

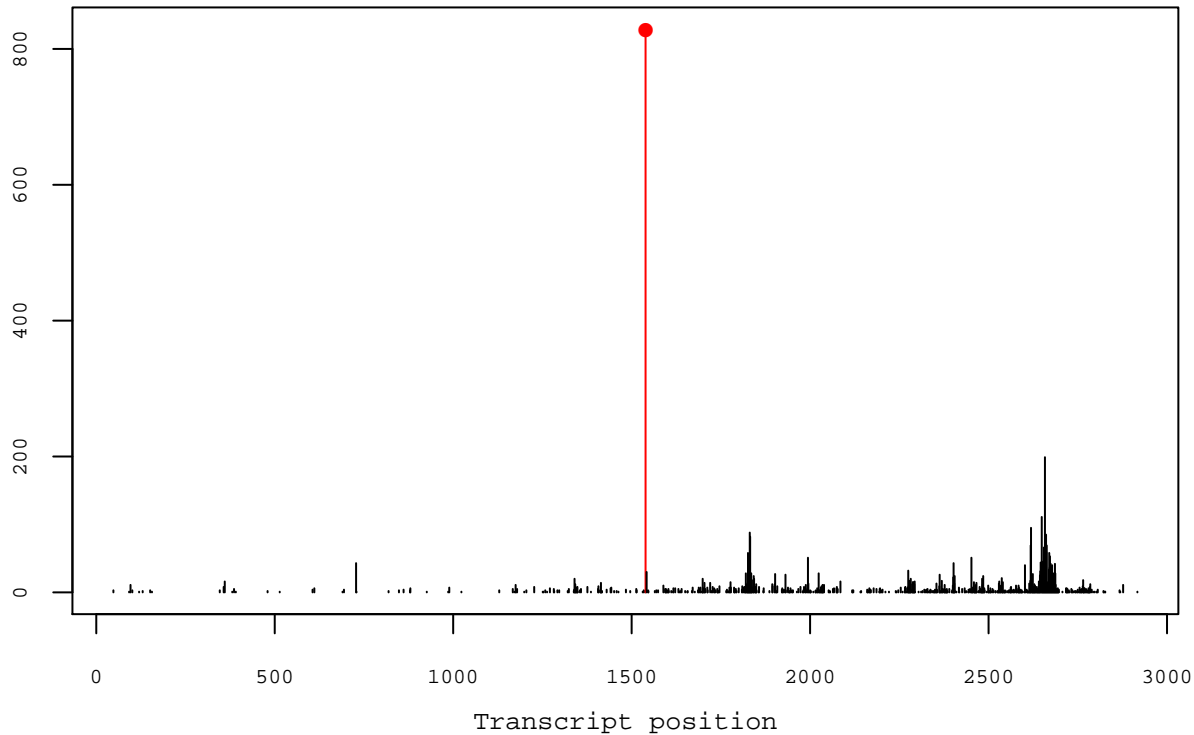
PggD2305\_c3\_g1\_i1

5' GCACATGG-GATATTGGCGCGGCTCAATCACCA '3

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

3' CCTCTATAACTGCGCCGAGTT '5

Fragment Abundance



Cleavage site: 1539

Tag abundance: 828

Weighted abundance: 770.5

Category: 0

sRNA abundance: 102

Alignment score: 2

MFE ratio: 0.857

p-value: 0.001