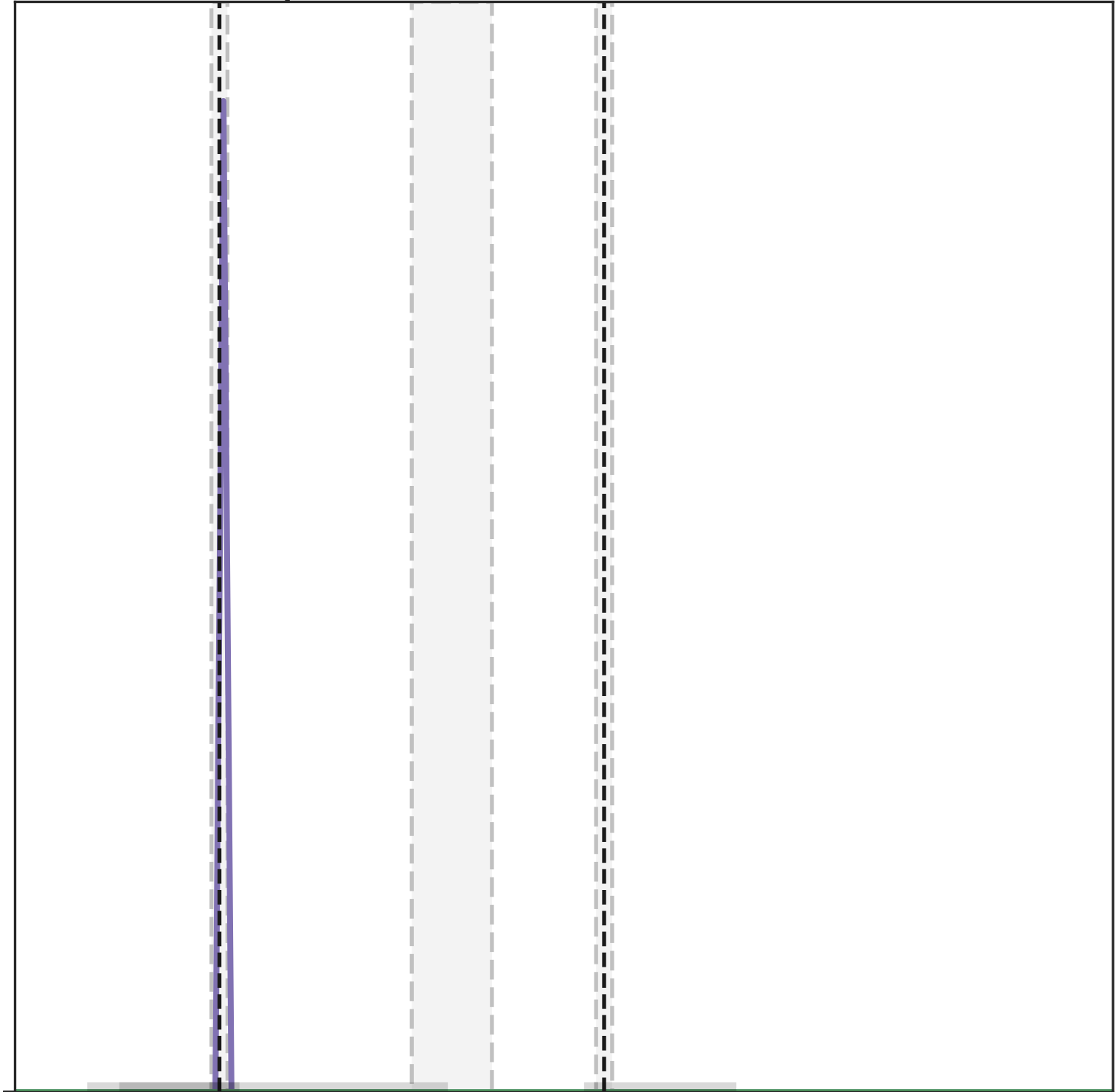


Mutation position distribution: Reference

Sequences: % Total ( % Reference, no. )

0.0% (0.0% , 0)



0 20 40 60 80 100 120

Reference amplicon position (bp)

- Quantification window
- Insertions
- Deletions
- Substitutions
- Predicted cleavage position
- sgRNA