

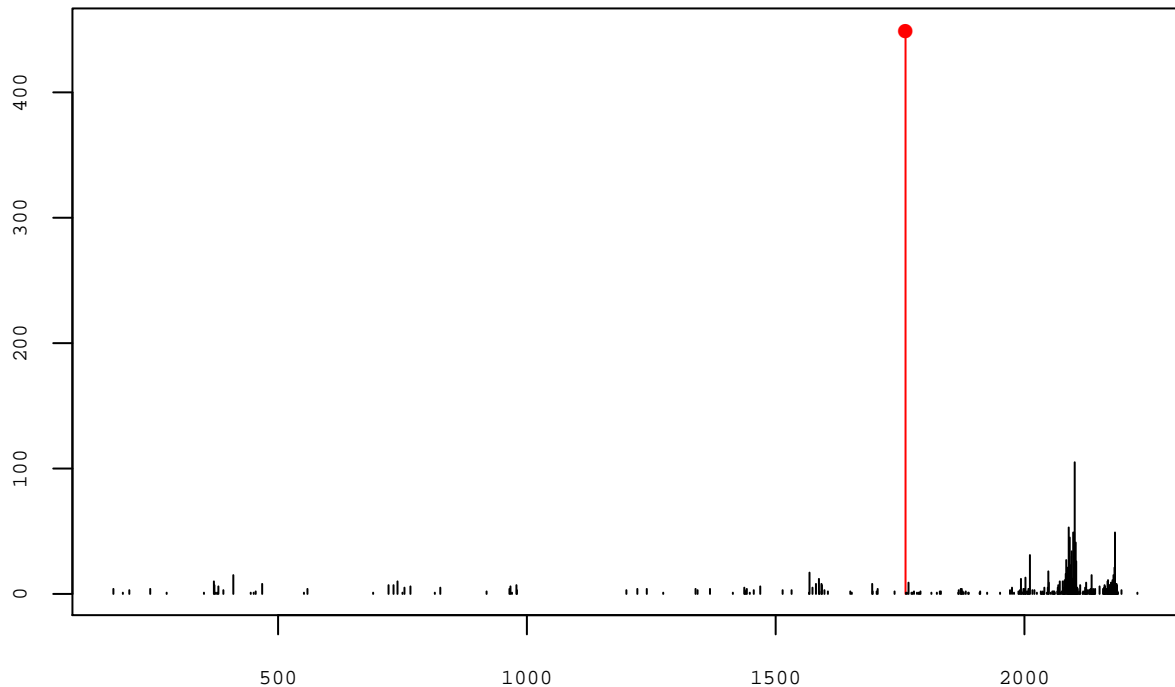
PggD5310_c30_g1_i1

5' TTCCACAGGGGGA-CCCTTCAGTCCAATCCATC '3

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3' CCTCGAGGGAAGTCAGGTT '5

Fragment Abundance



Transcript position

Cleavage site: 1761 Tag abundance: 449 Weighted abundance: 224.5 Category: 0
srRNA abundance: 168 Alignment score: 2.5 MFE ratio: 0.768 p-value: 0.005