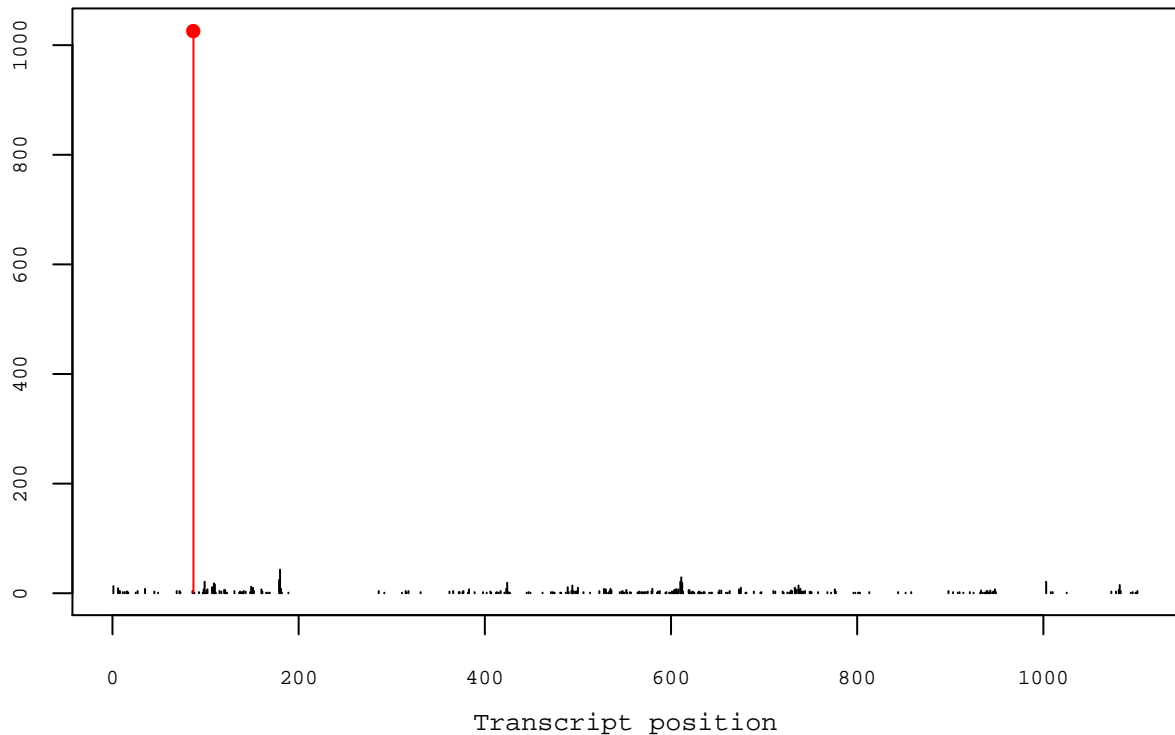


PggD153_c0_g1_i1

5' TCAGATAGAGCCCCCTTCAAACCAAAGCTACG '3
||||| ||||| |||||
3' ATCTCGAGGGAAGTTAGGTTT '5

Fragment Abundance



Cleavage site: 87 Tag abundance: 1026 Weighted abundance: 256.5 Category: 0
sRNA abundance: 106556 Alignment score: 3 MFE ratio: 0.74 p-value: 0.005