

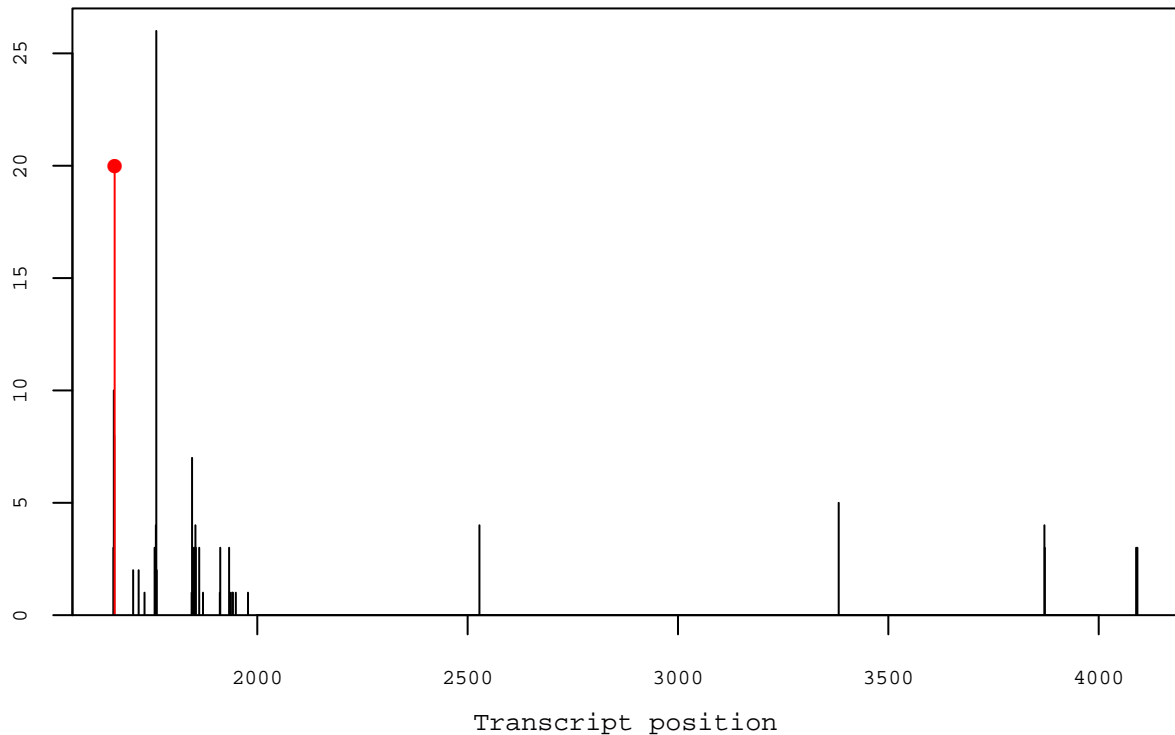
PggD226_c246_g1_i4

5' AAATGGCTTGTTG-TCGTGAACAGCTCATGTTTC '3

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3' CCGAGCAGCAAGCTCTTGTCGAG '5

Fragment Abundance



Cleavage site: 1661 Tag abundance: 20 Weighted abundance: 20 Category: 2
sRNA abundance: 5 Alignment score: 4 MFE ratio: 0.723 p-value: 0.003