

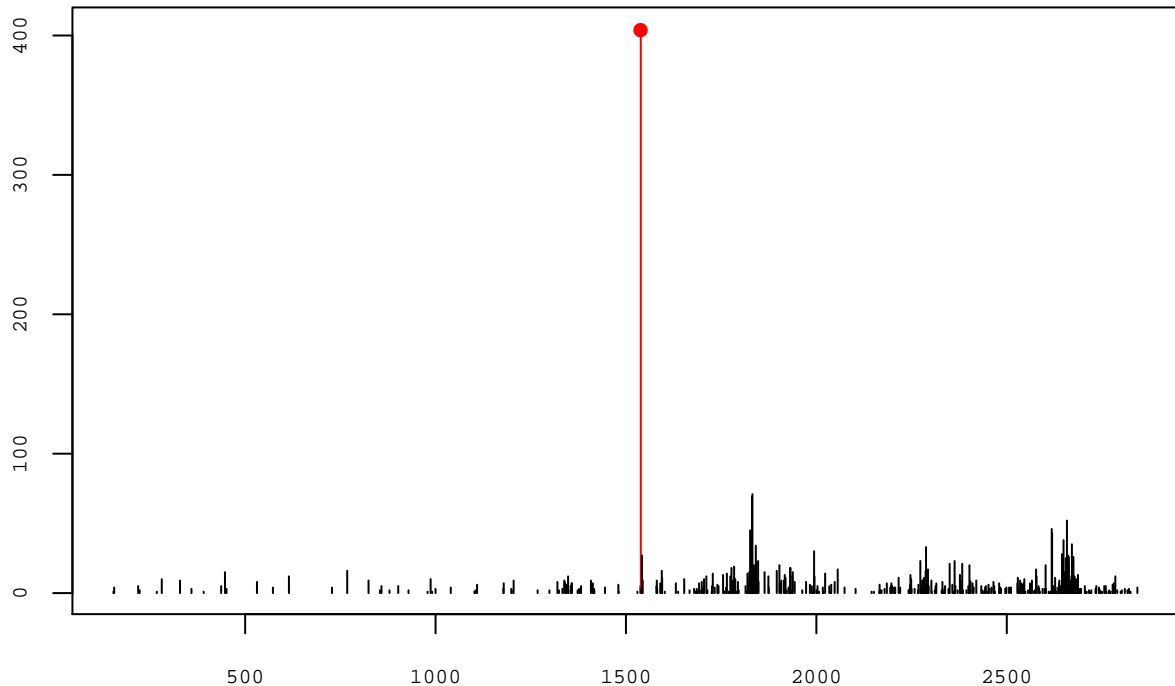
PggD2305_c3_g1_i1

5' GCACATGG-GATATTGGCGCGGCTCAATCACCA '3

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3' CCTCTATAACTGCGCCGAGTT '5

Fragment Abundance



Transcript position

Cleavage site: 1539 Tag abundance: 404 Weighted abundance: 371 Category: 0
sRNA abundance: 56 Alignment score: 2 MFE ratio: 0.857 p-value: 0.001