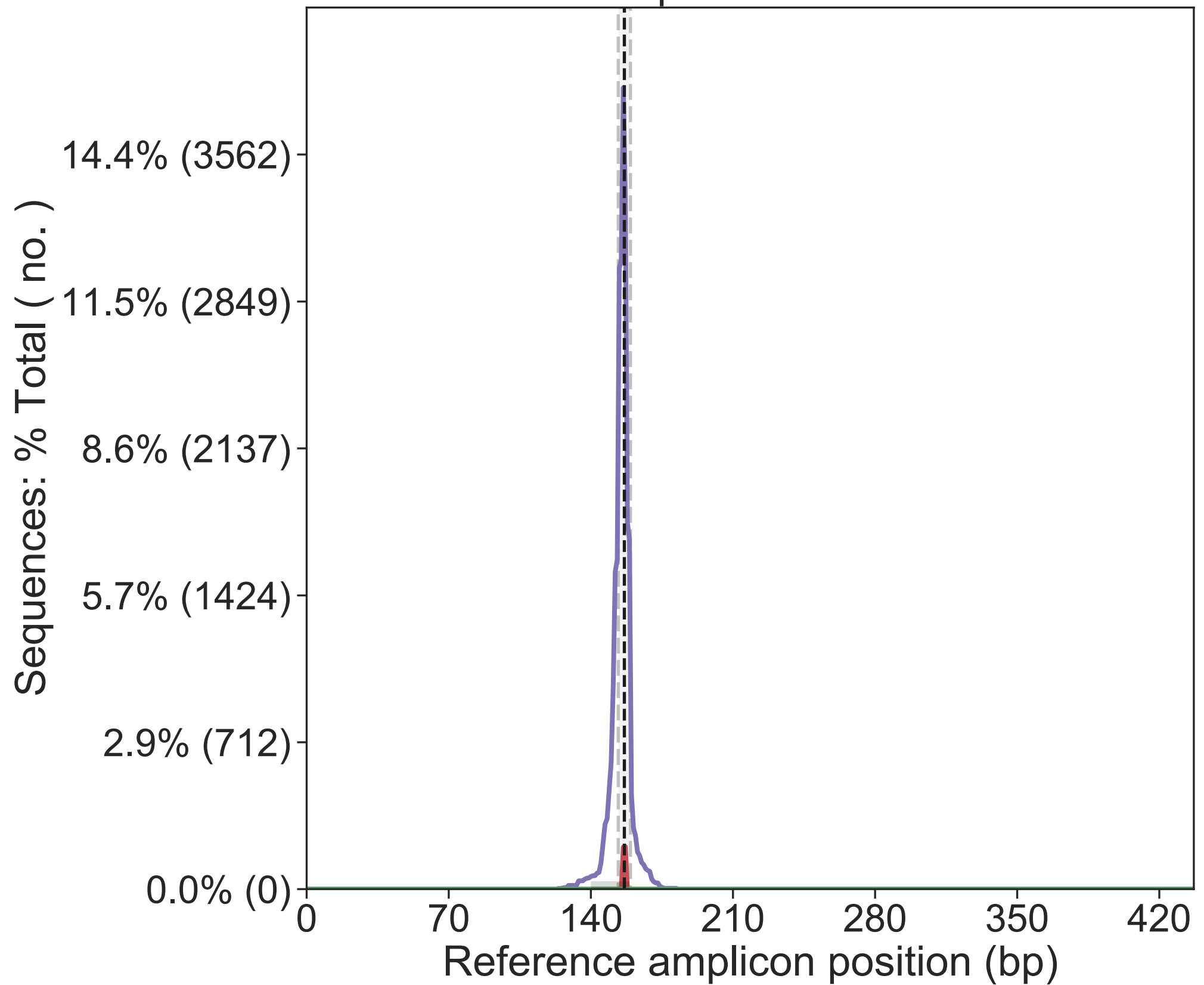


# Mutation position distribution



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