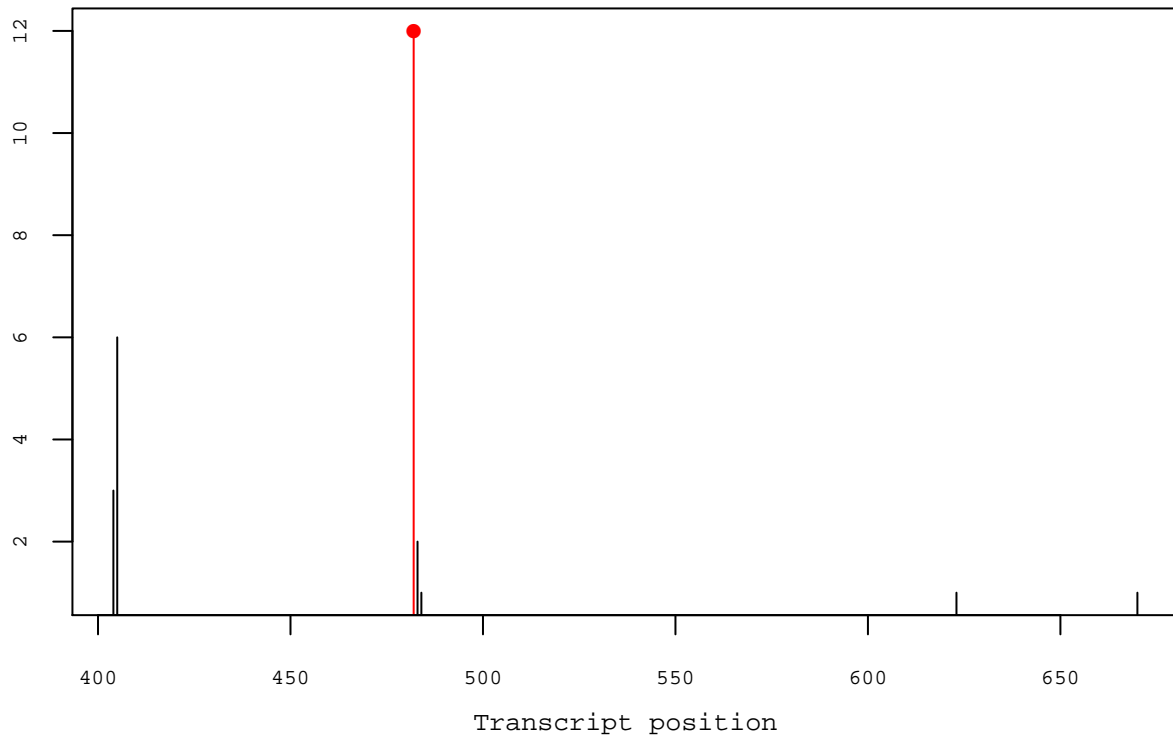


5' GGTTCGACAAAGATACCTACGTCCTCGGGAGC '3
|||||
3' AAGCTGTTTCTATGGATGTAGGAG '5

Fragment Abundance



Cleavage site: 482 Tag abundance: 12 Weighted abundance: 0.229 Category: 0
sRNA abundance: 7 Alignment score: 1 MFE ratio: 0.99 p-value: 0.001