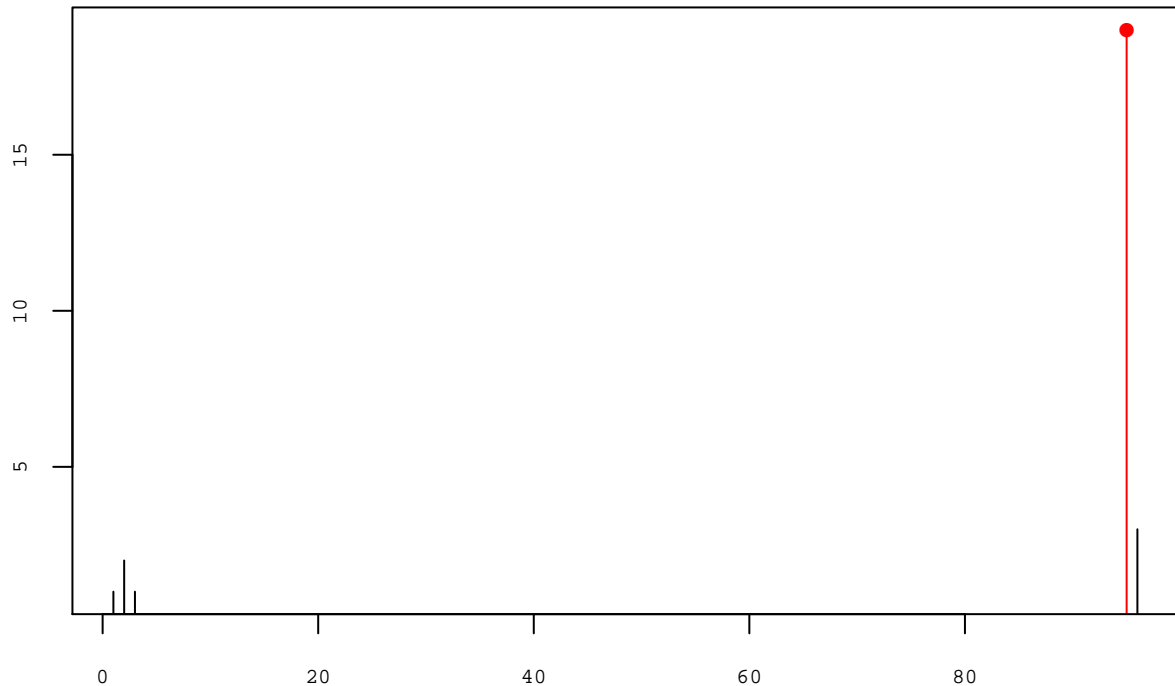


PggD8958_c40_g1_i1

5' TATTTGAGTCTCAGAGTCTATAA-AAGTGCCAA '3

||||○|○| ||||| ||
3' TCAGGGTTTCAGATATTGTT '5

Fragment Abundance



Transcript position

Cleavage site: 95 Tag abundance: 19 Weighted abundance: 1.9 Category: 0
sRNA abundance: 6 Alignment score: 3.5 MFE ratio: 0.753 p-value: 0.011