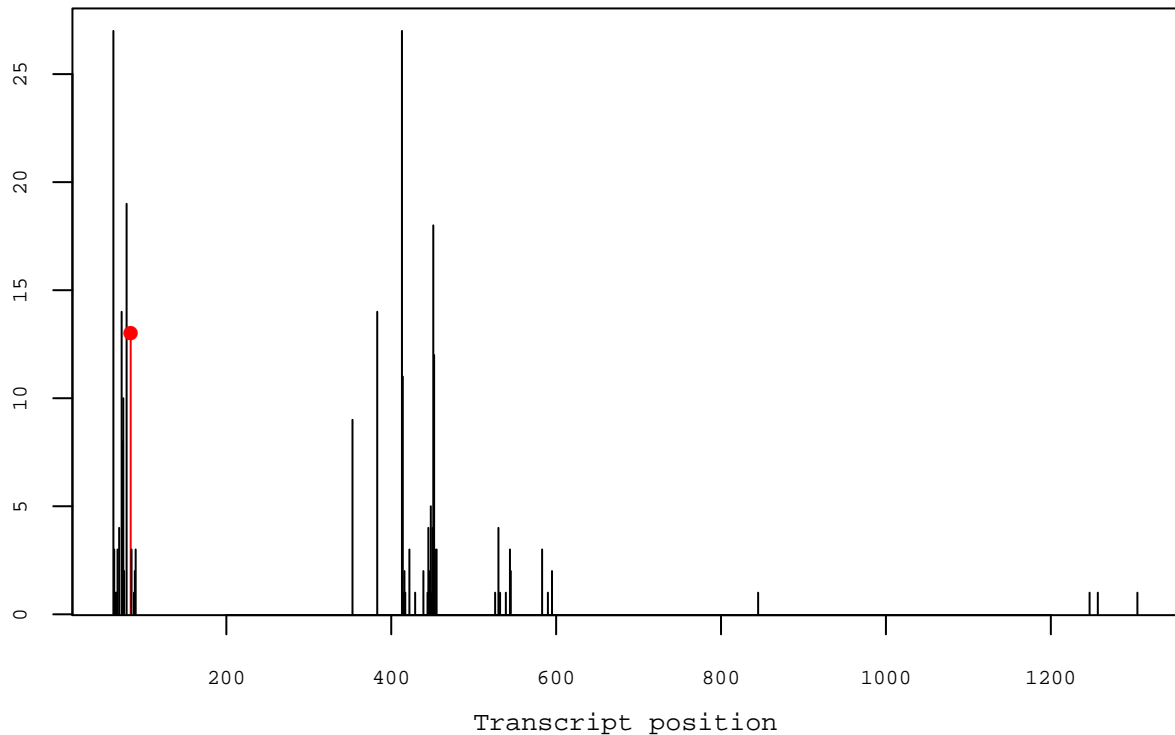


PggD12497_c103_g1_i1

5' TGCGGTGTCCGATCCTGACCGTAGATCACGGT '3
||||| ||||| ||||| o ||
3' GCCACAGACTAGGCCTGGCATTTA '5

Fragment Abundance



Cleavage site: 84 Tag abundance: 13 Weighted abundance: 6.333 Category: 2
sRNA abundance: 16 Alignment score: 4 MFE ratio: 0.764 p-value: 0.028