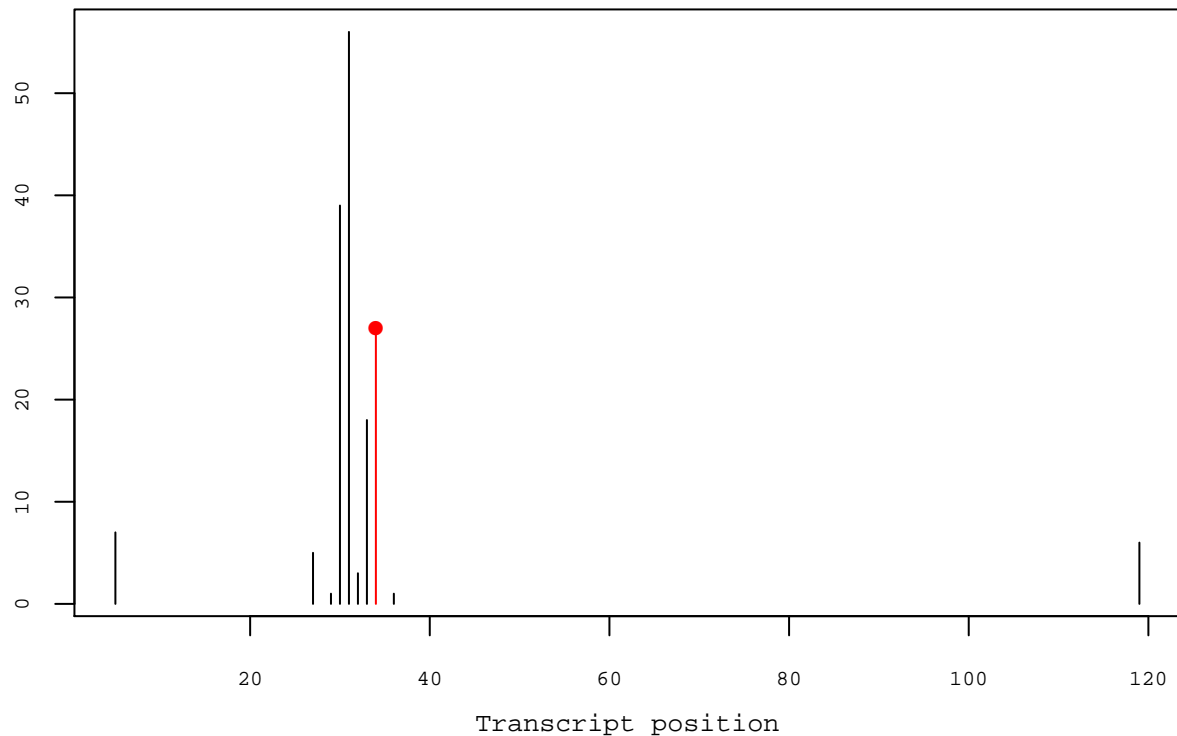


PggD5575_c505_g1_i1

5' TGTACCCTCCGAATAACCTAAAAATGTACGCAA '3
|||||
3' ATGGGAGGCTTATTGGATTATACA '5

Fragment Abundance



Cleavage site: 34 Tag abundance: 27 Weighted abundance: 13.5 Category: 2
sRNA abundance: 21 Alignment score: 2 MFE ratio: 0.891 p-value: 0.007