

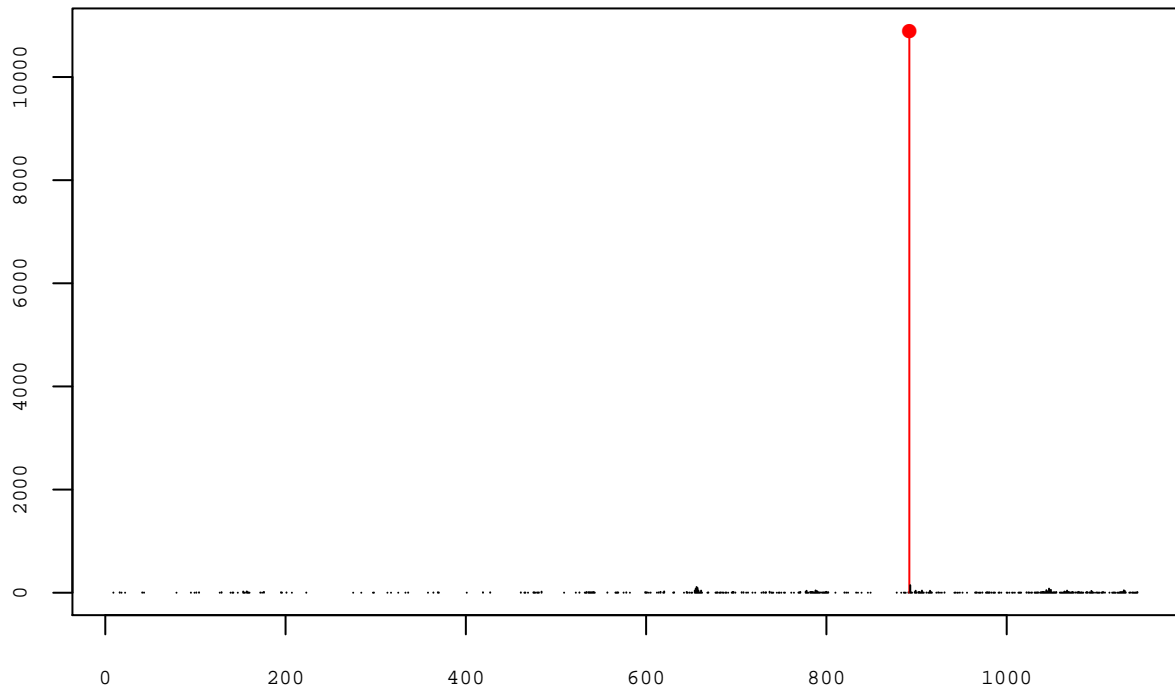
PggD4594_c6_g1_i1

5' GCACAGAGGGAAC-CCCTTCAGTCCAGTAACTT '3

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3' TCCCTCGAGGGAAGTCAGGTT '5

Fragment Abundance



Transcript position

Cleavage site: 892 Tag abundance: 10894 Weighted abundance: 3631.333 Category: 0
sRNA abundance: 13228 Alignment score: 2.5 MFE ratio: 0.72 p-value: 0.007