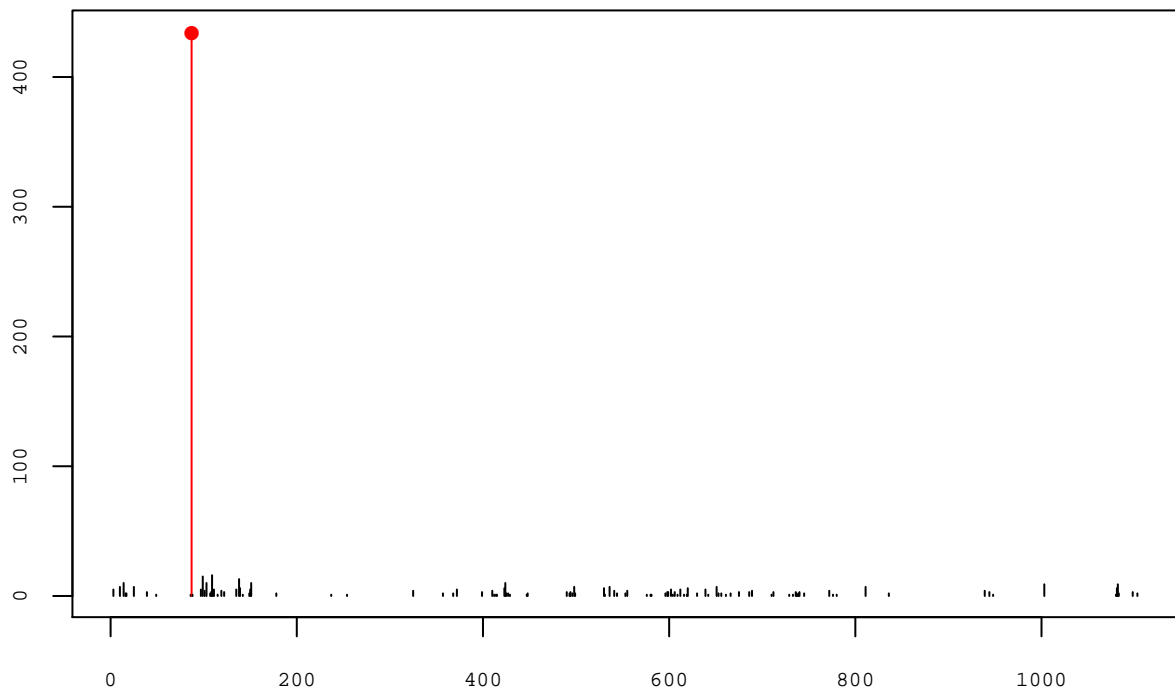


PggD153_c0_g1_i1

5' TCAGATAGAGCCCCCTTCAAACCAAAGCTACG '3
||||| ||||| |||||
3' TCTCGAGGGAAGTTAGGTTT '5

Fragment Abundance



Transcript position

Cleavage site: 87 Tag abundance: 434 Weighted abundance: 108.5 Category: 0
sRNA abundance: 8438 Alignment score: 3 MFE ratio: 0.73 p-value: 0.011