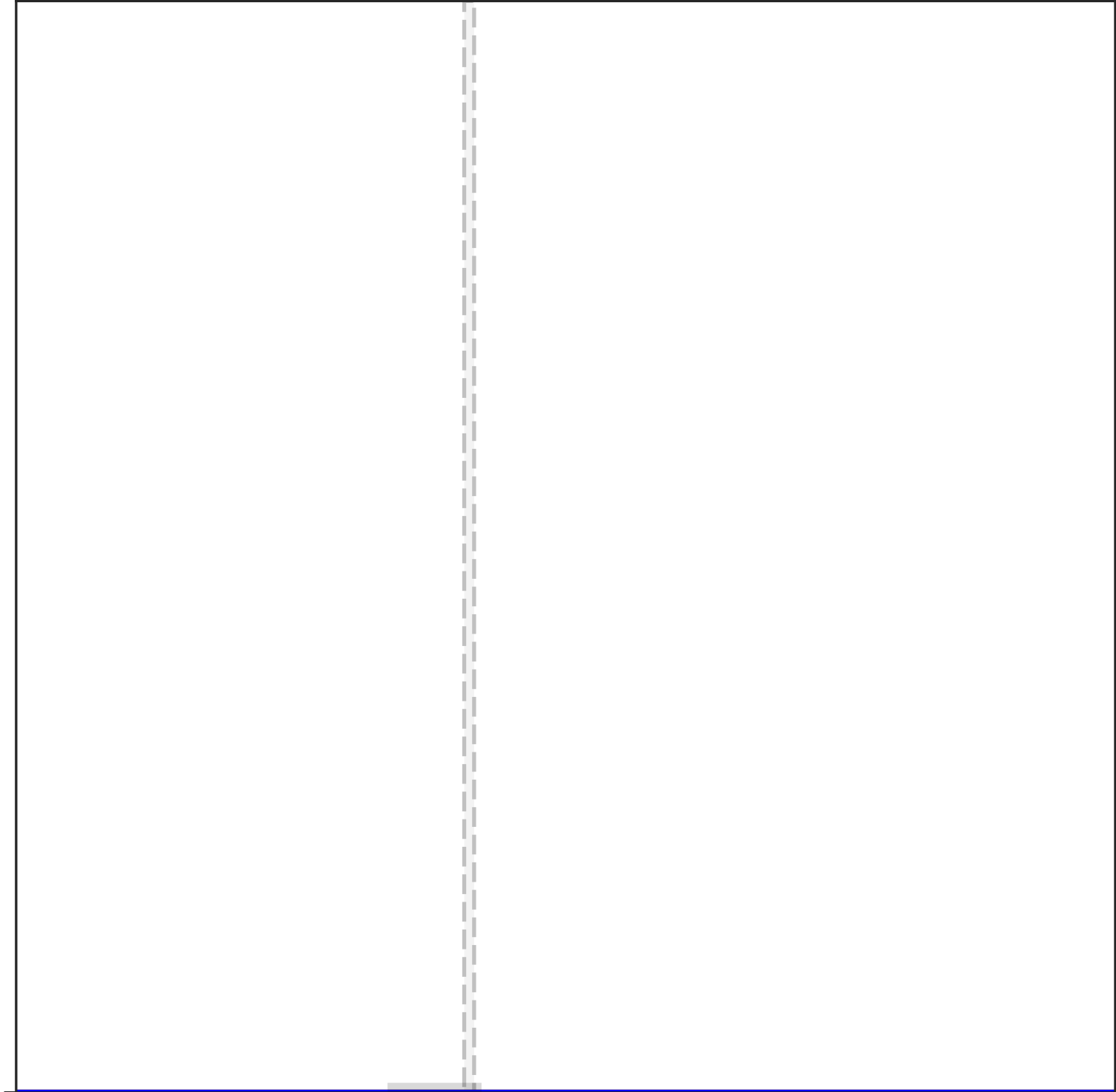


Mutation position distribution

Sequences: % Total (no.)

0.0% (0)



0 35 70 105 140 175 210

Reference amplicon position (bp)



Quantification window



Insertions



Deletions



Substitutions



sgRNA