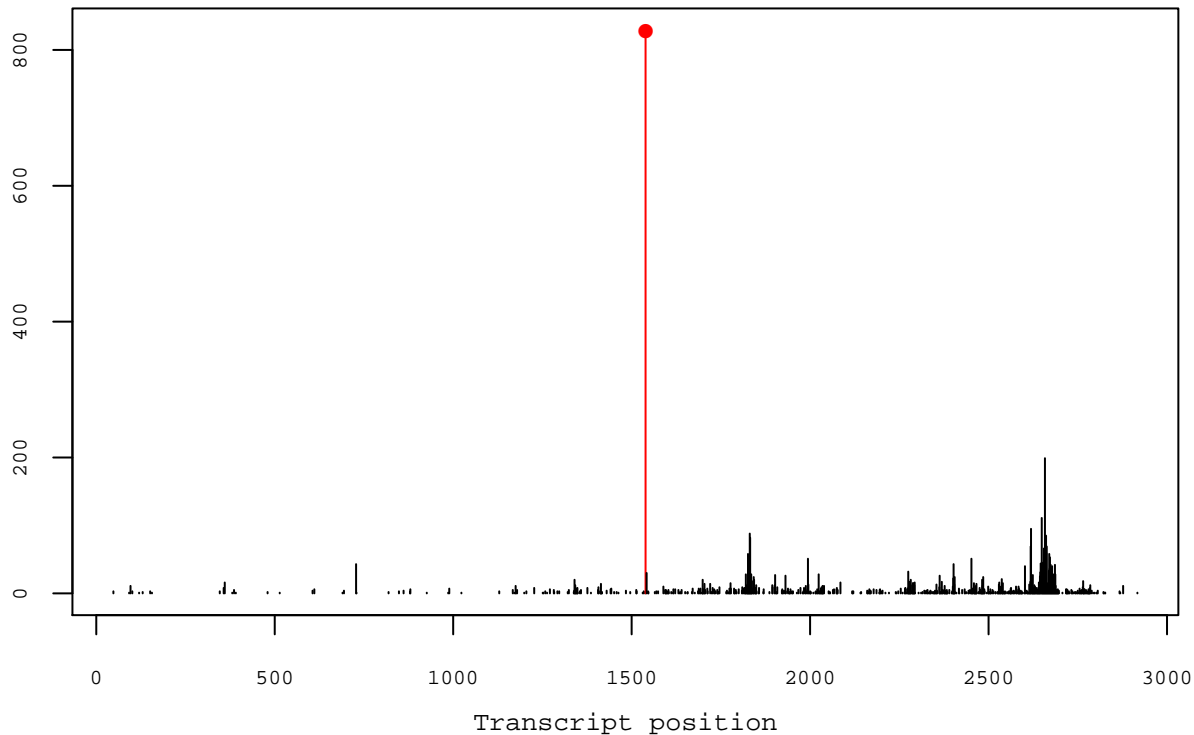


PggD2305\_c3\_g1\_i1

5' GCACATGGGATATTGGCGGGCTCAATCACCA '3  
| | | | | | | | | | | | | | | | | |  
3' CGCTATAACCGTGCCGAGTT '5

Fragment Abundance



Cleavage site: 1539 Tag abundance: 828 Weighted abundance: 770.5 Category: 0  
sRNA abundance: 387 Alignment score: 2 MFE ratio: 0.894 p-value: 0.001