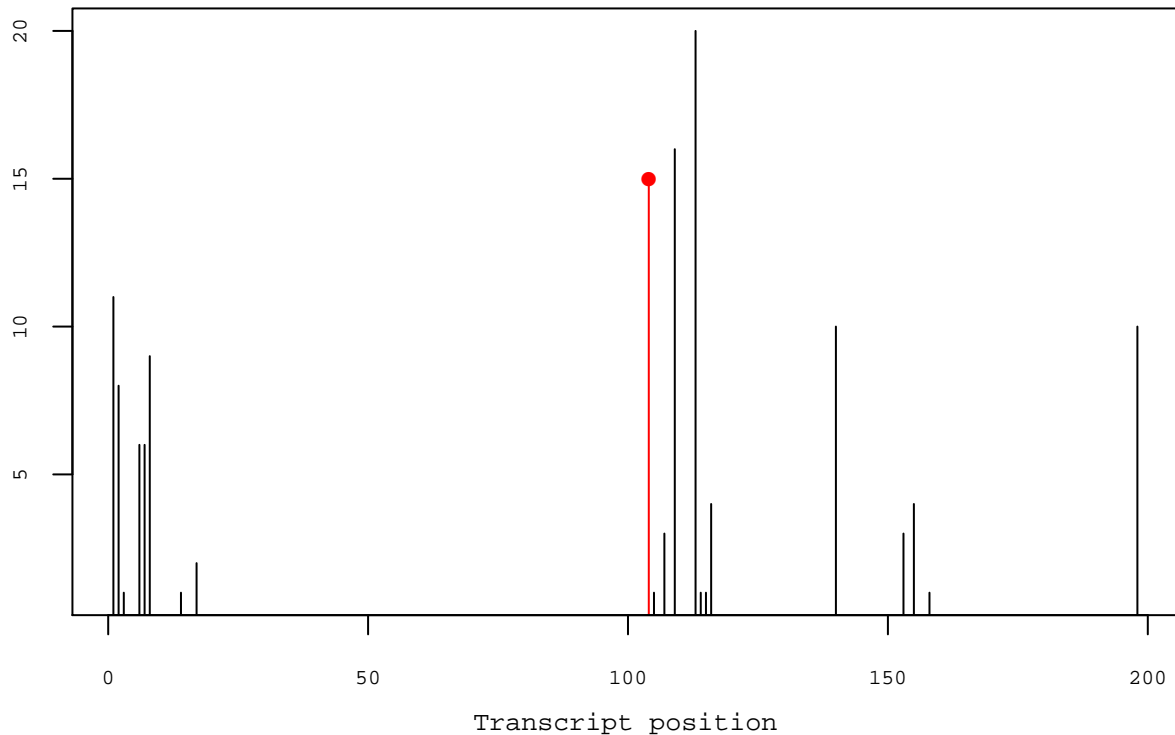


PggD15357_c29_g1_i1

5' CCTTAA-GTCCATGTCATTTTGCTTTAGTTATG '3
| | | | | | | | | | | | | | | | | |
3' TGCAGGTACAGTAAAAGGAGAT '5

Fragment Abundance



Cleavage site: 104

Tag abundance: 15

Weighted abundance: 0.061

Category: 2

sRNA abundance: 5

Alignment score: 4

MFE ratio: 0.739

p-value: 0.019