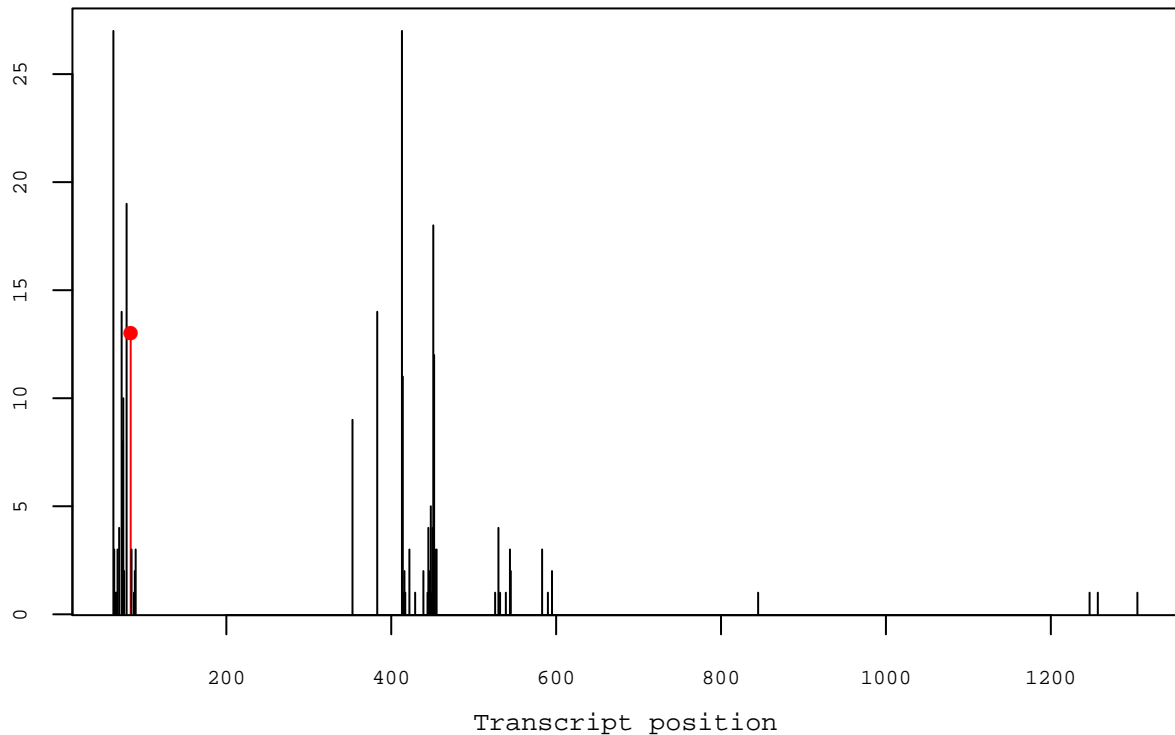


PggD12497\_c103\_g1\_i1

5' TCGGGTGTCCGATCCTGACCGTAGATCACGGT '3  
|||||  
3' CCACAGGCTAGGACTGGCATTTA '5

Fragment Abundance



Cleavage site: 84 Tag abundance: 13 Weighted abundance: 6.333 Category: 2  
sRNA abundance: 8 Alignment score: 1 MFE ratio: 1 p-value: 0.028