

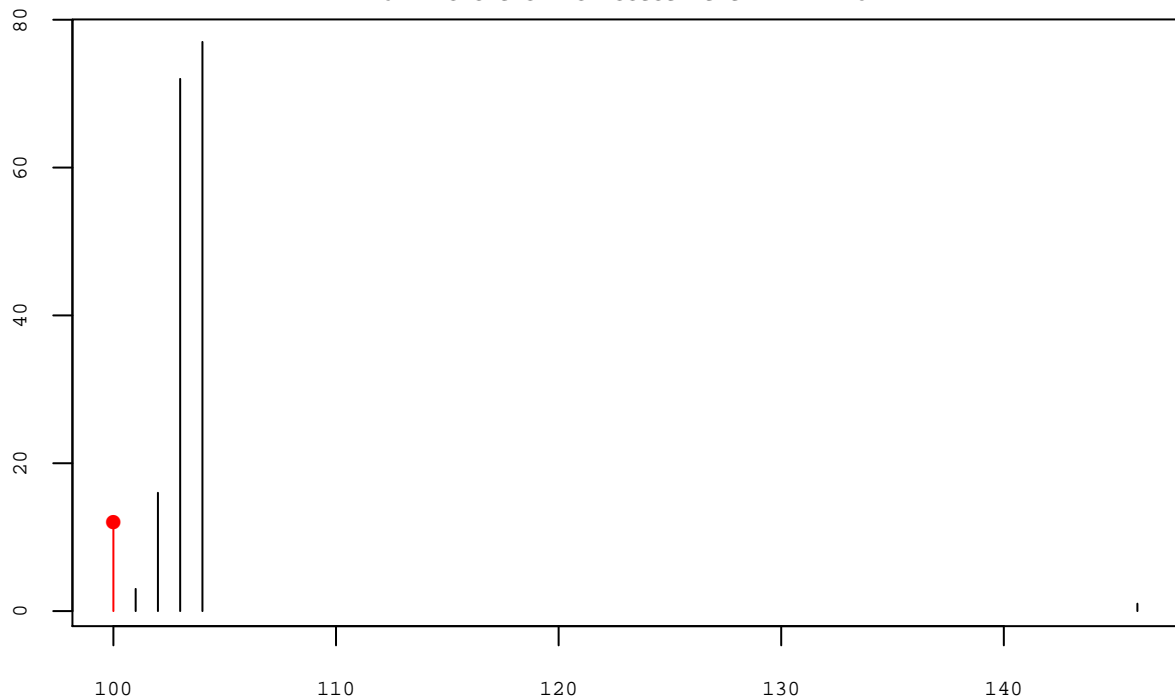
PggD128\_c121\_g1\_i1

5' GCTGCTGTCATACAGTCGCGATGTGCGGCTGT '3

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3' GTGACAGTATGTTGGCGCTACACA '5

Fragment Abundance



Transcript position

Cleavage site: 100

Tag abundance: 12

Weighted abundance: 3.917

Category: 3

sRNA abundance: 31

Alignment score: 4

MFE ratio: 0.825

p-value: 0.024