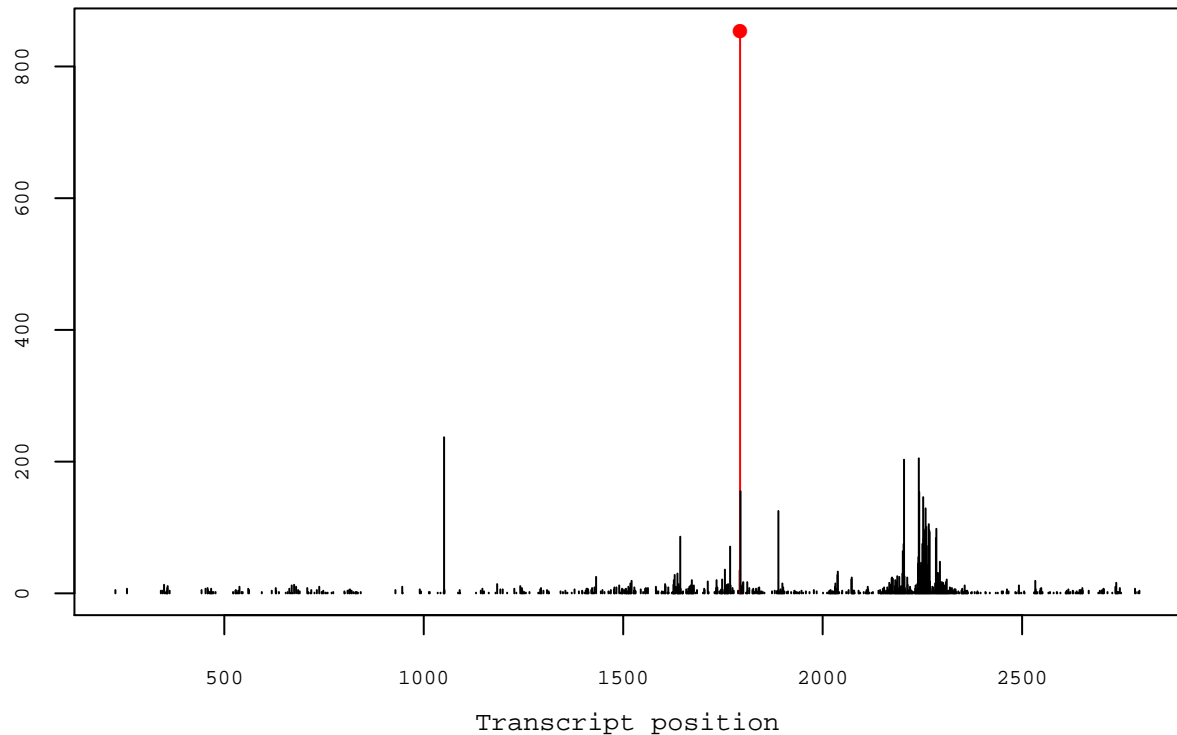


PggD2930_c0_g1_i1

5' ACACAGTGGGGAACCCTTCAGTCCAGTAACCC 3'
||o| |||||o
3' CCTCGAGGGAAGTCAGGTT 5'

Fragment Abundance



Cleavage site: 1793 Tag abundance: 854 Weighted abundance: 213.5 Category: 0
sRNA abundance: 282 Alignment score: 3 MFE ratio: 0.768 p-value: 0.007