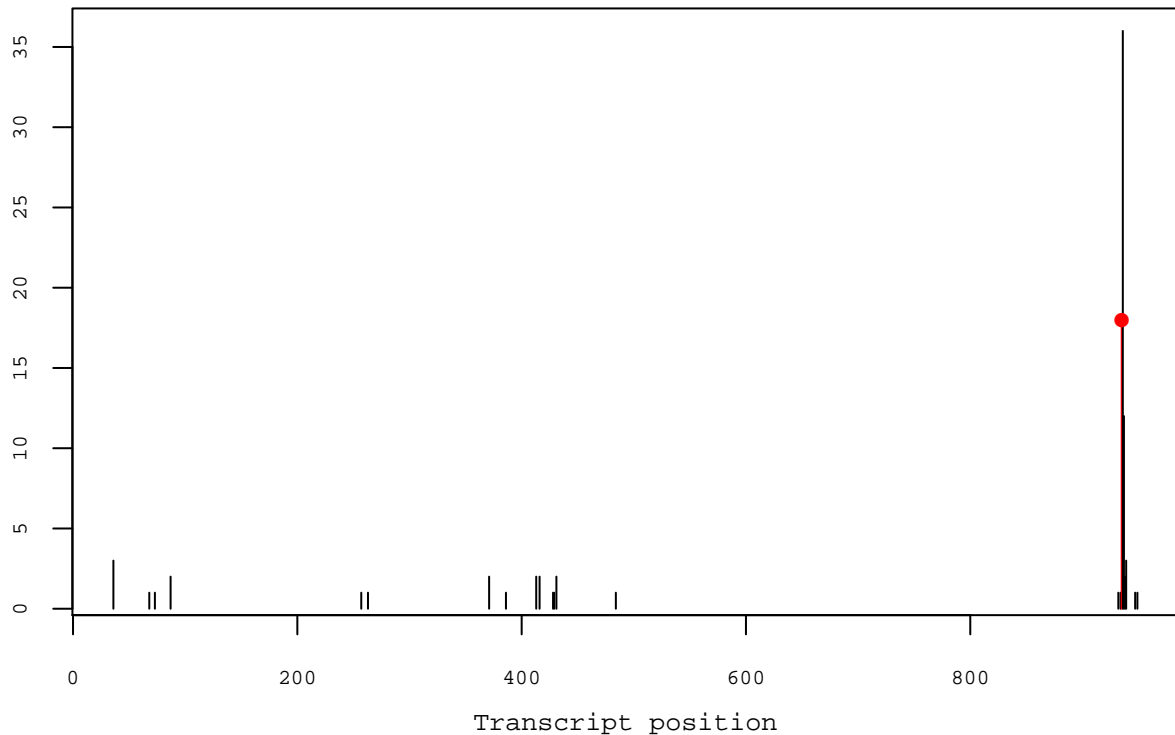


PggD5575_c104_g1_i21

5' AACAGCTTGAGTTCGGCTCGTTAAGAAGCTCG '3
||| |o| ||| |o| ||| |
3' GTCGAGCACAAAGCTGAGCAATTCG '5

Fragment Abundance



Cleavage site: 935 Tag abundance: 18 Weighted abundance: 0.947 Category: 2
sRNA abundance: 20 Alignment score: 3.5 MFE ratio: 0.777 p-value: 0.031