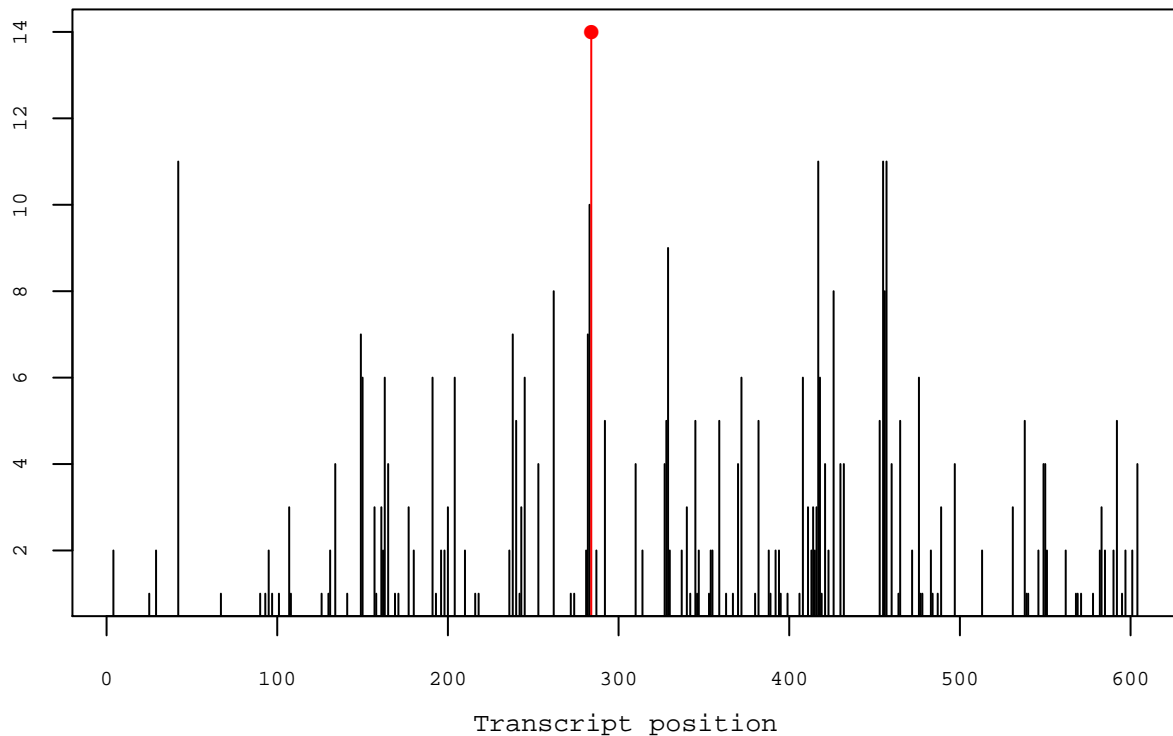


5' AAATCTTCTAAGTCATTCCAAAATTATCTTT '3
|||
3' TAGAAGATTTCAGTAAGGTTTAA '5

Fragment Abundance



Cleavage site: 284 Tag abundance: 14 Weighted abundance: 0.406 Category: 0
sRNA abundance: 29 Alignment score: 0 MFE ratio: 1 p-value: 0.003