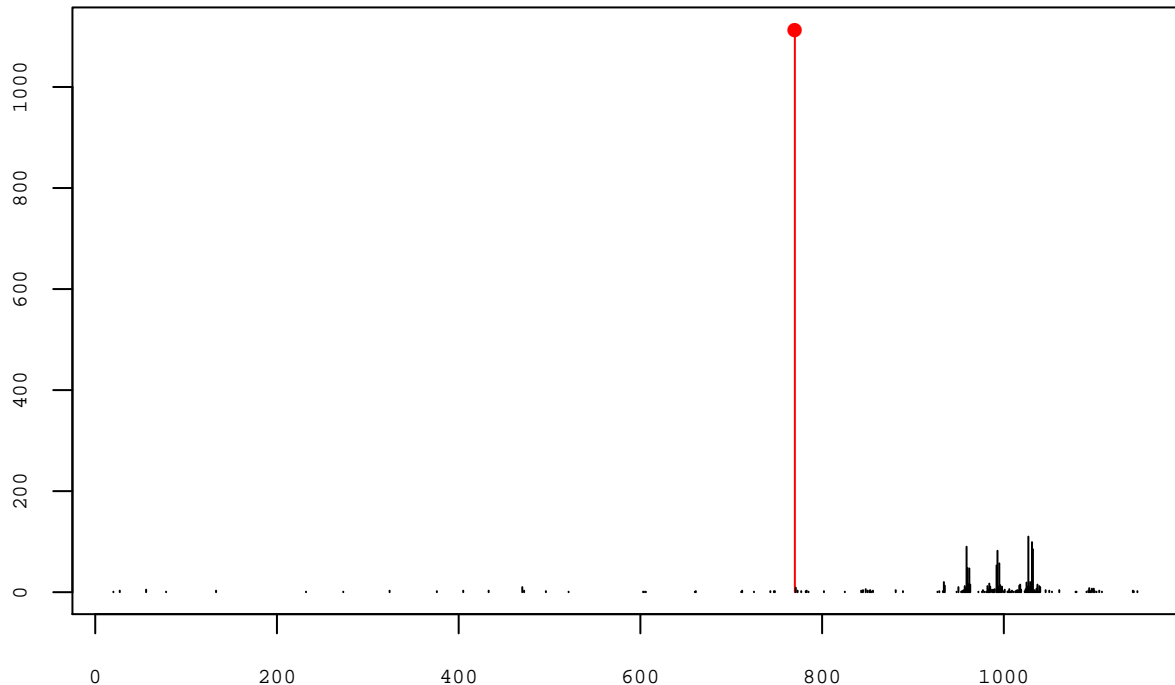


PggD1928_c0_g1_i2

5' AAGGCATTACGTGCCCTGCTTCTCCACAGTGG 3'
o |||||
3' GTGCACGGGACGAAGAGGT 5'

Fragment Abundance



Transcript position

Cleavage site: 770	Tag abundance: 1113	Weighted abundance: 556.5	Category: 0
sRNA abundance: 17	Alignment score: 0.5	MFE ratio: 0.959	p-value: 0.004