

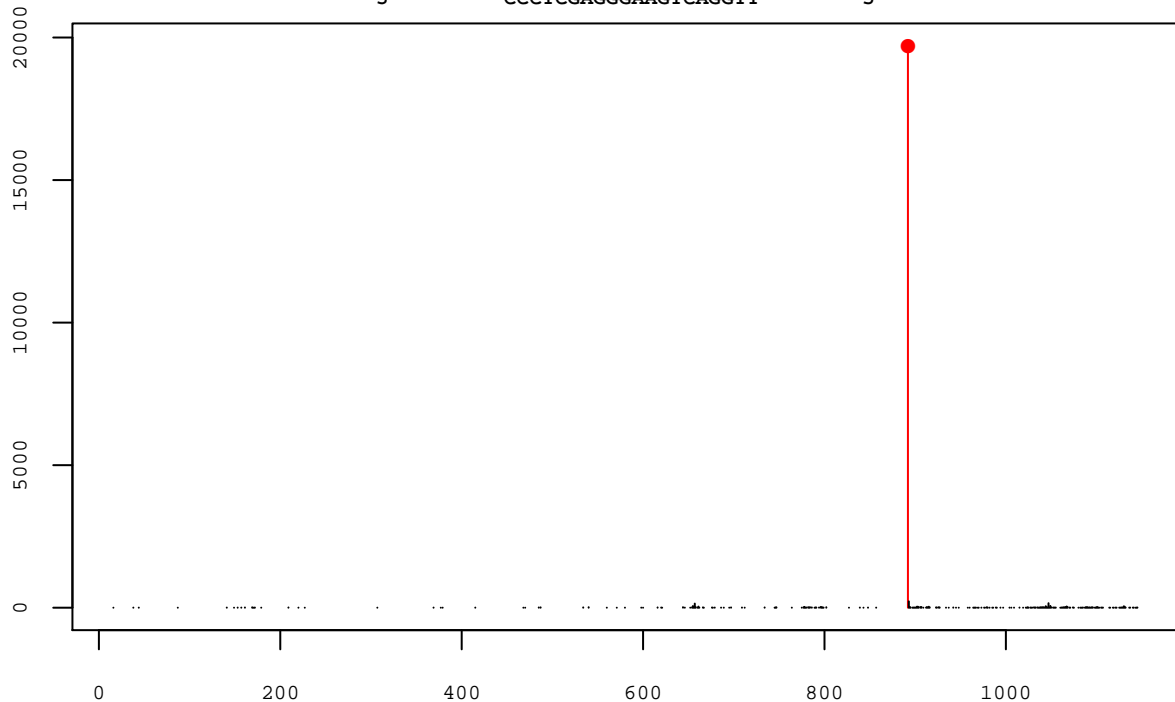
PggD4594\_c6\_g1\_i1

5' GCACAGAGGGAAC-CCCTTCAGTCCAGTAACTT '3

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3' CCCTCGAGGGAAGTCAGGTT '5

Fragment Abundance



Transcript position

Cleavage site: 892 Tag abundance: 19706 Weighted abundance: 6568.667 Category: 0  
sRNA abundance: 1130 Alignment score: 2.5 MFE ratio: 0.709 p-value: 0.007