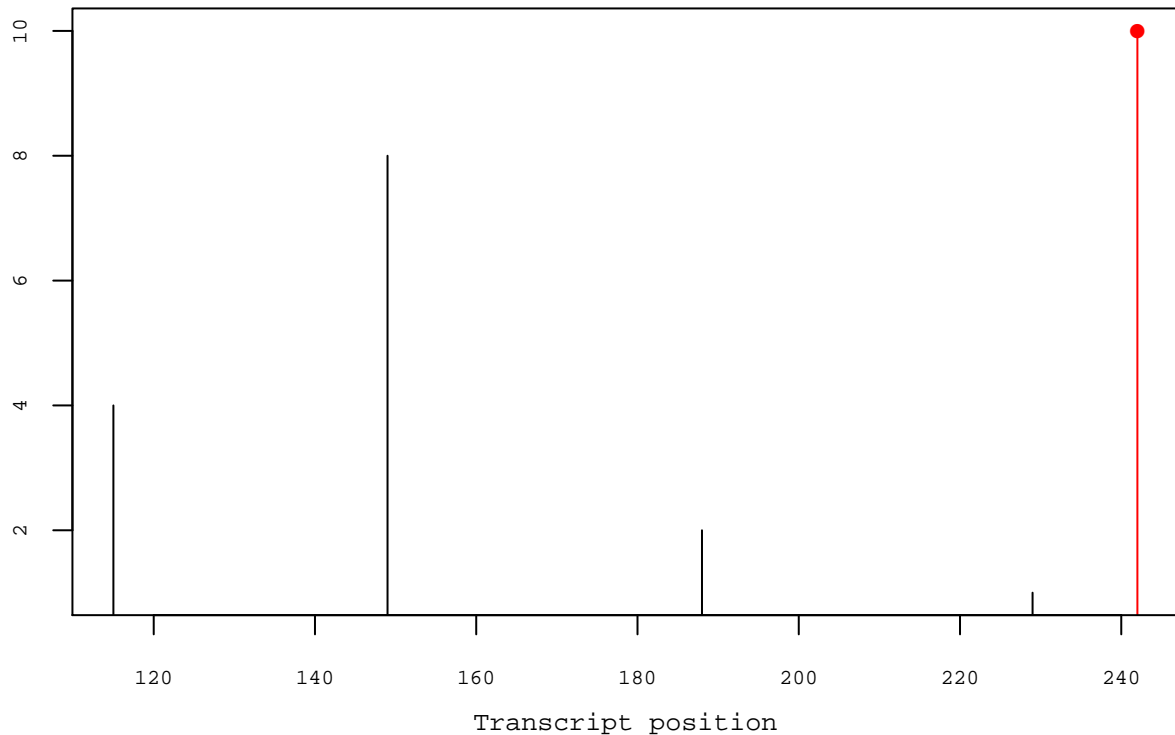


PggD14795_c125_g1_i3

5' TAGCTGGTGTATCCTTCCTGAGCTTTTCTCT '3

||o|o|||||o|||||||
3' CCGCGATAGGGAGGACTCGAA '5

Fragment Abundance



Cleavage site: 242 Tag abundance: 10 Weighted abundance: 10 Category: 0
sRNA abundance: 712 Alignment score: 2 MFE ratio: 0.842 p-value: 0.004