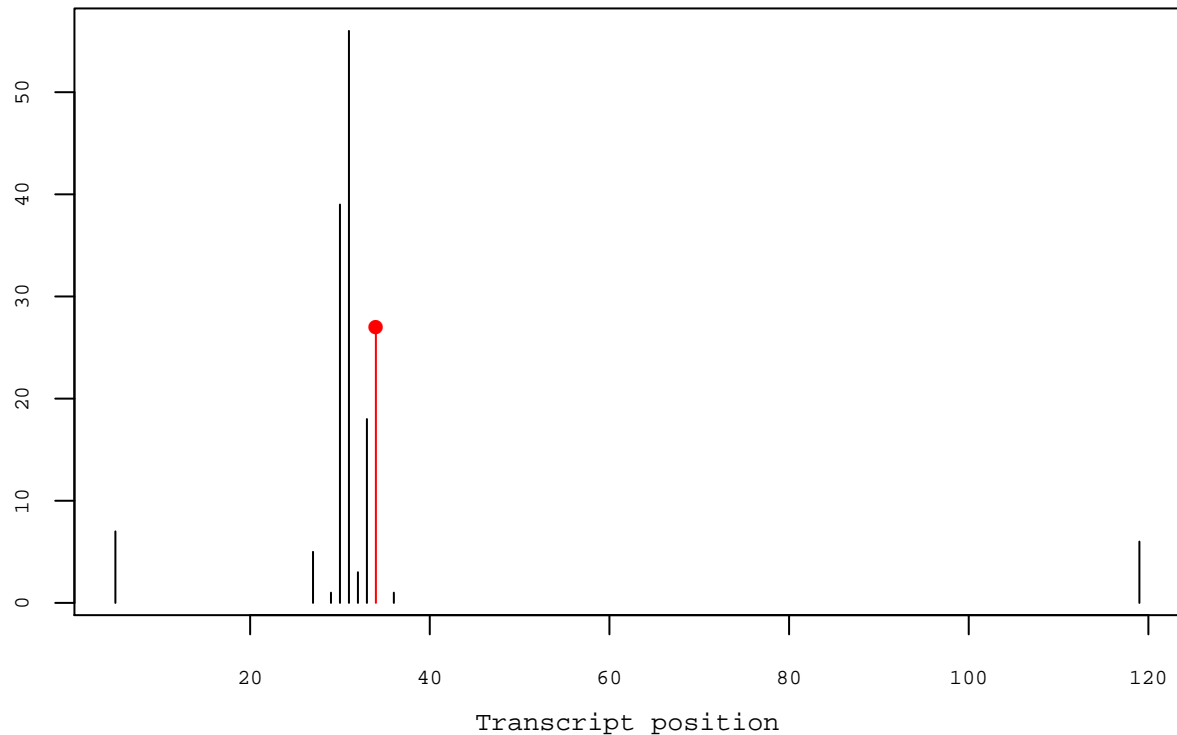


PggD5575\_c505\_g1\_i1

5' TGTACCCTCCGAATAACCTAAAAATGTACGCAA '3  
||||| ||||| |||||  
3' ATGGGAGGATTATTGGATTATACA '5

Fragment Abundance



Cleavage site: 34 Tag abundance: 27 Weighted abundance: 13.5 Category: 2  
sRNA abundance: 10 Alignment score: 3 MFE ratio: 0.766 p-value: 0.022