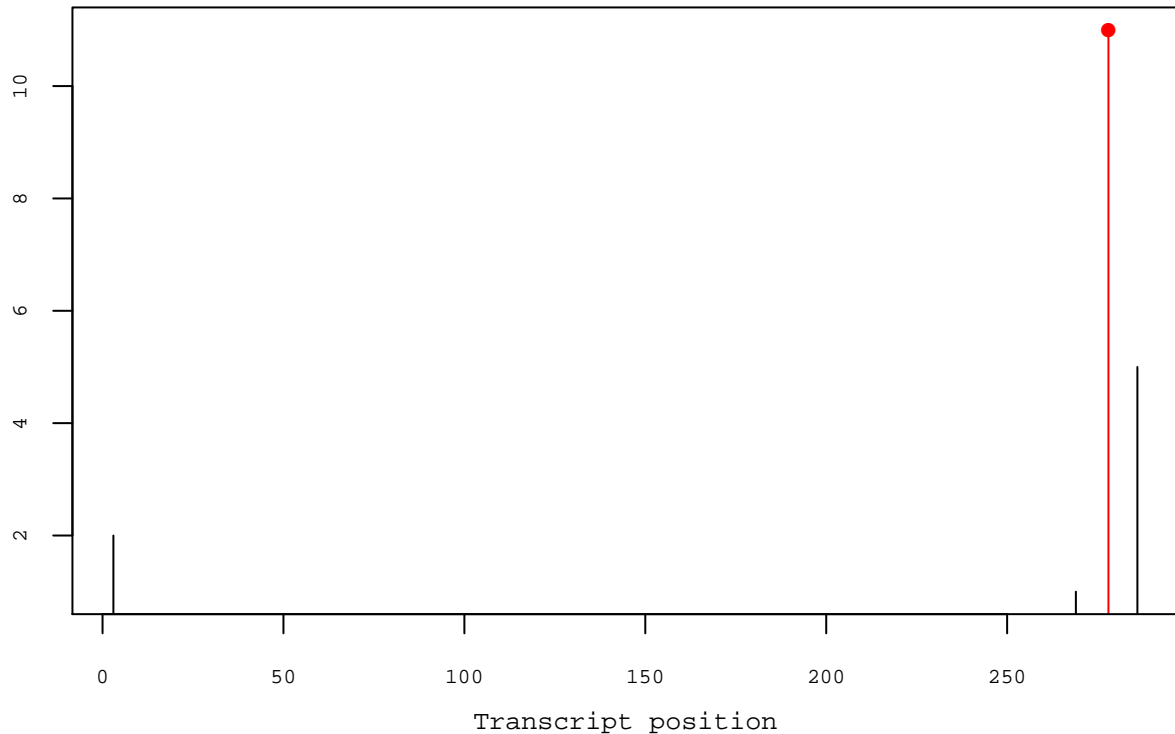


PggD12332_c23_g1_i1

5' GGTTCATTTTAGTCCCAGAGTTTATAAAAGT '3
| | ||||| |||||
3' CTA-GTAAAATCAGAGTCTCAAATA '5

Fragment Abundance



Cleavage site: 278 Tag abundance: 11 Weighted abundance: 2.2 Category: 0
sRNA abundance: 11 Alignment score: 4 MFE ratio: 0.713 p-value: 0.019