

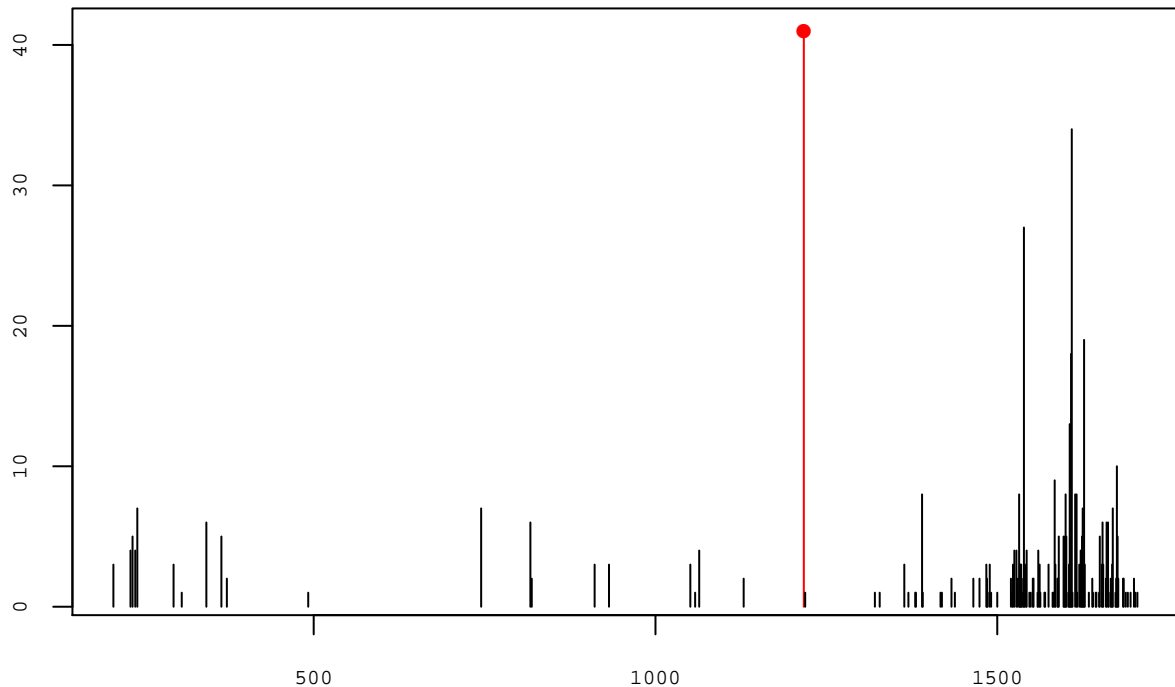
PggD4707_c6_g1_i1

5' TTTAGTAGGGGGA-CCCTTCAGTCCAATTCACC '3

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3' CCTCGAGGGAAGTCAGGTT '5

Fragment Abundance



Cleavage site: 1217 Tag abundance: 41 Weighted abundance: 41 Category: 0
sRNA abundance: 168 Alignment score: 2.5 MFE ratio: 0.768 p-value: 0.007