

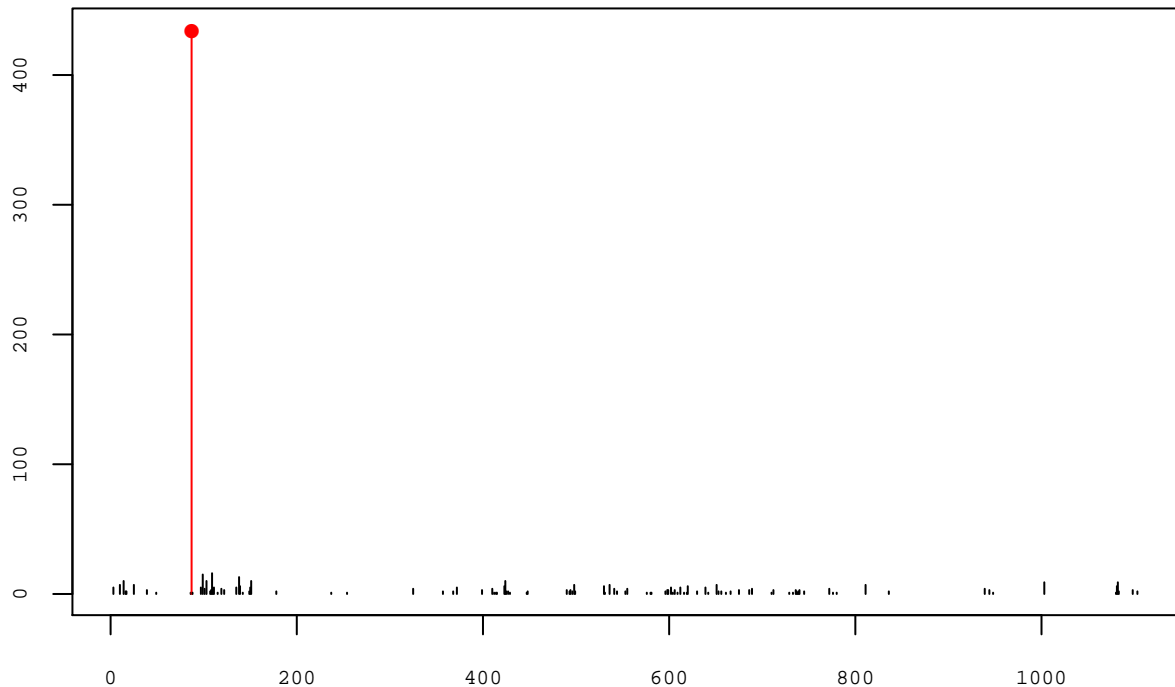
PggD153_c0_g1_i1

5' TCAGATAGAGCCCCCTTCAAACCAAAGCTACG '3

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3' GTCTCGAGGGAAGTTAGGTTT '5

Fragment Abundance



Transcript position

Cleavage site: 87 Tag abundance: 434 Weighted abundance: 108.5 Category: 0
srNA abundance: 5795 Alignment score: 3.5 MFE ratio: 0.706 p-value: 0.01