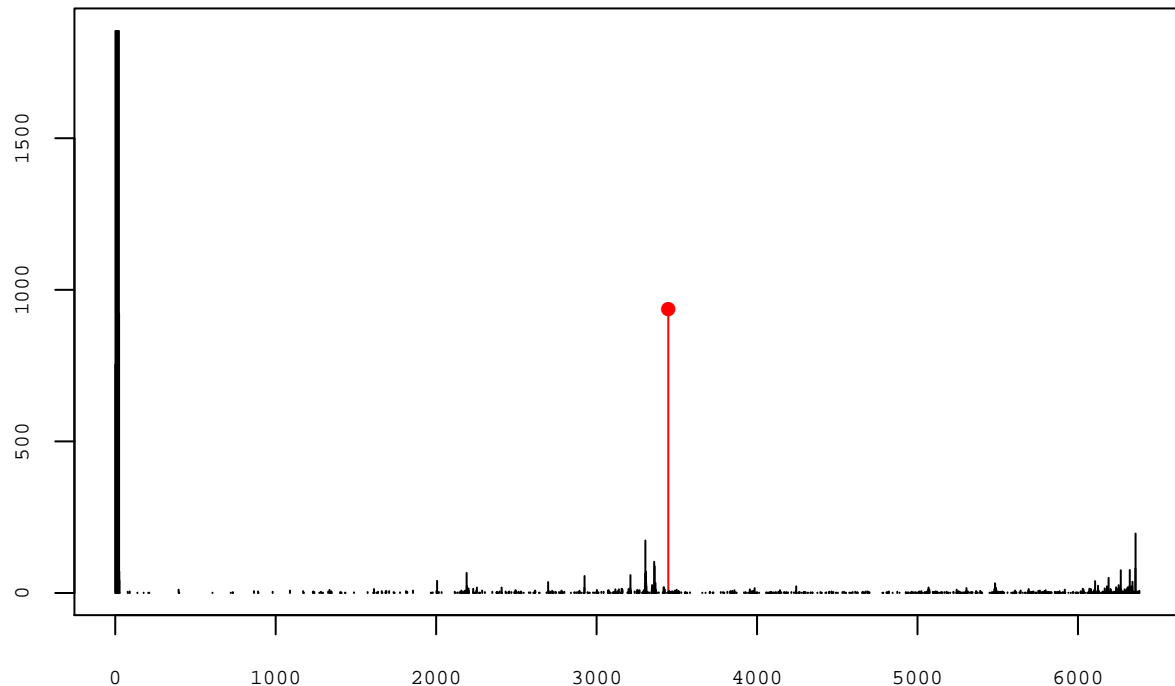


PggD11644_c3_g1_i1

5' ATGAGCTGGATGCAGAGGTCTTATCGATGTCA '3
|||||
3' CCTACGTCTCCA-AATAGCT '5

Fragment Abundance



Transcript position

Cleavage site: 3447 Tag abundance: 937 Weighted abundance: 937 Category: 0
sRNA abundance: 262 Alignment score: 2 MFE ratio: 0.916 p-value: 0