

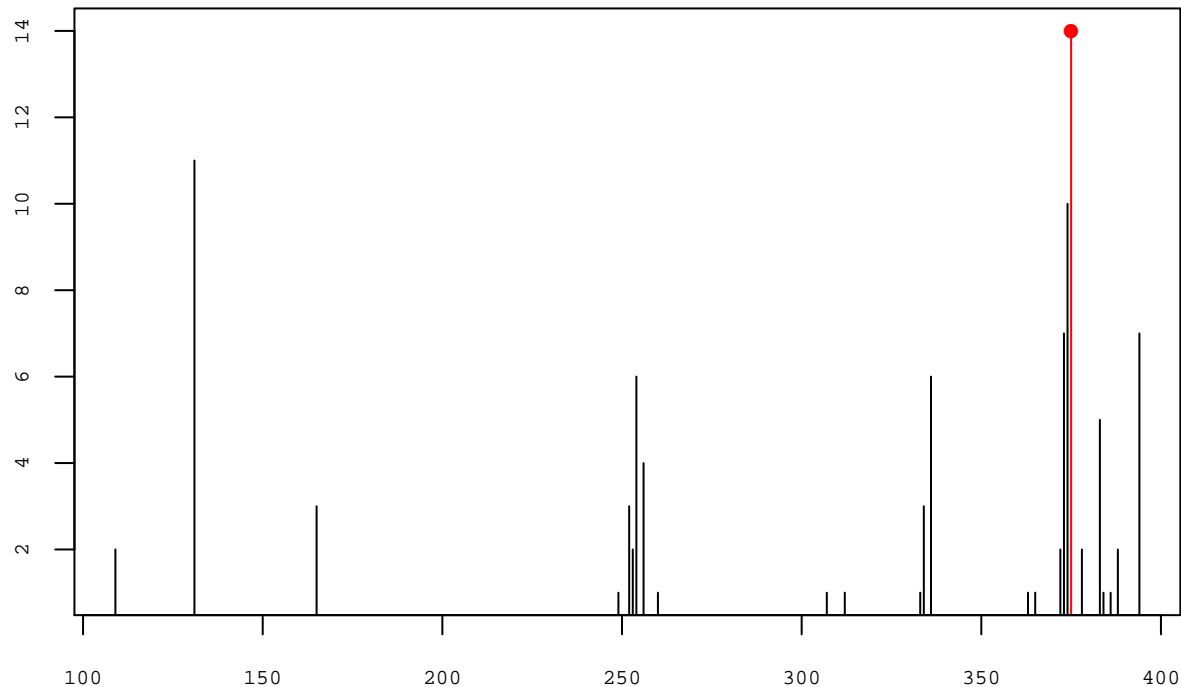
PggD9781_c122_g1_i1

5' AAGTCTTCTAAGTCATTCCAAAATTATCTTT '3

o |||||||||||||||||||

3' TAGAAGATTTCAGTAAGGTTTTAAA '5

Fragment Abundance



Transcript position

Cleavage site: 375 Tag abundance: 14 Weighted abundance: 0.406 Category: 0

sRNA abundance: 29 Alignment score: 0.5 MFE ratio: 1 p-value: 0.05