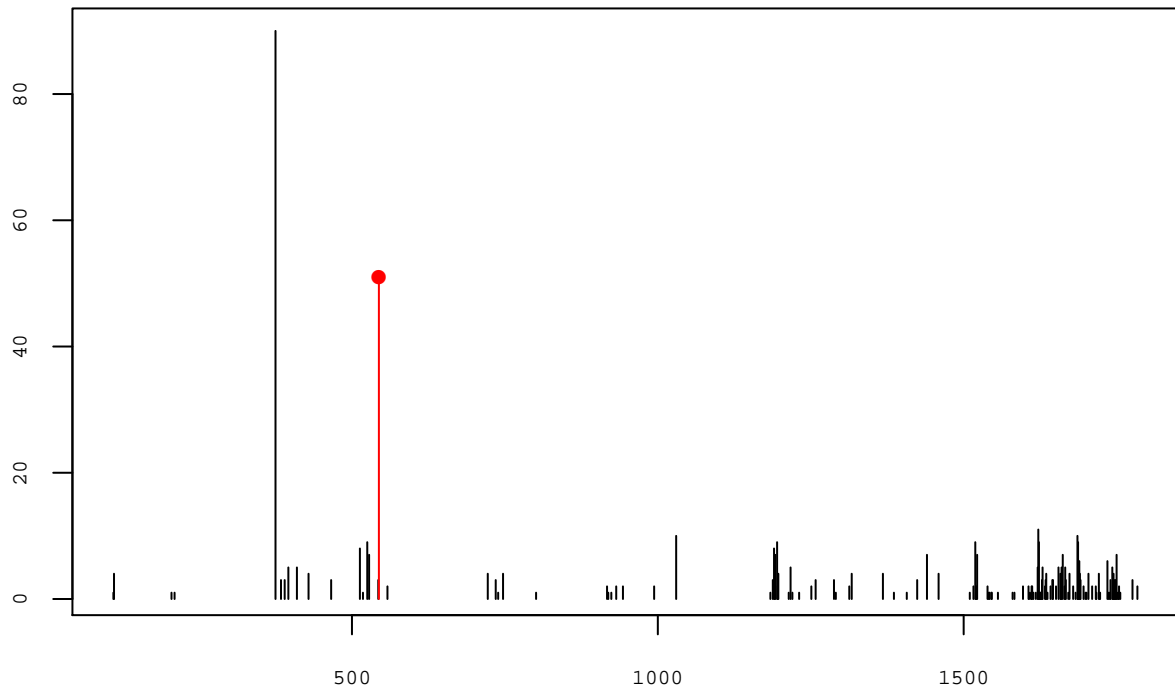


PggD7691_c15_g1_i1

5' TCCCTCTCGATATTGGCTCGGCTCAATTCCT 3'
|||||
3' CGCTATAACCGTGCCGAGTT 5'

Fragment Abundance



Transcript position

Cleavage site: 544	Tag abundance: 51	Weighted abundance: 51	Category: 2
sRNA abundance: 387	Alignment score: 3	MFE ratio: 0.812	p-value: 0.04