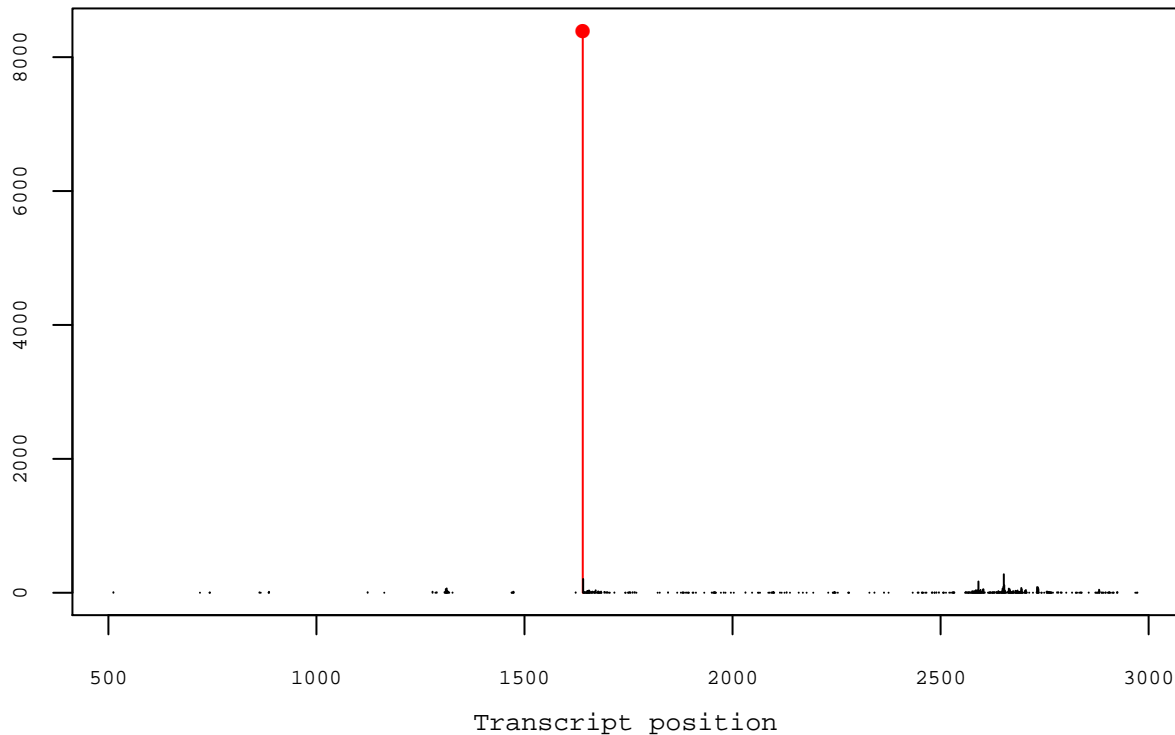


PggD1242_c13_g1_i2

5' GCAGCTAGCATAACAGGGAGCCAGGCAAATTCT 3'
| | | | | | | | | | | | | | |
3' ACCGTATGTCCCTCGGTCCGT 5'



Cleavage site: 1640 Tag abundance: 8392 Weighted abundance: 4196 Category: 0
sRNA abundance: 102 Alignment score: 1 MFE ratio: 0.914 p-value: 0.001