

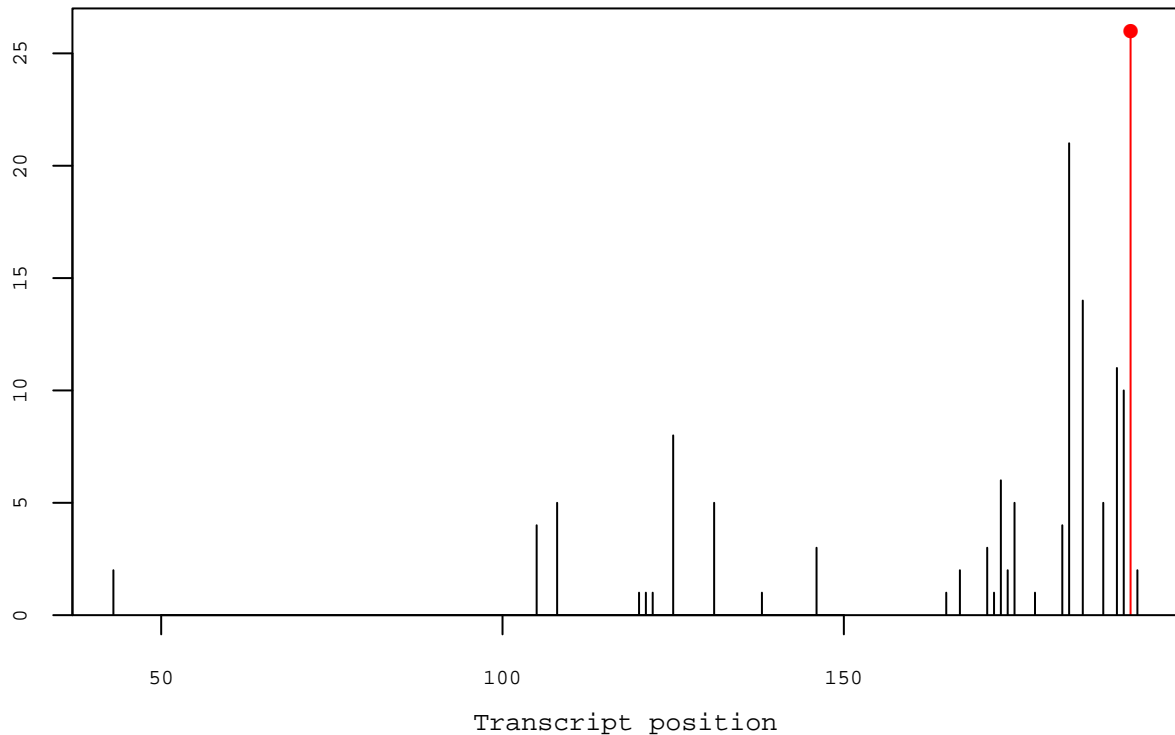
PggD9980_c244_g1_i1

5' GAACACGAGCTGTTACGAACAAA-GAGTCCTT '3

|||||

3' TGTGCTCGACAAGTGCTTGTAT '5

Fragment Abundance



Cleavage site: 192 Tag abundance: 26 Weighted abundance: 4.143 Category: 0
sRNA abundance: 23 Alignment score: 1.5 MFE ratio: 0.955 p-value: 0.025