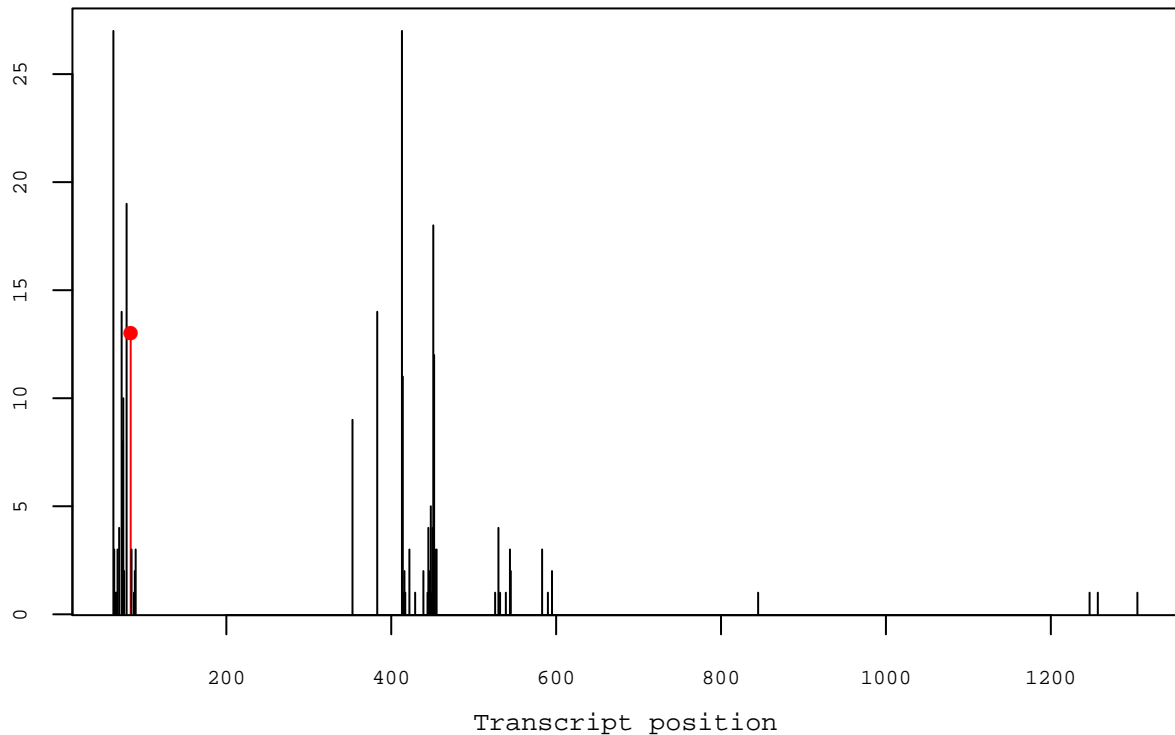


PggD12497\_c103\_g1\_i1

5' TGCGGTGTCCGATCCTGACCGTAGATCACGGT '3  
|||||  
3' GCCACAGGCTAAGACTGGCATCTA '5

Fragment Abundance



Cleavage site: 84

sRNA abundance: 7

Tag abundance: 13

Alignment score: 2

Weighted abundance: 6.333

MFE ratio: 0.902

Category: 2

p-value: 0.028