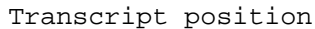


PggD1457_c16_g2_i2

5' GCTTGAGCAGGTGCCCTGCTTCTCCAGCTTTT '3

3' ||| ||||| ||||| ||||| |||||

ACGTGCACGGGACGAAGAGGT



Cleavage site: 664 Tag abundance: 133 Weighted abundance: 66.5 Category: 0
sRNA abundance: 235 Alignment score: 2 MFE ratio: 0.926 p-value: 0.005