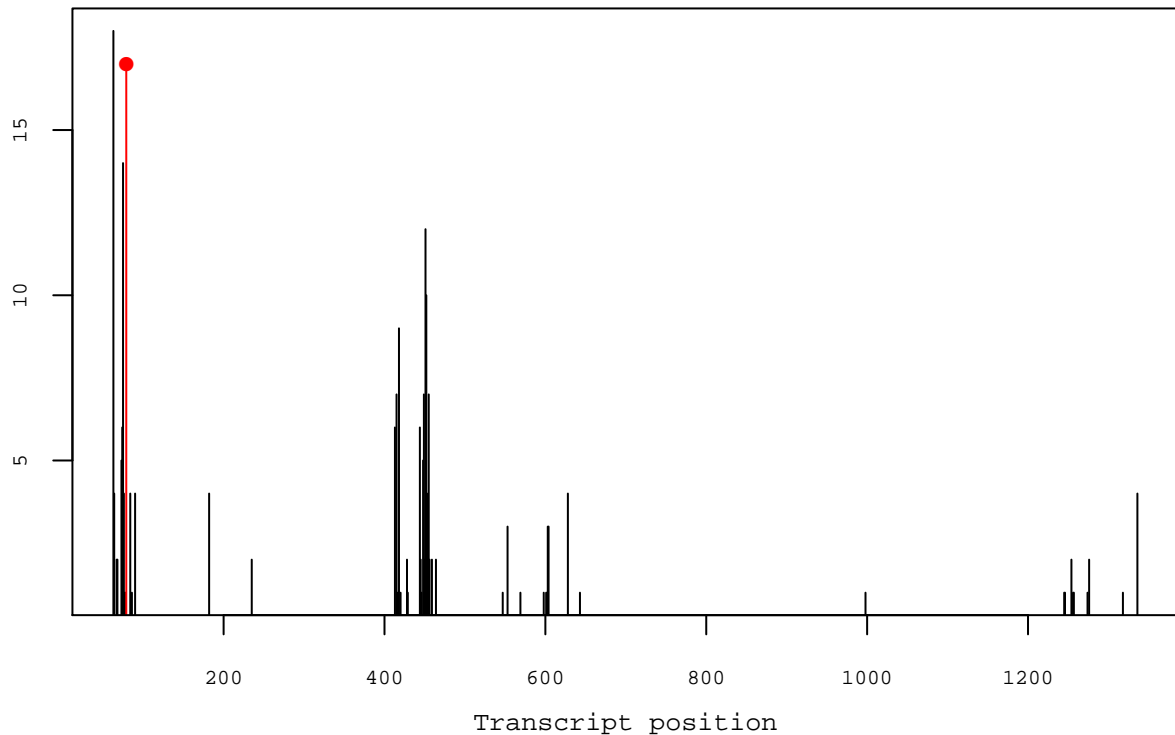


PggD12497_c103_g1_i1

5' ATGCATGCGGTGTCCGATCCTGACCGTAGATC '3
|||||
3' CGTACGCCACAGGCTAAGACTGGC '5

Fragment Abundance



Cleavage site: 79 Tag abundance: 17 Weighted abundance: 5.667 Category: 2
sRNA abundance: 8 Alignment score: 2 MFE ratio: 0.909 p-value: 0.023