

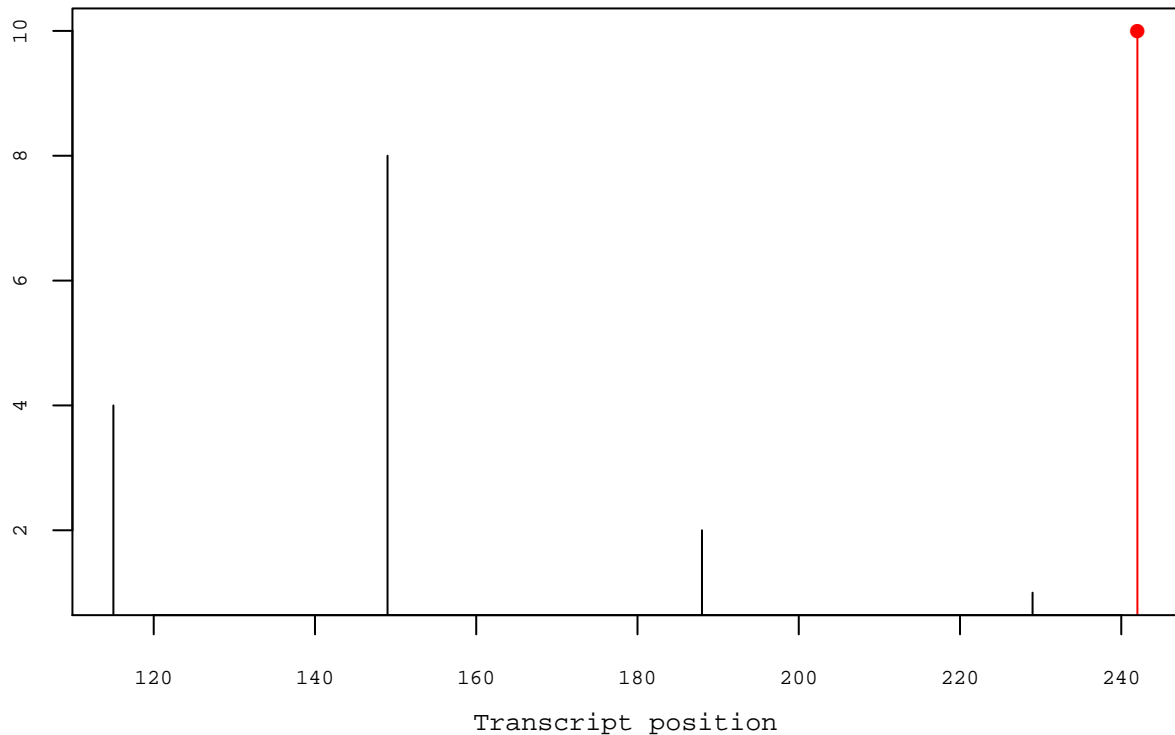
PggD14795_c125_g1_i3

5' TAGC-TGGTGTATCCTTCCTGAGCTTTTCTCT '3

| |||o|o|||||o|||||||

3' GTACCGCGATAGGGAGGACTCGAA '5

Fragment Abundance



Cleavage site: 242 Tag abundance: 10 Weighted abundance: 10 Category: 0
sRNA abundance: 13 Alignment score: 3 MFE ratio: 0.796 p-value: 0.004