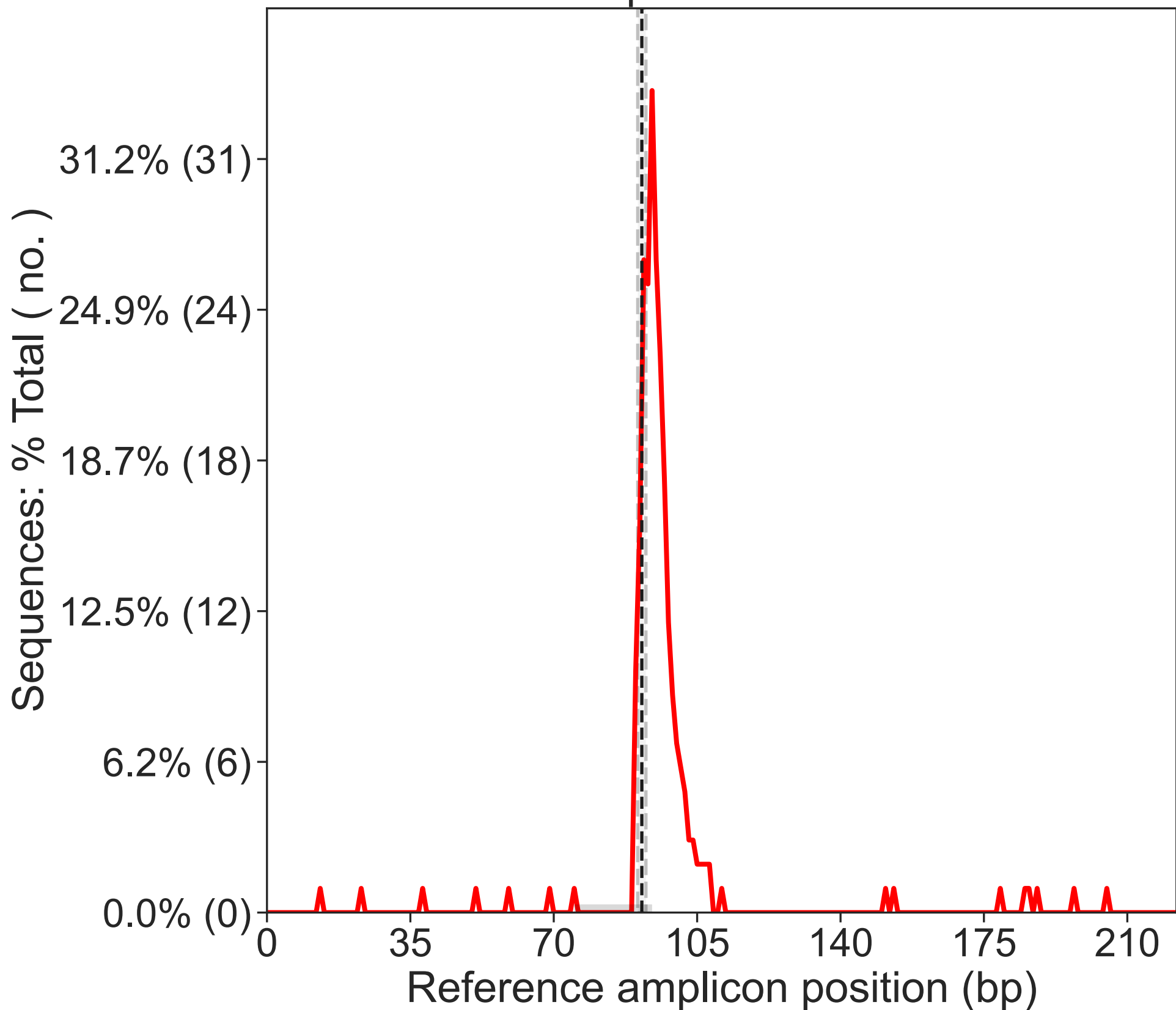


# Mutation position distribution



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