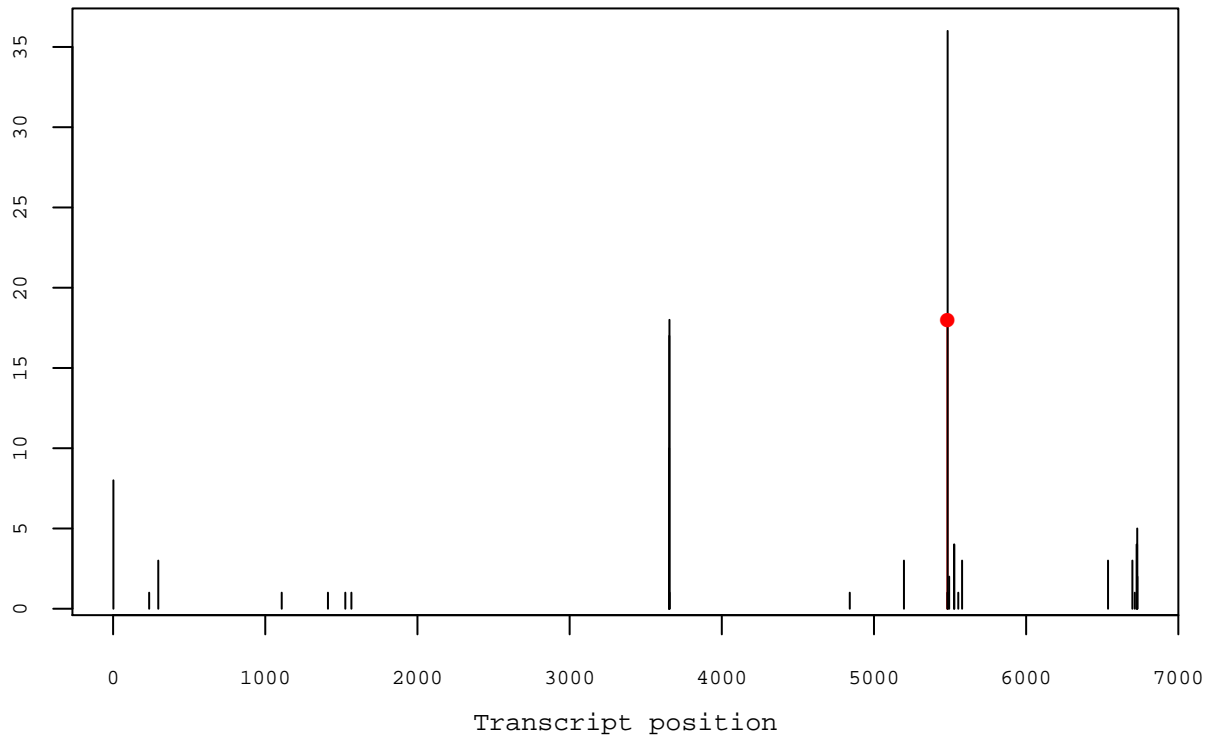


PggD15546_c13_g1_i2

5' AACAGCTCGTGTTCCGCTCGTTAAGAAGCTCG '3

[illegible]

3' GTCGAGCACAAGCTGAGCAATTCG 5'



Cleavage site: 5483

Tag abundance: 18

Weighted abundance: 0.947

Category: 2

srna abundance: 20

Alignment score: 4

MFE ratio: 0.815

p-value: 0.003