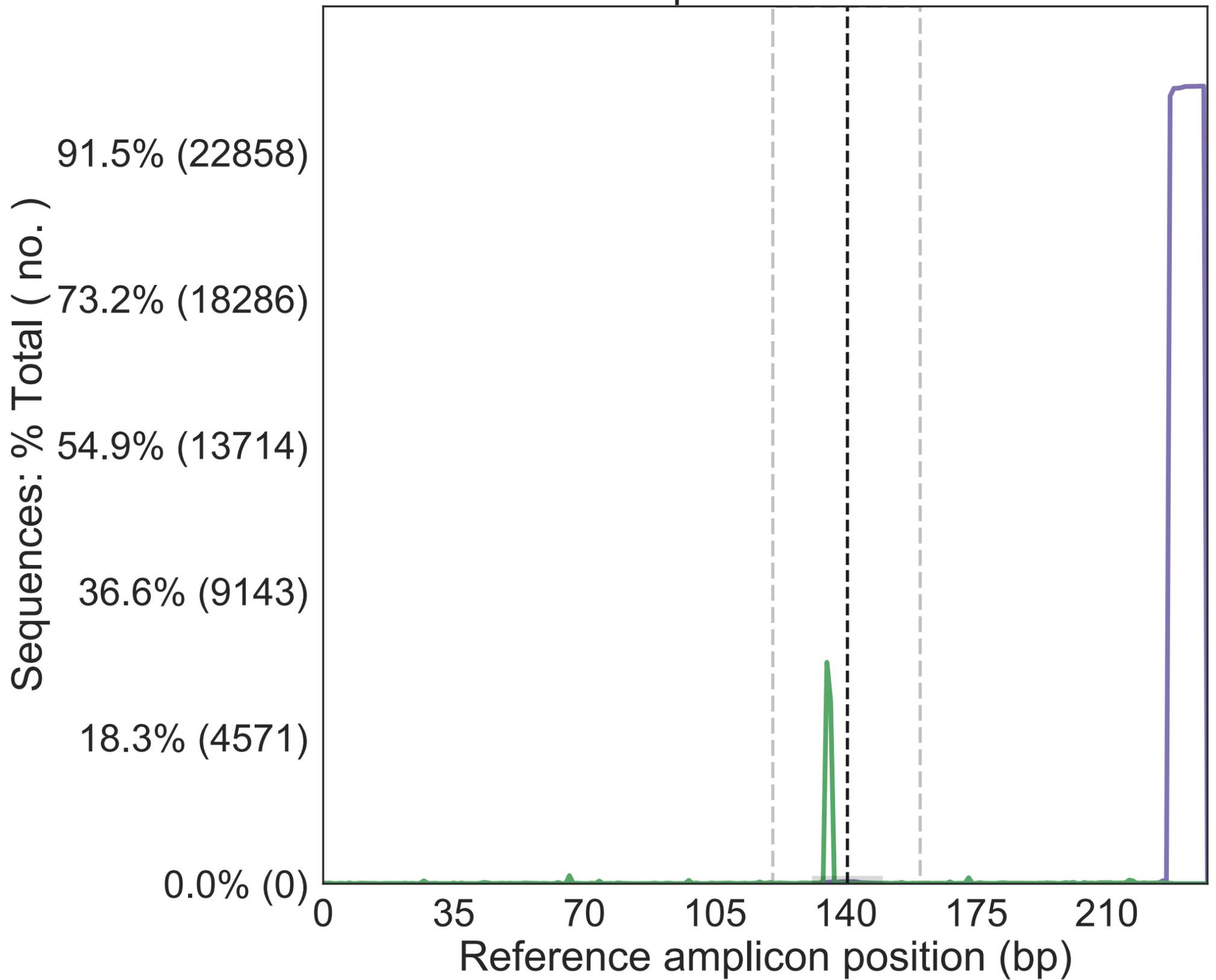


# Mutation position distribution



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