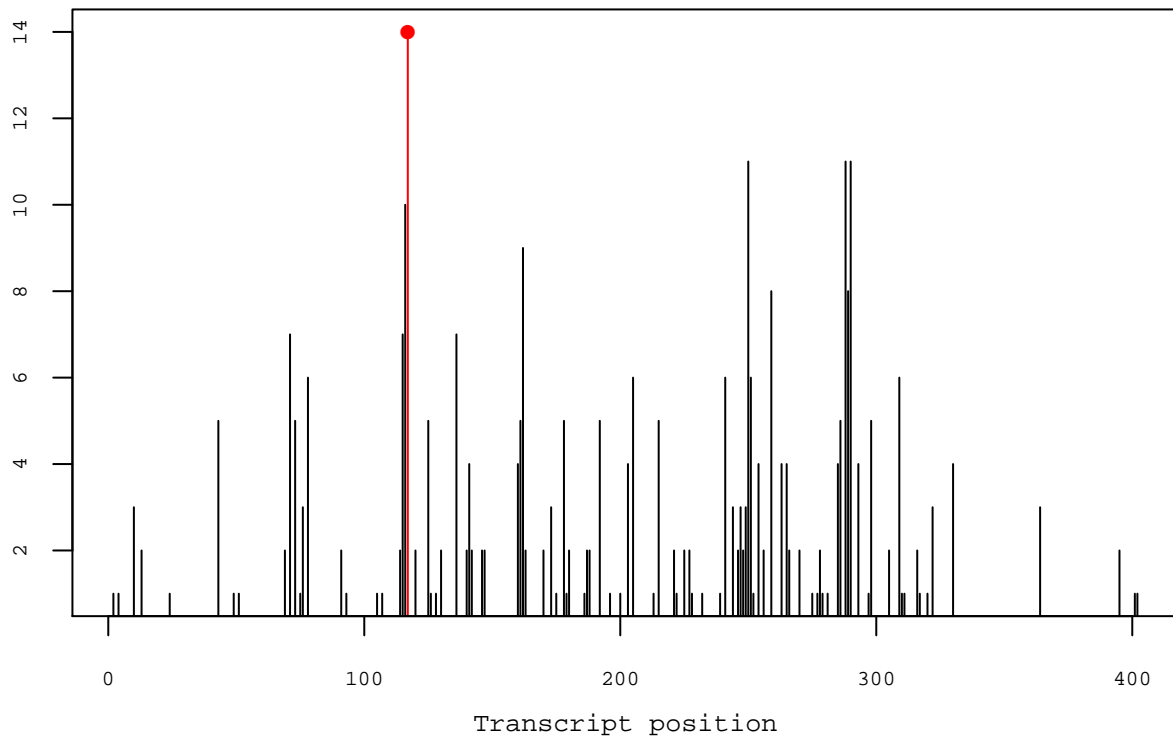


5' AAGTCTTCTAAGTCATTCCAAAATTATCTTT '3
o |||||||||||||||||||||
3' TAGAAGATTTCAGTAAGGTTTTAAA '5

Fragment Abundance



Cleavage site: 117 Tag abundance: 14 Weighted abundance: 0.406 Category: 0
sRNA abundance: 29 Alignment score: 0.5 MFE ratio: 1 p-value: 0.048