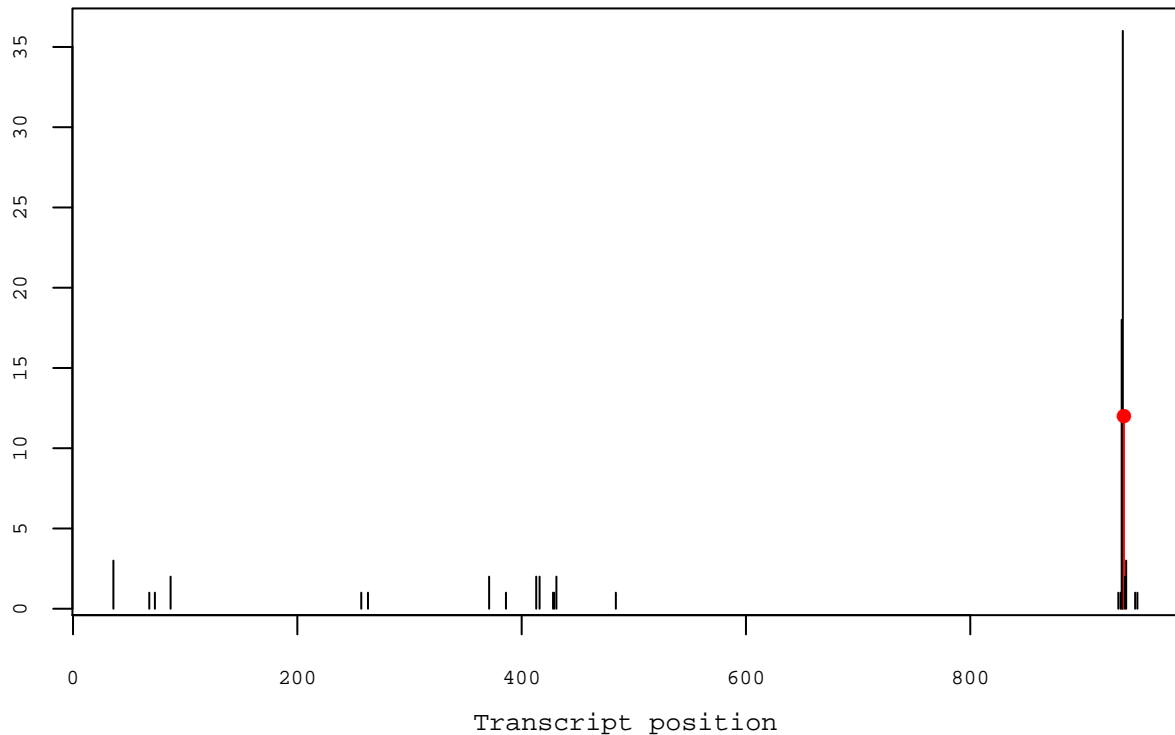


PggD5575\_c104\_g1\_i21

5' CAGCTTGAGTTCGGCTCGTTAAGAAGCTCGTG '3  
|||o||| ||||| ||||| |||||  
3' CGAGCACAAAGCCAAGCAATTCTTC '5

Fragment Abundance



Cleavage site: 937 Tag abundance: 12 Weighted abundance: 0.983 Category: 2  
sRNA abundance: 9 Alignment score: 3.5 MFE ratio: 0.726 p-value: 0.02