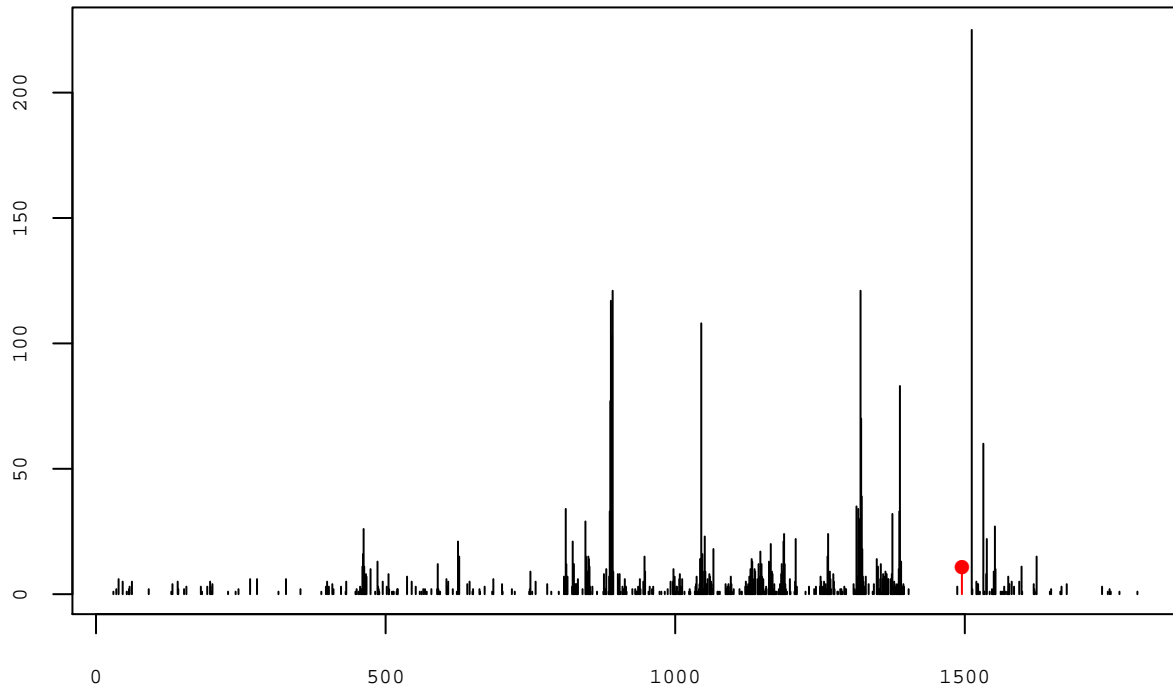


PggD6653_c50_g1_i1

5' GAGCTGCGTATGTCATAGGTTATTCAGAGGGT '3
||| ||| |o| ||| ||| ||| ||| |||
3' CGACGCATGCAGTCTCCAATAAGT '5

Fragment Abundance



Transcript position

Cleavage site: 1495 Tag abundance: 11 Weighted abundance: 1.21 Category: 2
sRNA abundance: 12 Alignment score: 2.5 MFE ratio: 0.815 p-value: 0.042