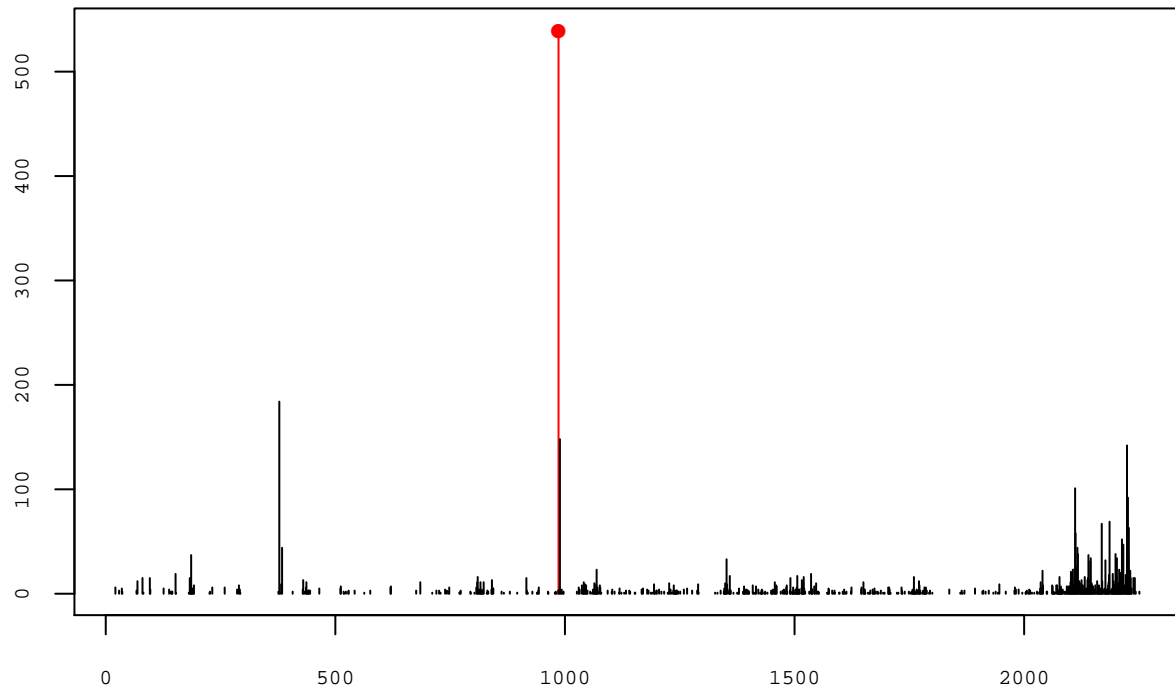


PggD5898_c0_g1_i3

5' GCCCATA-GGATATTGGCGCGGCTCAATCATCA '3

3' TCACTATAACCGCGCCGAGTT '5

Fragment Abundance



Transcript position

Cleavage site: 986 Tag abundance: 539 Weighted abundance: 269.5 Category: 0
sRNA abundance: 71 Alignment score: 2 MFE ratio: 0.886 p-value: 0.002