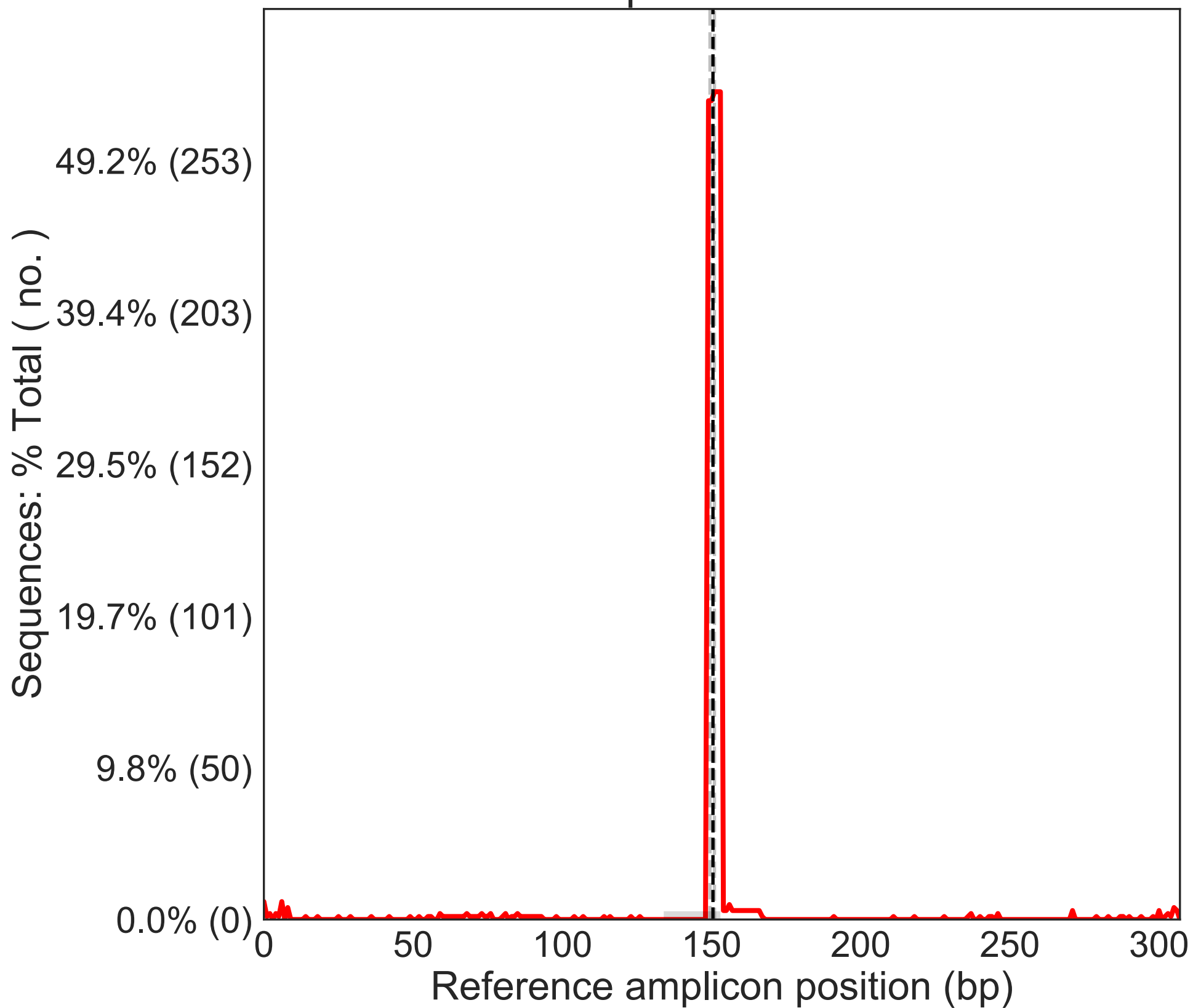


# Mutation position distribution



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