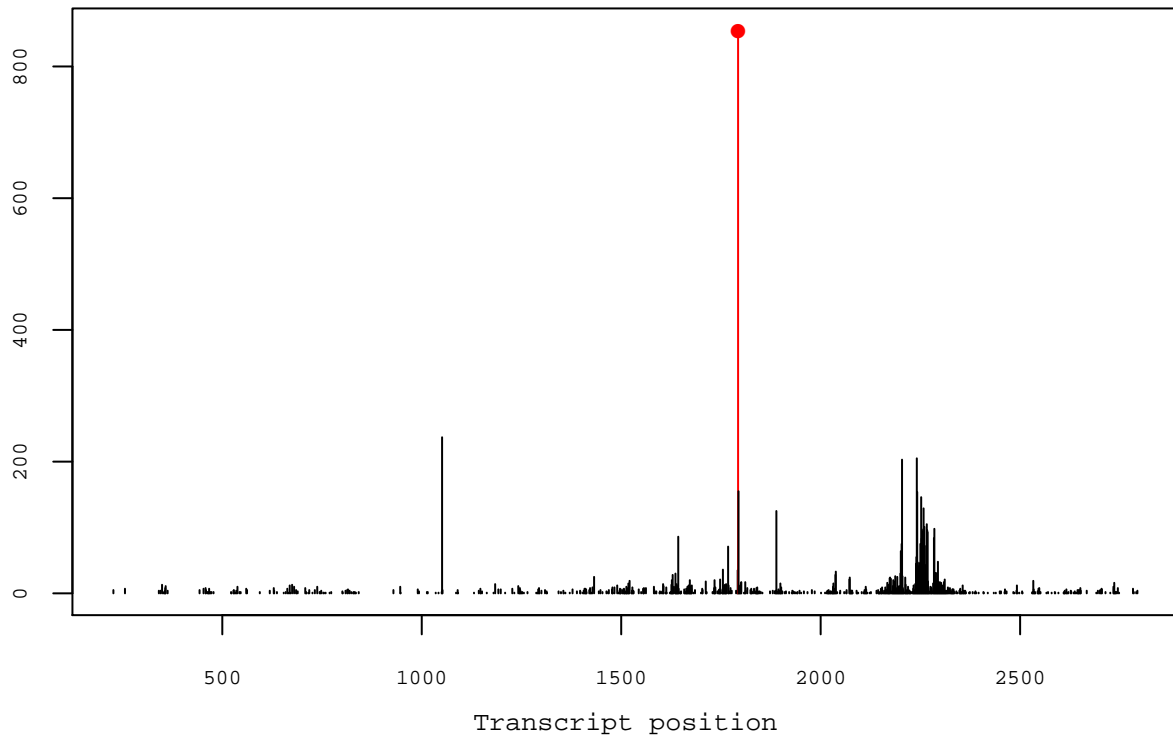


PggD2930_c0_g1_i1

5' ACACAGTGGGGAACCCTTCAGTCCAGTAACCC '3
|| |||o| |||||o
3' TC-CCTCGAGGGAAGTCAGGTT '5

Fragment Abundance



Cleavage site: 1793 Tag abundance: 854 Weighted abundance: 213.5 Category: 0
sRNA abundance: 13228 Alignment score: 4 MFE ratio: 0.704 p-value: 0.008