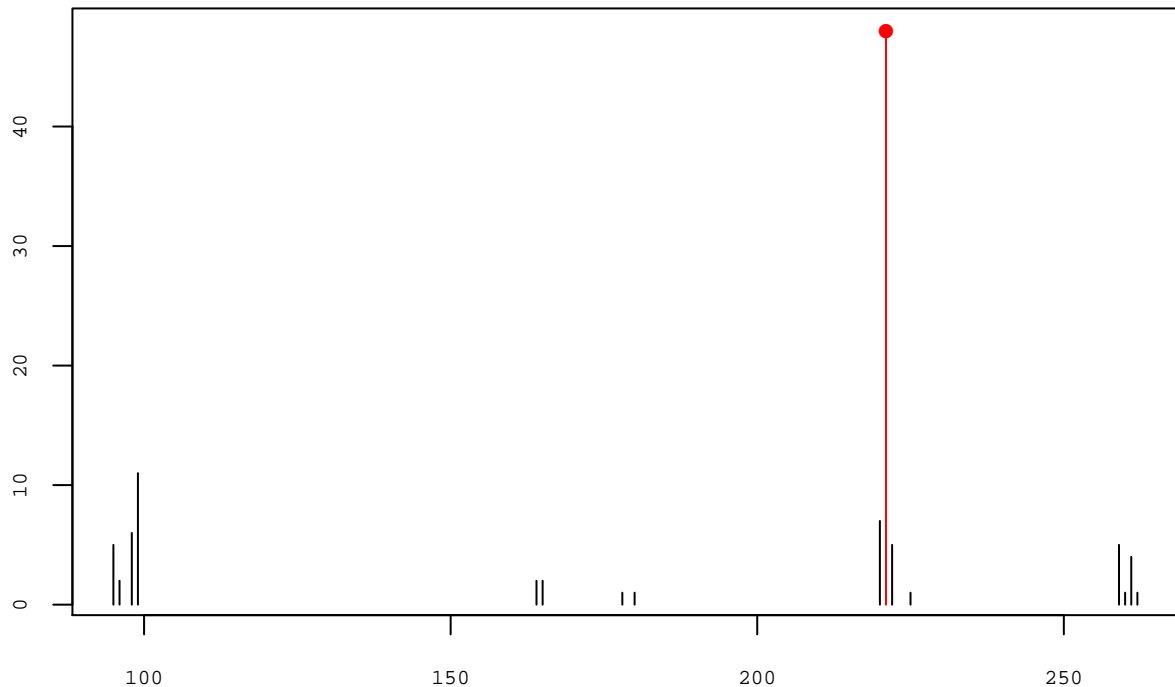


PggD14093_c110_g1_i1

5' AATTTGGGTGTTAAATATGTAAATACACAAG '3
|||oo||||||||||||||||||
3' AAAGTCCACAATTATACATTAT '5

Fragment Abundance



Transcript position

Cleavage site: 221 Tag abundance: 48 Weighted abundance: 1.505 Category: 0
sRNA abundance: 5 Alignment score: 1 MFE ratio: 0.885 p-value: 0.016