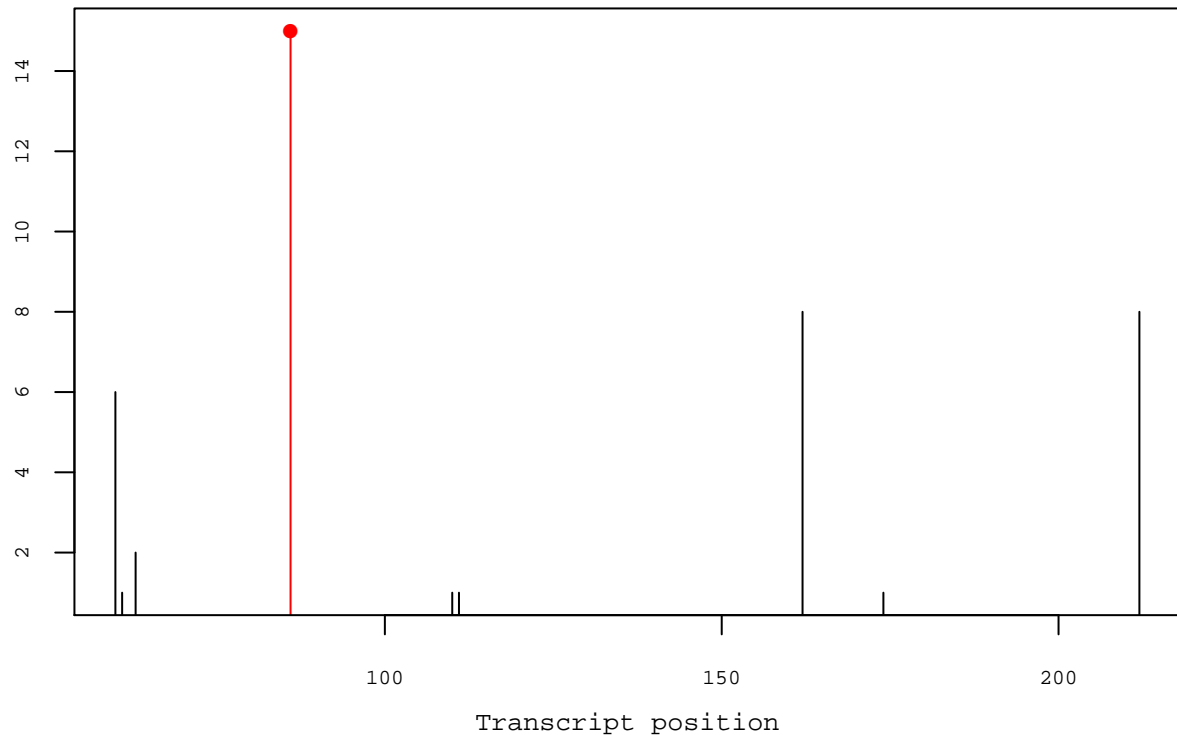


5' CAAGCAGATGCCCCGACCAACTGAT-CTGAGTAT '3
|||||
3' TCGTCTACGGGCTGGTTGACTATA '5

Fragment Abundance



Cleavage site: 86 Tag abundance: 15 Weighted abundance: 15 Category: 0
sRNA abundance: 7 Alignment score: 2 MFE ratio: 0.959 p-value: 0.013