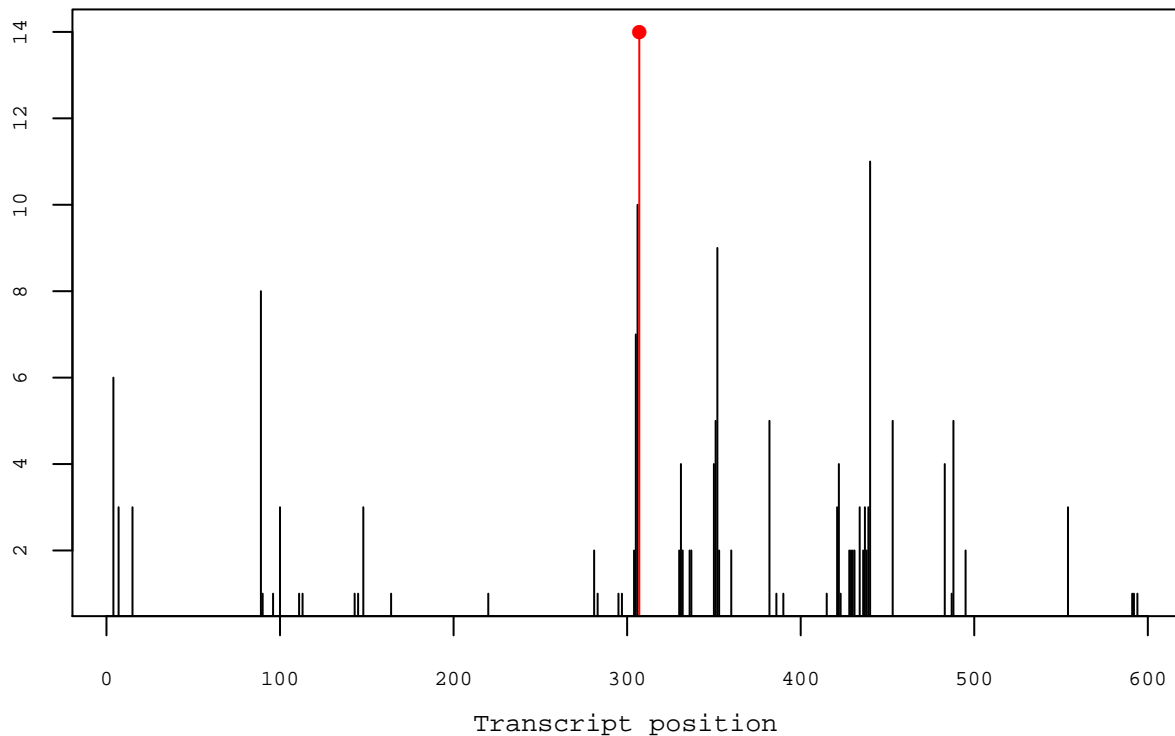


PggD10464_c283_g1_i1

5' AAGTCTTCTAAGTCATTCCAAAATTATCTTT 3'
o |||||||||||||||||||||
3' TAGAAGATTCAGTAAGGTTTAA 5'

Fragment Abundance



Cleavage site: 307 Tag abundance: 14 Weighted abundance: 0.406 Category: 0
sRNA abundance: 29 Alignment score: 0.5 MFE ratio: 1 p-value: 0.031