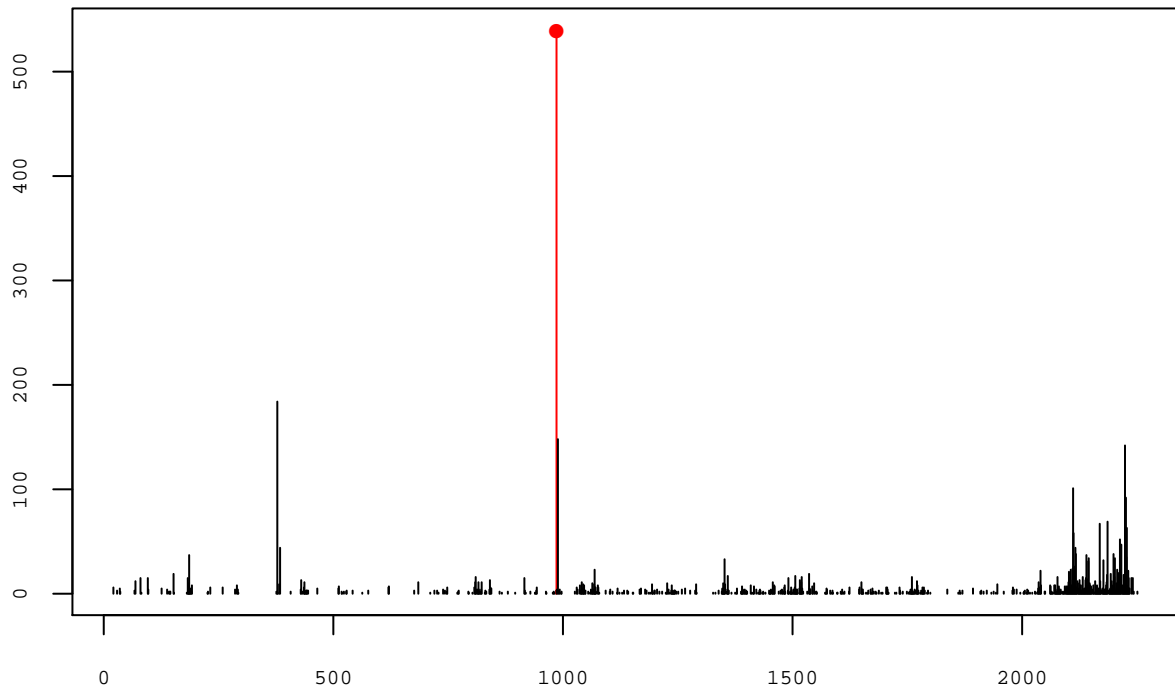


PggD5898_c0_g1_i3

5' GCCCATAGGATATTGGCGCGGCTCAATCATCA '3

|||||||o|||||||
3' CGCTATAACCGTGCCGAGTT '5

Fragment Abundance



Transcript position

Cleavage site: 986 Tag abundance: 539 Weighted abundance: 269.5 Category: 0
sRNA abundance: 387 Alignment score: 3 MFE ratio: 0.878 p-value: 0.002