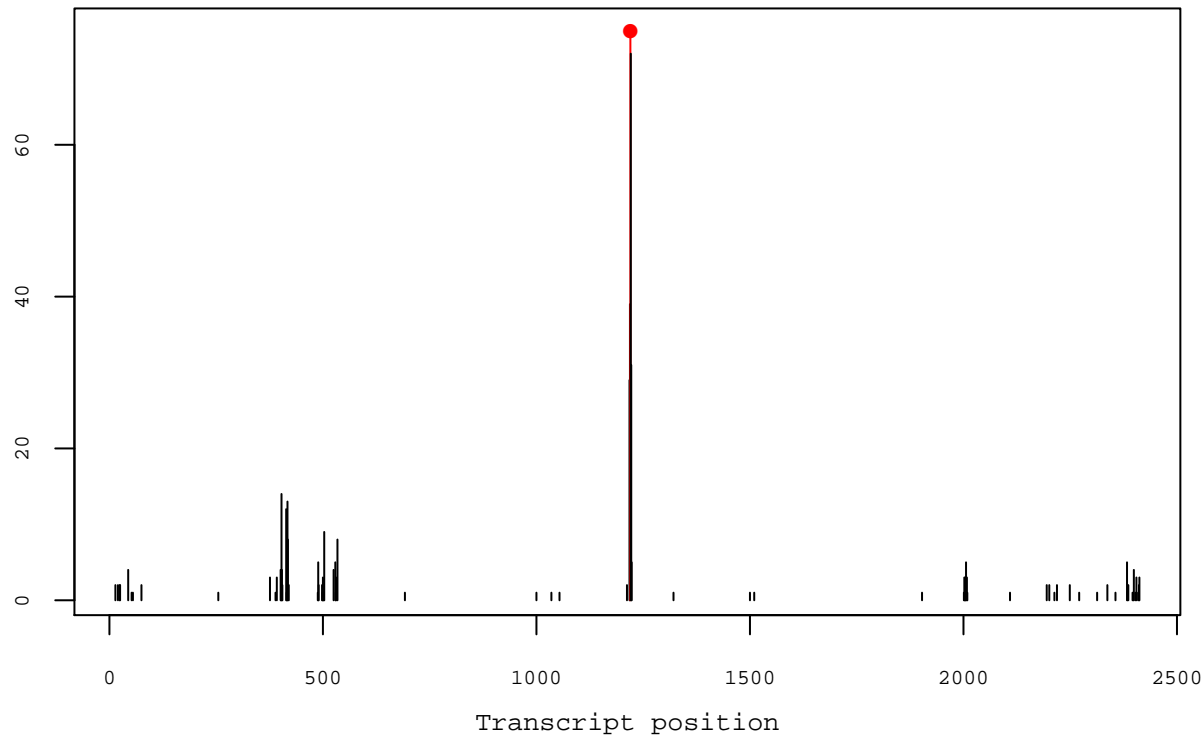


PggD4164_c31_g1_i2

5' ACACCACCCAAATCTTATGTATTTACATGTAT '3
||| ||| ||| |o||| ||| |o|||
3' TGGTGGGTTTAGGATACATAGATG '5

Fragment Abundance



Cleavage site: 1220 Tag abundance: 75 Weighted abundance: 25 Category: 0
sRNA abundance: 8 Alignment score: 2 MFE ratio: 0.873 p-value: 0.015