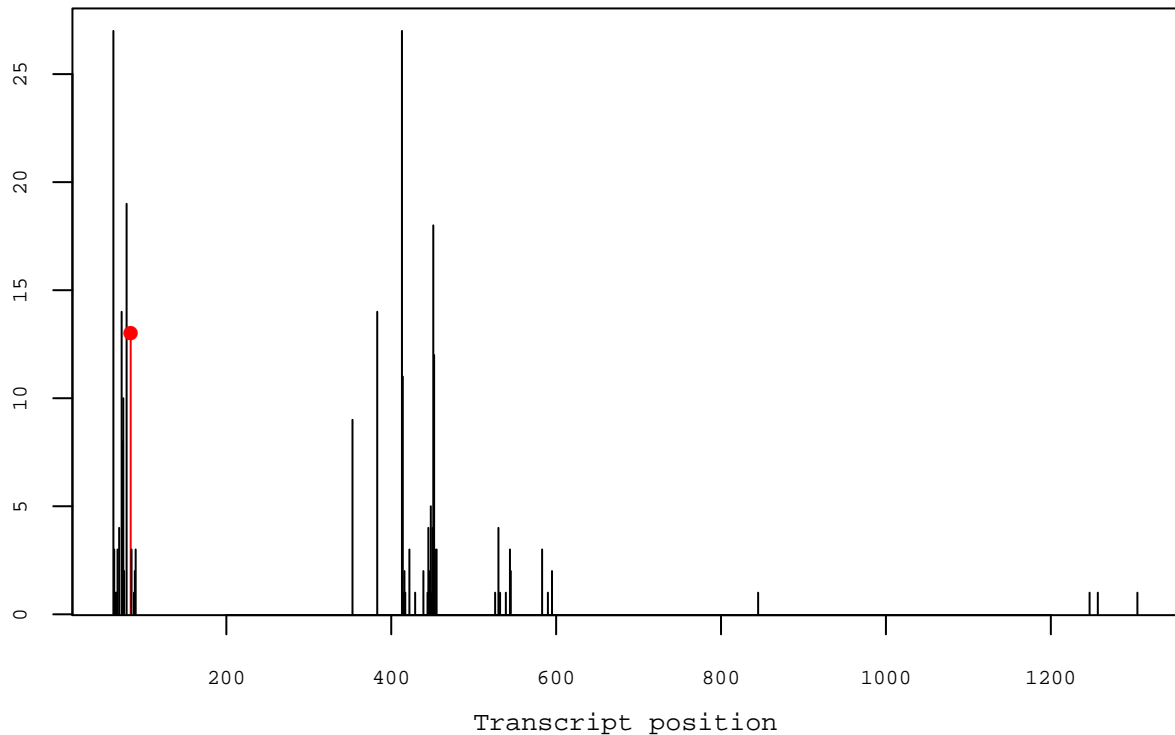


PggD12497_c103_g1_i1

5' TGCGGTGTCCGATCCTGACCGTAGATCACGGT '3
|||||
3' GCCACAGGCTAGGACTGGCATTTA '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