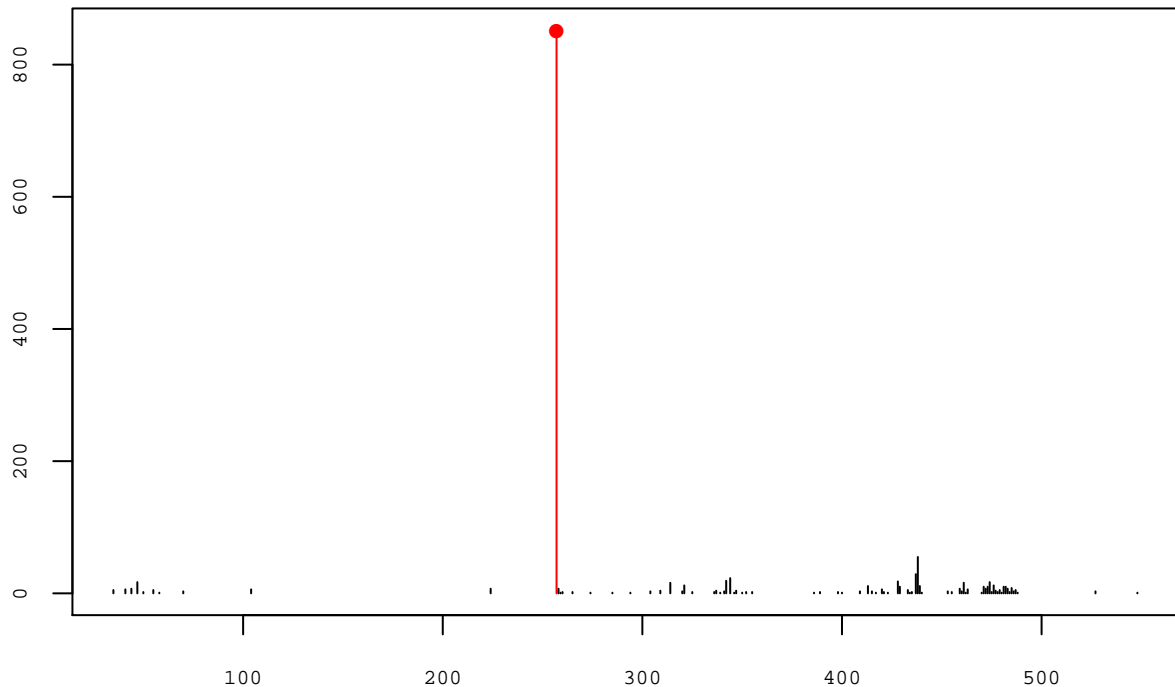


PggD10737_c261_g1_i1

5' GAGAGATCACGTGCCCTGCTTCTCCACATCCG '3

3' ACGTGCACGGGACGAAGAGGT 5'



Transcript position

Cleavage site: 257 Tag abundance: 851 Weighted abundance: 425.5 Category: 0

sRNA abundance: 235 Alignment score: 2 MFE ratio: 0.898 p-value: 0.009