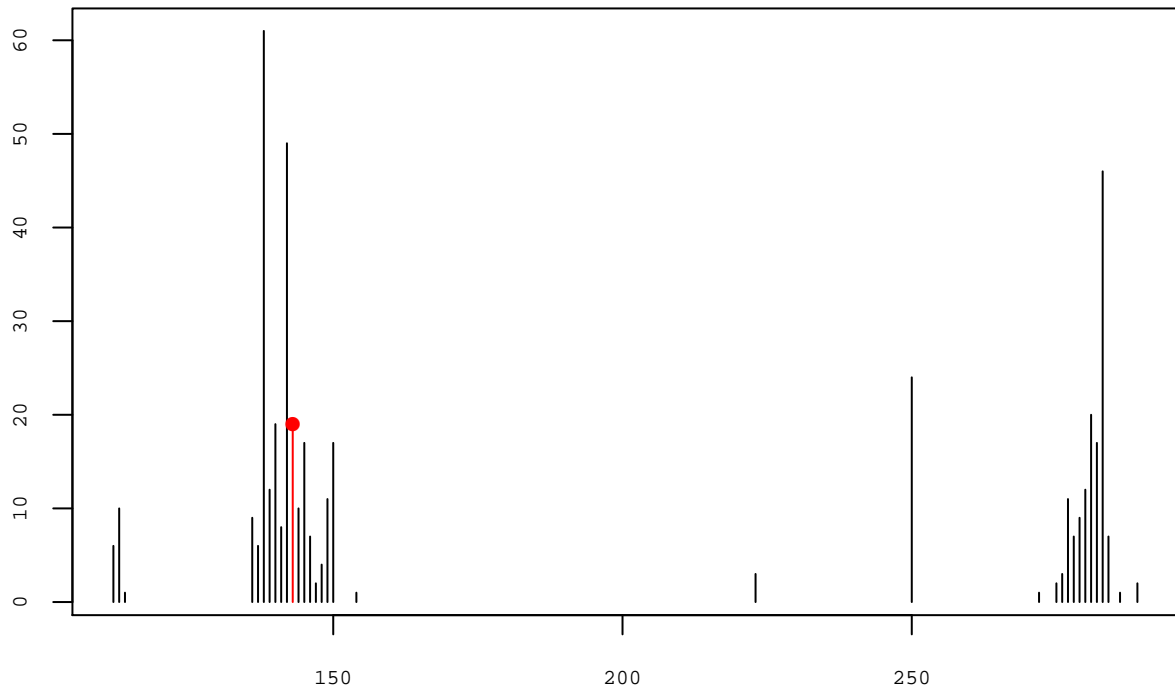


PggD13902_c79_g1_i1

5' CAATTCTCATTATTTACATATCTAACACCTA '3
|||||||o|||||o|||||
3' AGAGTAAATGAATGTGTAGATT '5

Fragment Abundance



Cleavage site: 143 Tag abundance: 19 Weighted abundance: 0.056 Category: 2
sRNA abundance: 12 Alignment score: 2 MFE ratio: 0.865 p-value: 0.014