

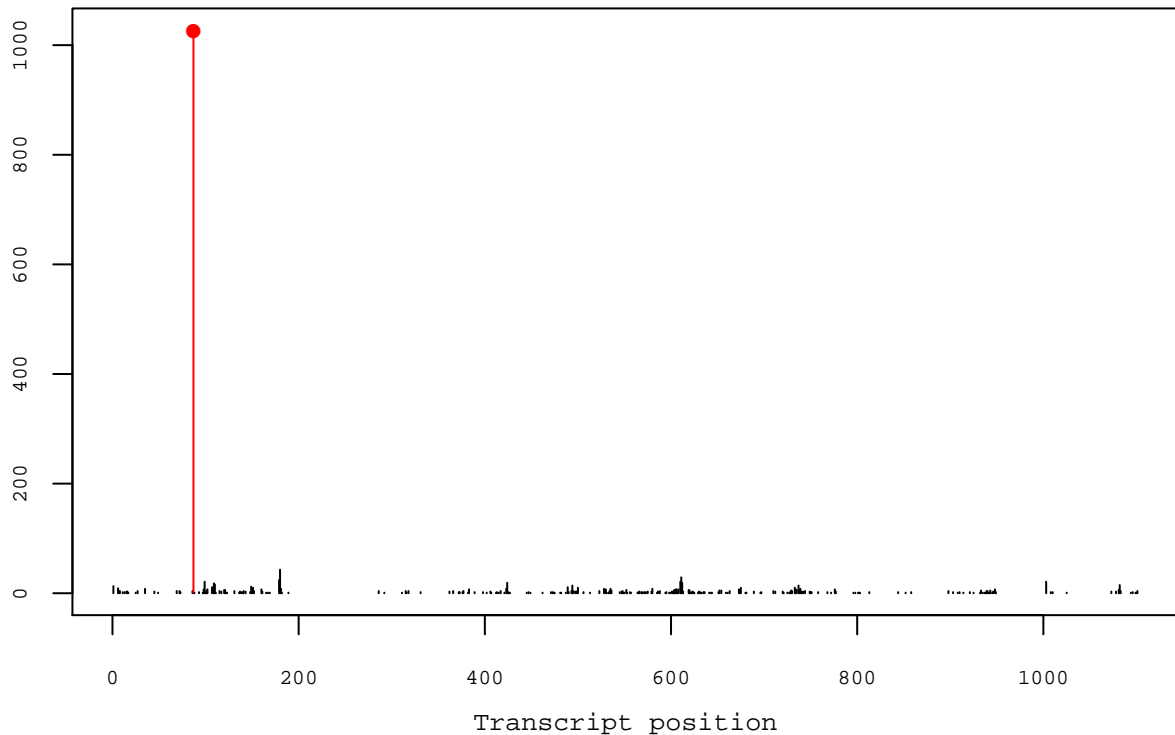
PggD153_c0_g1_i1

5' TCAGATAGAGCCCCCTTCAAACCAAAGCTACG '3

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

3' GTCTCGAGGGAAGTTAGGTTT '5

Fragment Abundance



Cleavage site: 87 Tag abundance: 1026 Weighted abundance: 256.5 Category: 0
sRNA abundance: 8957 Alignment score: 3.5 MFE ratio: 0.706 p-value: 0.009