

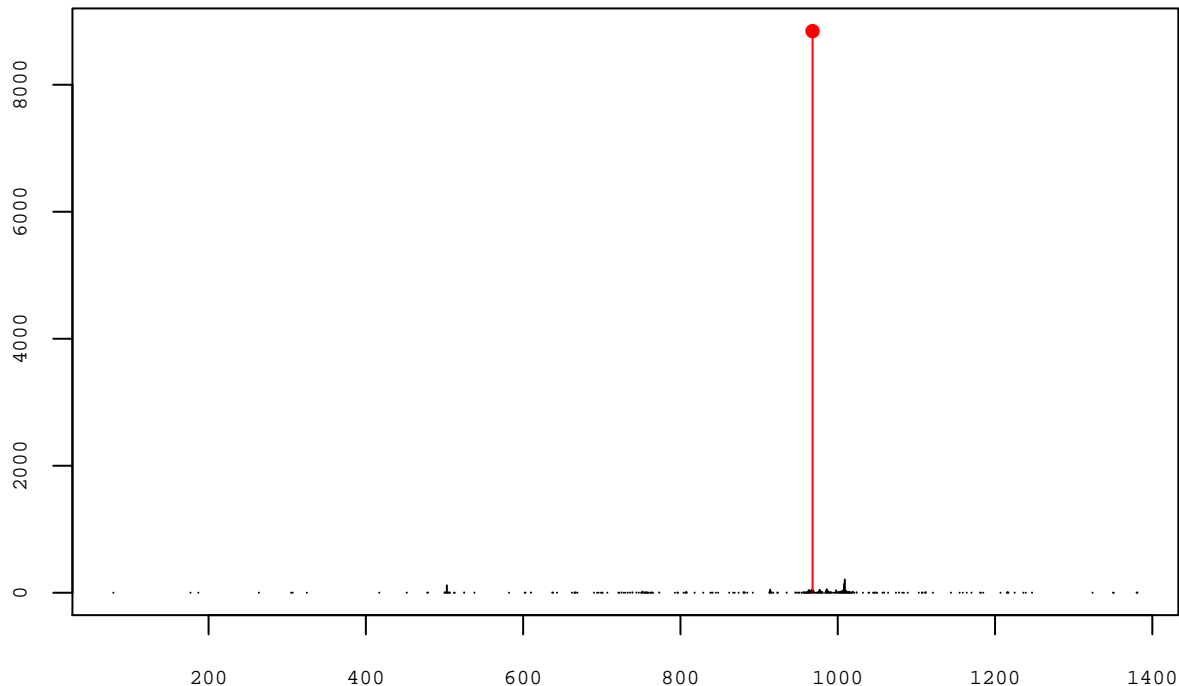
PggD14469\_c62\_g1\_i2

5' TTCCCCTTGCTATCCCTCCTGAGCTGAGCTAG '3

o |||||

3' CCGCGATAGGGAGGACTCGAA '5

Fragment Abundance



Transcript position

Cleavage site: 968	Tag abundance: 8848	Weighted abundance: 8848	Category: 0
srRNA abundance: 712	Alignment score: 3.5	MFE ratio: 0.835	p-value: 0.004