

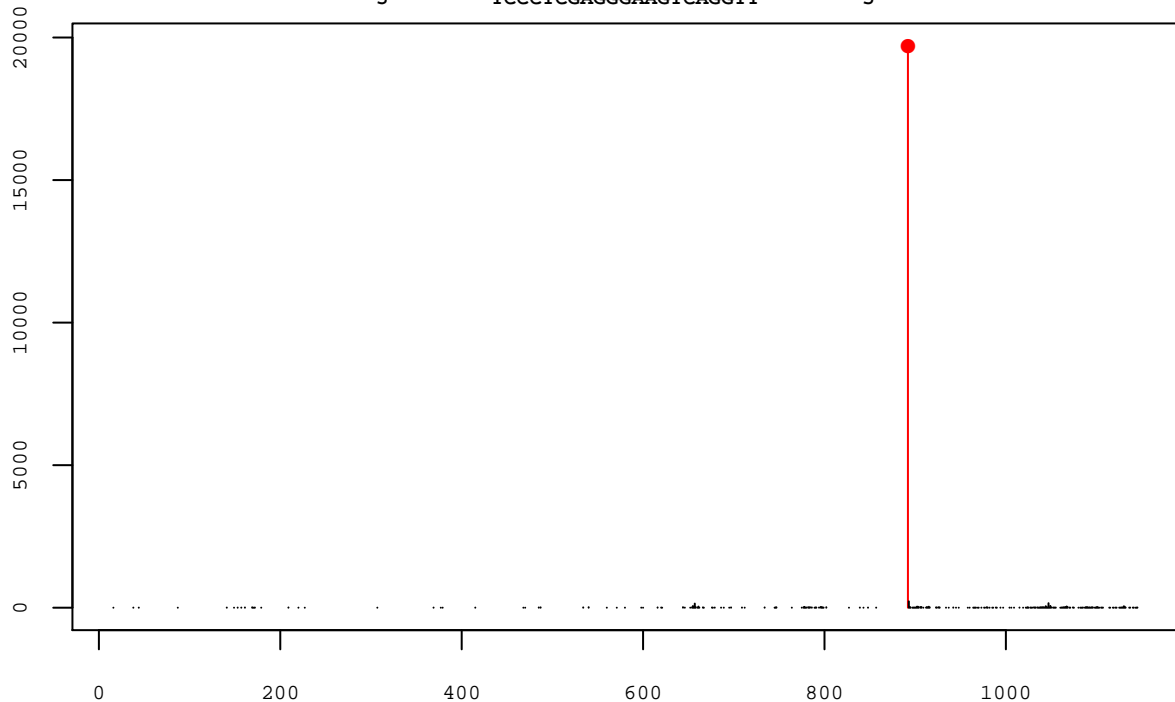
PggD4594_c6_g1_i1

5' GCACAGAGGGAAC-CCCTTCAGTCCAGTAACTT '3

||||| | ||||| ||||| |||○

3' TCCCTCGAGGGAAGTCAGGTT '5

Fragment Abundance



Cleavage site: 892 Tag abundance: 19706 Weighted abundance: 6568.667 Category: 0
sRNA abundance: 8354 Alignment score: 2.5 MFE ratio: 0.72 p-value: 0.007