

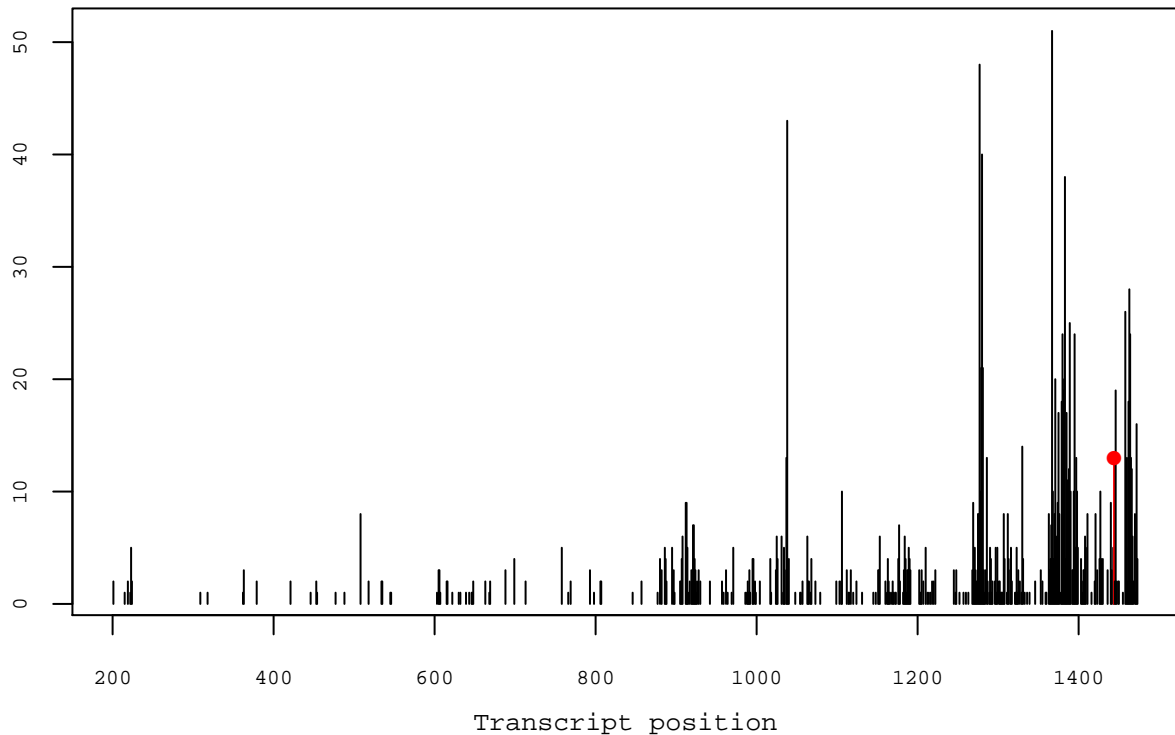
PggD12012\_c1\_g1\_i1

5' TTGTAATA-CGATTCCATGTACTTGCATTGAAG '3

|| | |||||

3' ATCTCAAAGGTACATGAACGT '5

Fragment Abundance



Cleavage site: 1444 Tag abundance: 13 Weighted abundance: 13 Category: 2  
sRNA abundance: 5 Alignment score: 3 MFE ratio: 0.701 p-value: 0.04