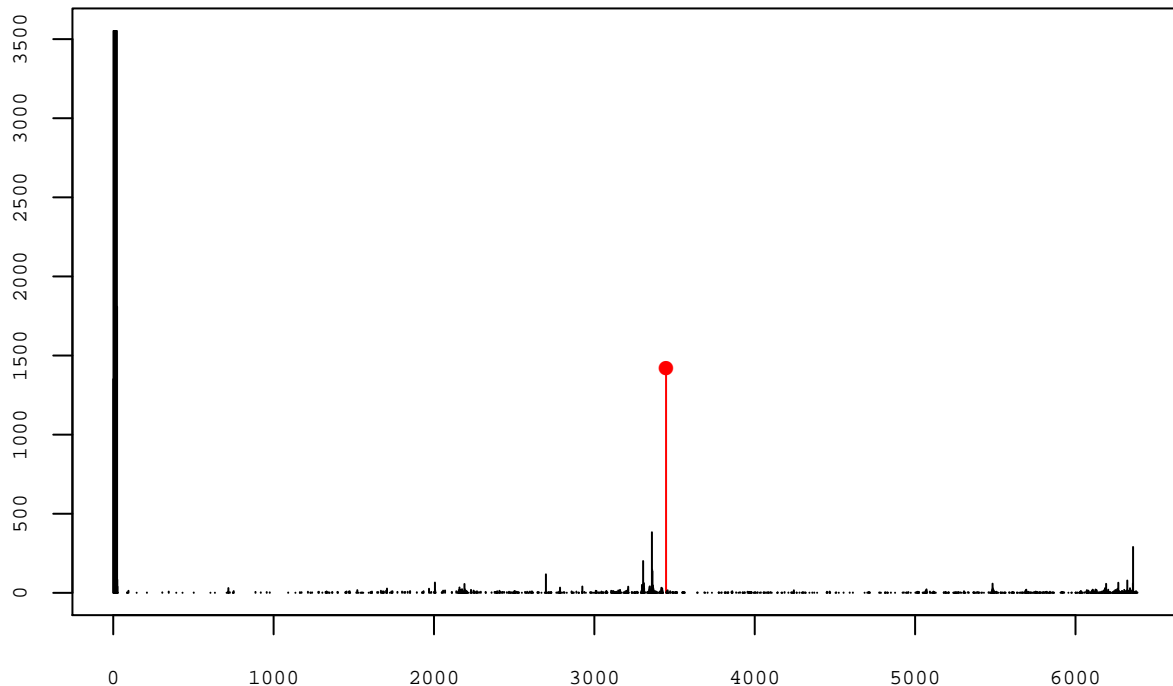


PggD11644_c3_g1_i1

5' ATGAGCTGGATGCAGAGGTCTTATCGATGTCA '3
|o||||||||| |||||
3' GGCCTACGTCTCCA-AATAGCT '5

Fragment Abundance



Transcript position

Cleavage site: 3447 Tag abundance: 1422 Weighted abundance: 1422 Category: 0
sRNA abundance: 1985 Alignment score: 2.5 MFE ratio: 0.872 p-value: 0