

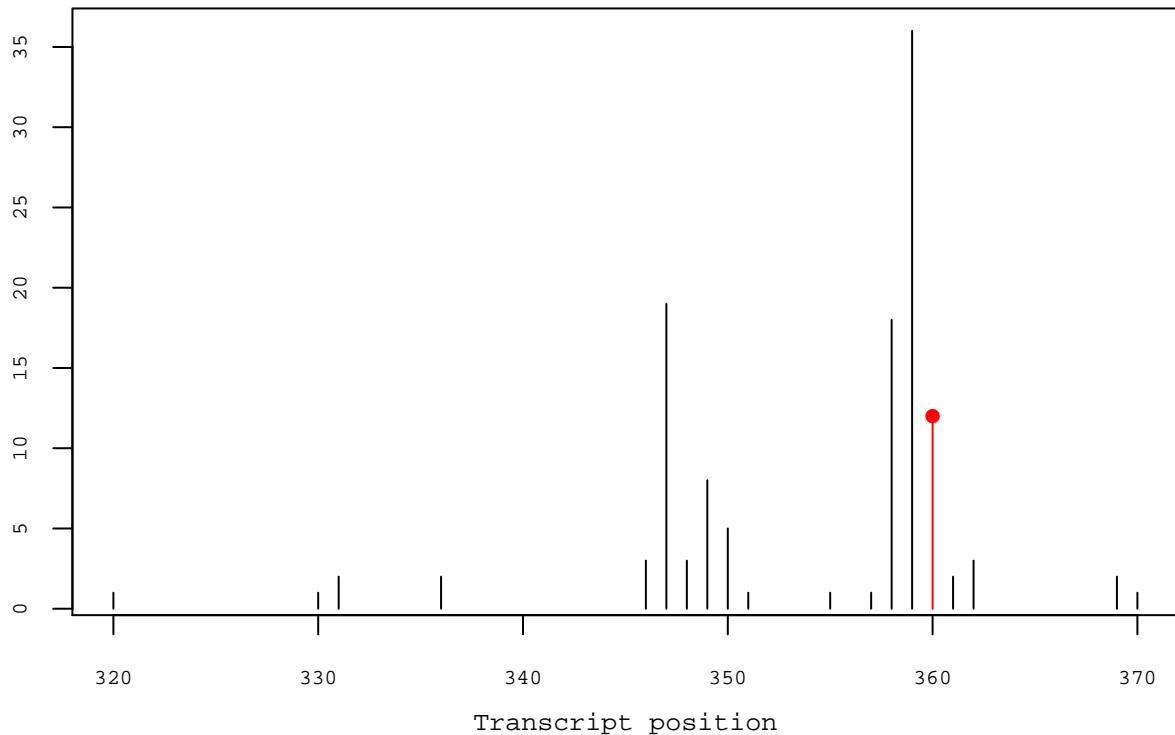
PggD15562_c117_g1_i1

5' CAGCTCGTGTTCGGCTCGTTAAG-AAGCTCGTG '3

o|||||||o||||o|||||

3' TGAGCACAAGTCGAGCGATTCGTT '5

Fragment Abundance



Cleavage site: 360

Tag abundance: 12

Weighted abundance: 0.983

Category: 2

sRNA abundance: 18

Alignment score: 4

MFE ratio: 0.827

p-value: 0.04