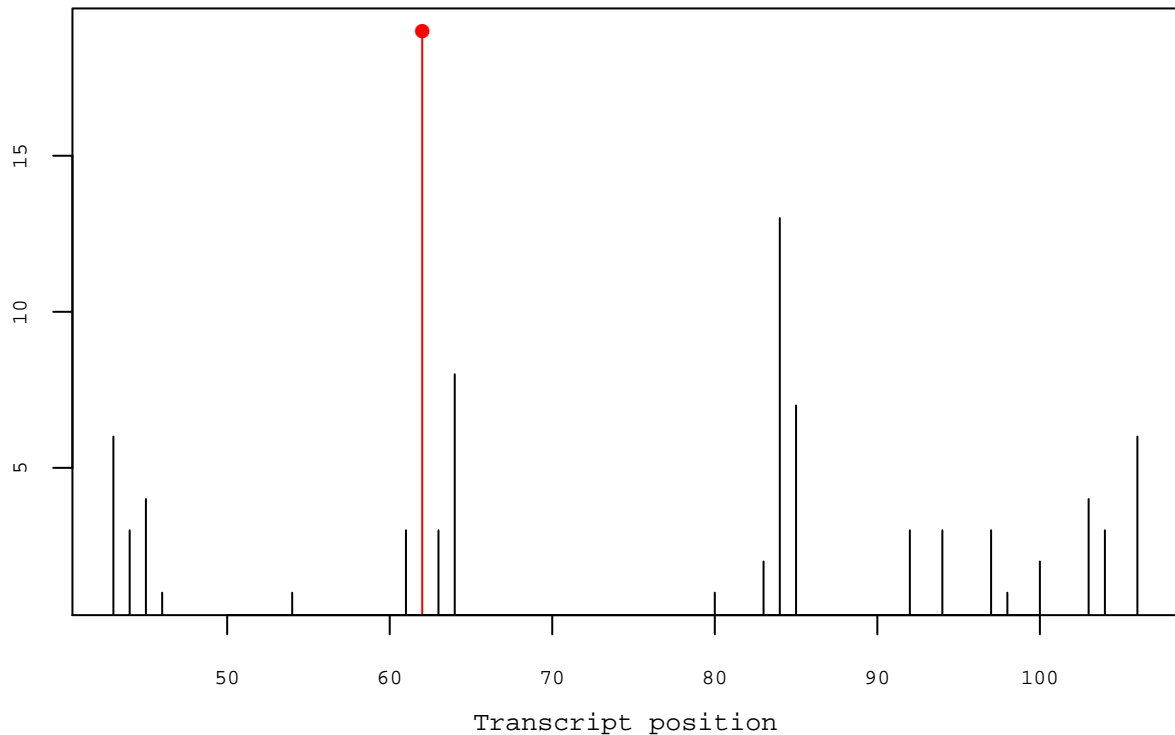


5' CTTTGTTCGTGAACAGCTCGTGTTCGGCTCGT '3
|||||
3' ACAAGCACTTGTGAGCACAAAC '5

Fragment Abundance



Cleavage site: 62 Tag abundance: 19 Weighted abundance: 0.589 Category: 0
sRNA abundance: 13 Alignment score: 2 MFE ratio: 0.93 p-value: 0.045