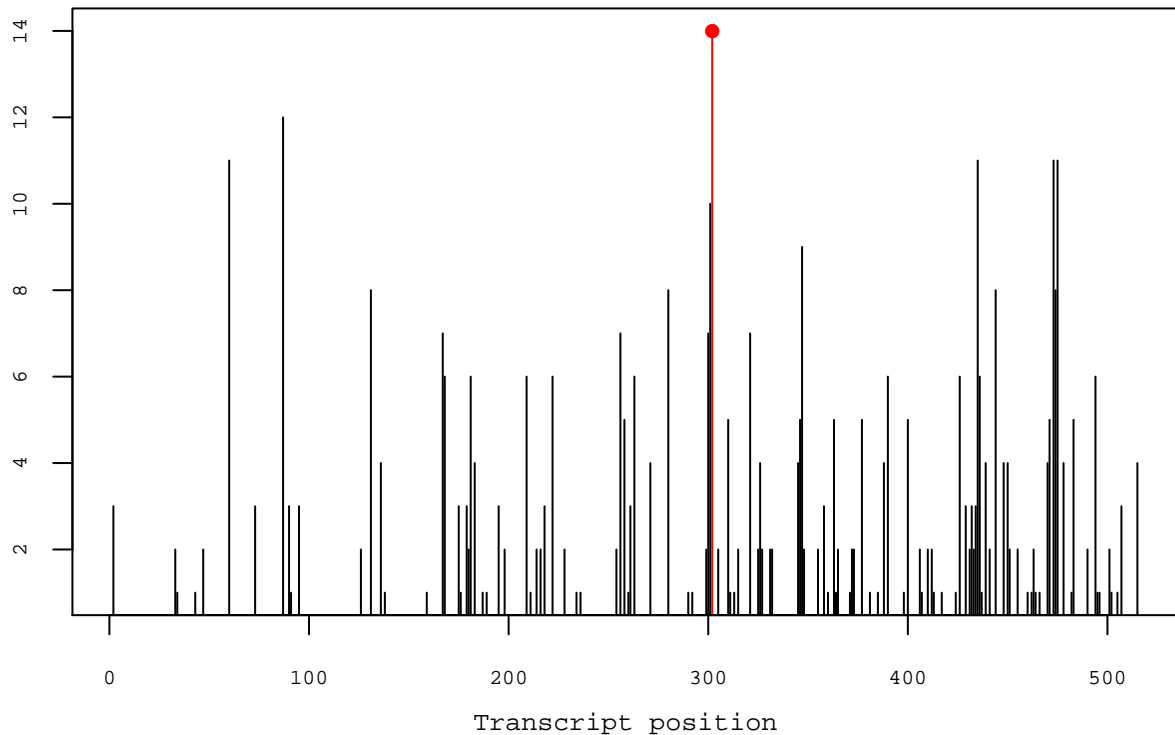


PggD13459_c90_g1_i1

5' AAATCTTCTAAGTCATTCCAAAATTTATCTTT '3
|||
3' TAGAAGATTTCAGTAAGGTTTTAAA '5

Fragment Abundance



Cleavage site: 302 Tag abundance: 14 Weighted abundance: 0.406 Category: 0
sRNA abundance: 29 Alignment score: 0 MFE ratio: 1 p-value: 0.004