

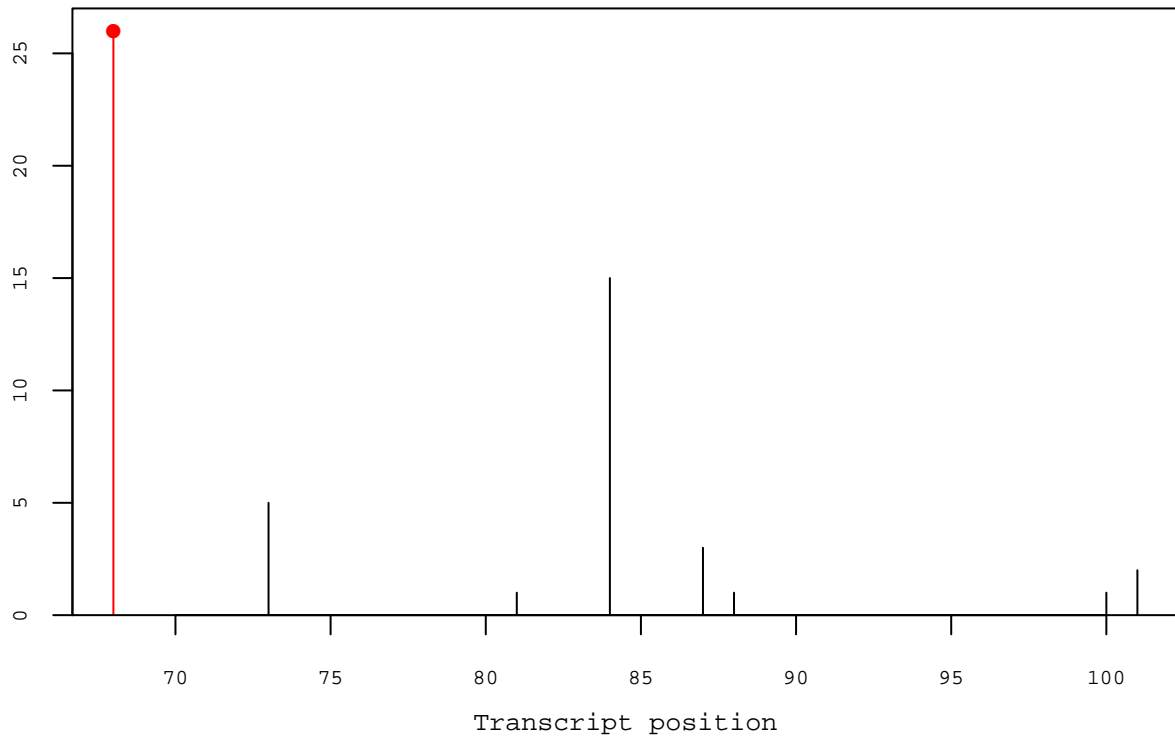
PggD1888_c134_g1_i1

5' CATGTAAC TTGTTTGCAGGTTACAGACGTACG 3'

|||||

3' TTGTACAAACGTCCAATGTCT 5'

Fragment Abundance



Cleavage site: 68 Tag abundance: 26 Weighted abundance: 1.083 Category: 0
sRNA abundance: 10 Alignment score: 1 MFE ratio: 0.881 p-value: 0.016