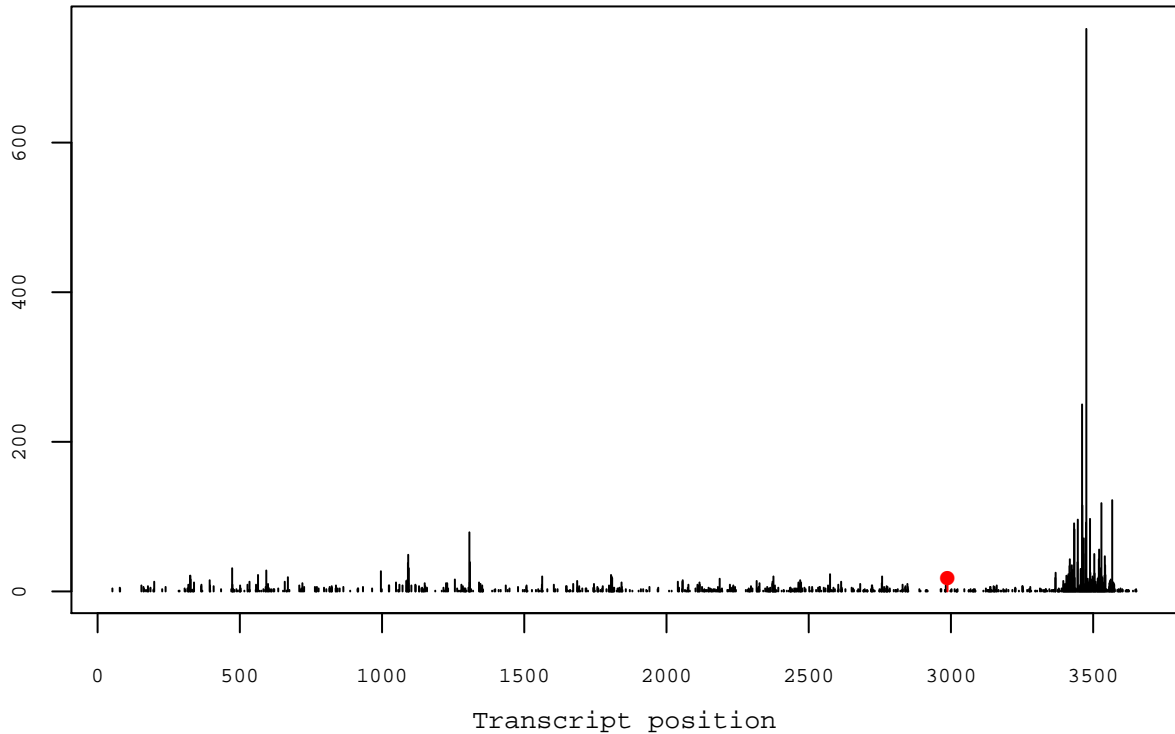


PggD11872_c0_g1_i2

5' CCTCTTAGATCAGGCTGGCAGCTT-GTATTTGT '3

3' ATCTAGTACGACCGTCGAAGT

'5



Cleavage site: 2988	Tag abundance: 18	Weighted abundance: 18	Category: 2
sRNA abundance: 766	Alignment score: 2.5	MFE ratio: 0.789	p-value: 0.035