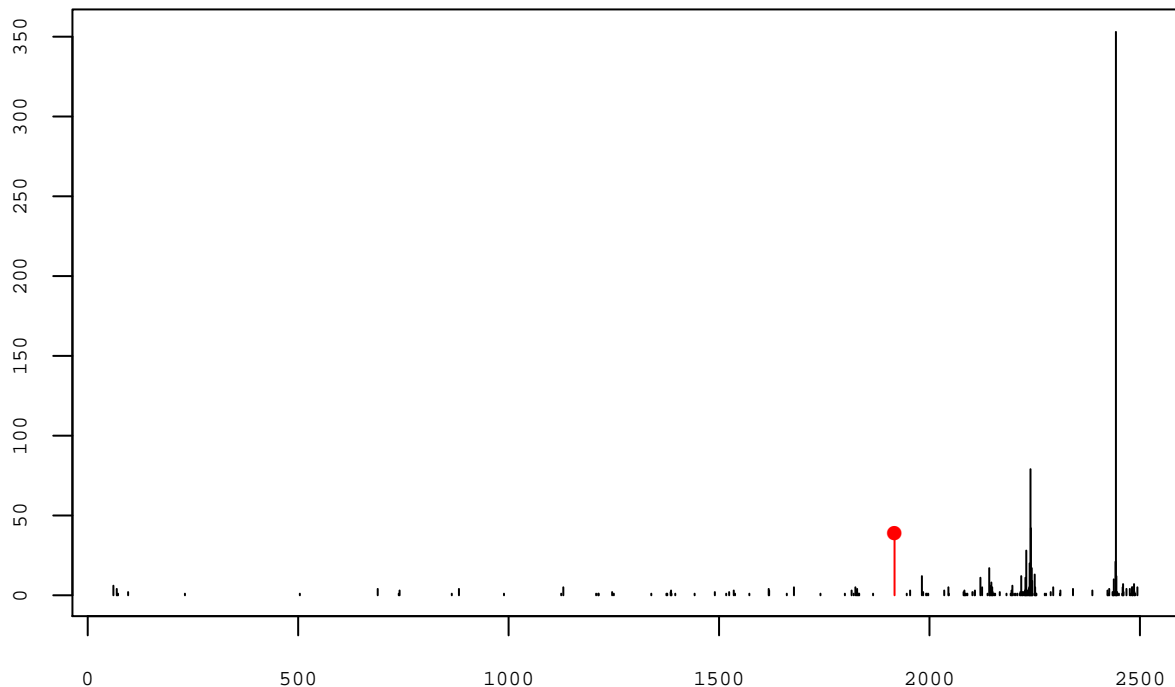


PggD12654_c9_g1_i1

5' CTCTTCTGCAGCATCATCAGGATTCTCAGCCT '3
|||||o|||||
3' TACGTCGTAGTAGTTCTAAGA '5

Fragment Abundance



Transcript position

Cleavage site: 1917 Tag abundance: 39 Weighted abundance: 13 Category: 2
sRNA abundance: 37 Alignment score: 2 MFE ratio: 0.983 p-value: 0.042