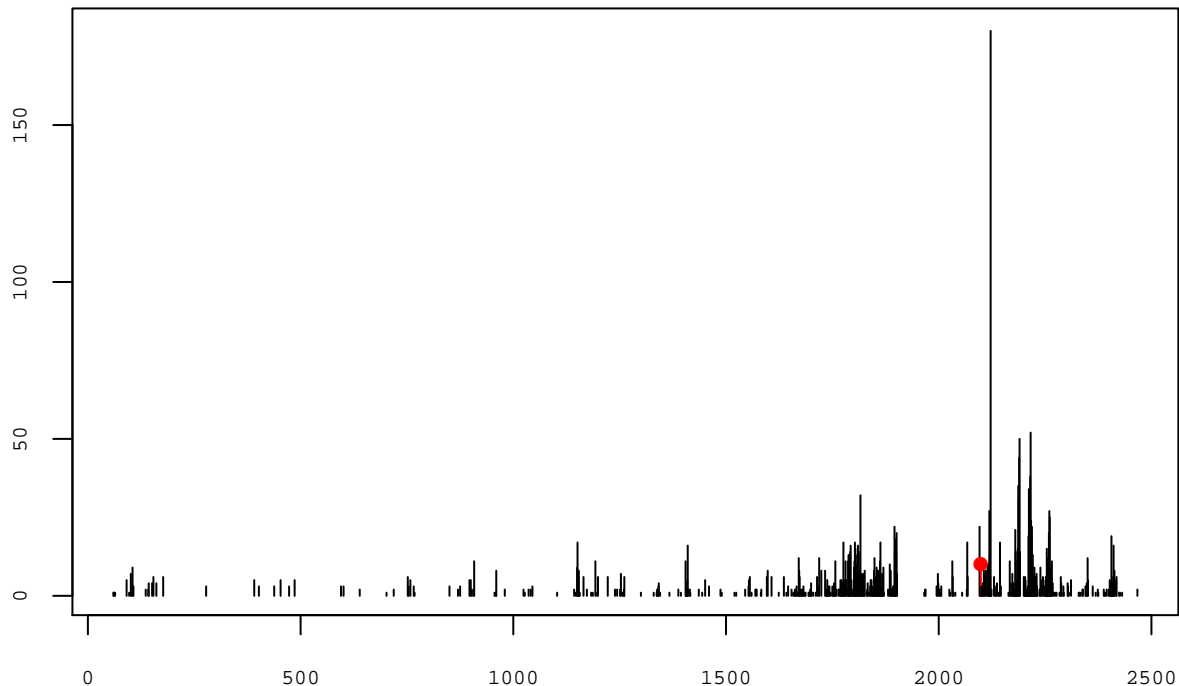


PggD7047_c0_g1_i1

5' GAGGTTACAGGGGGTATAACACGGATATAACT '3
||| ||| |o| ||| ||| |o| |||
3' CCAATGTCTCCCATATTGCGTCTA '5

Fragment Abundance



Transcript position

Cleavage site: 2099 Tag abundance: 10 Weighted abundance: 10 Category: 2
sRNA abundance: 10 Alignment score: 3.5 MFE ratio: 0.838 p-value: 0.036