

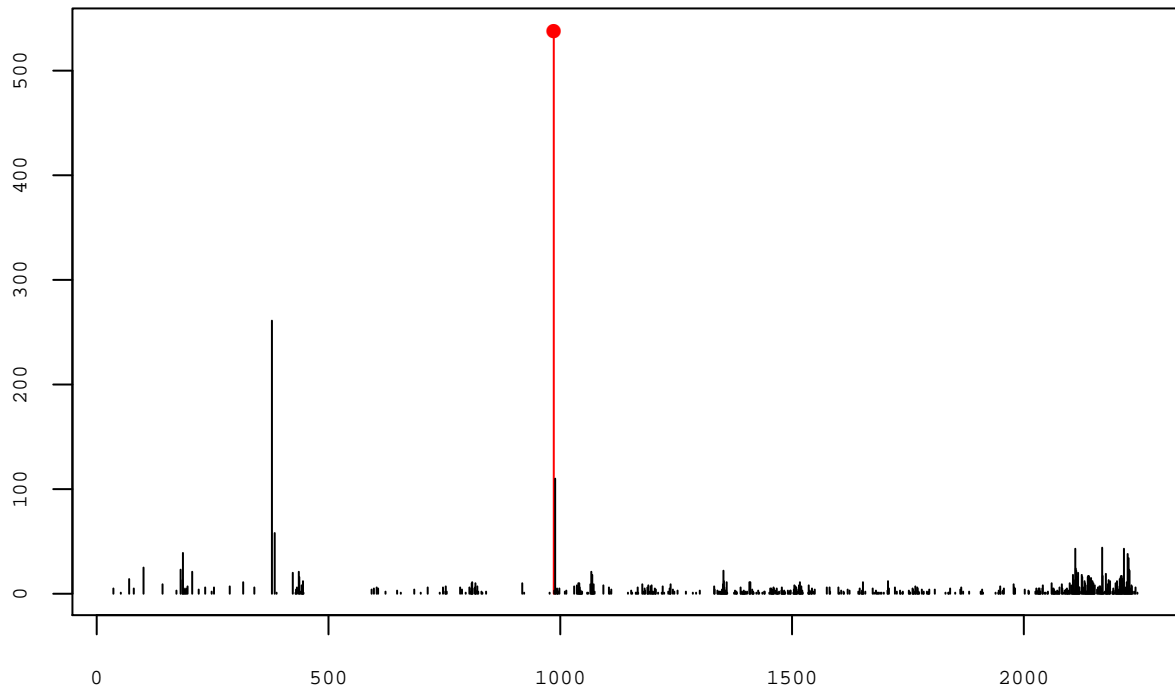
PggD5898_c0_g1_i3

5' GCCCATAGGATATTGGCGGGCTCAATCATCA '3

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3' CCTCTATAACTGCGCCGAGTT '5

Fragment Abundance



Transcript position

Cleavage site: 986 Tag abundance: 538 Weighted abundance: 269 Category: 0
sRNA abundance: 56 Alignment score: 3.5 MFE ratio: 0.844 p-value: 0.002