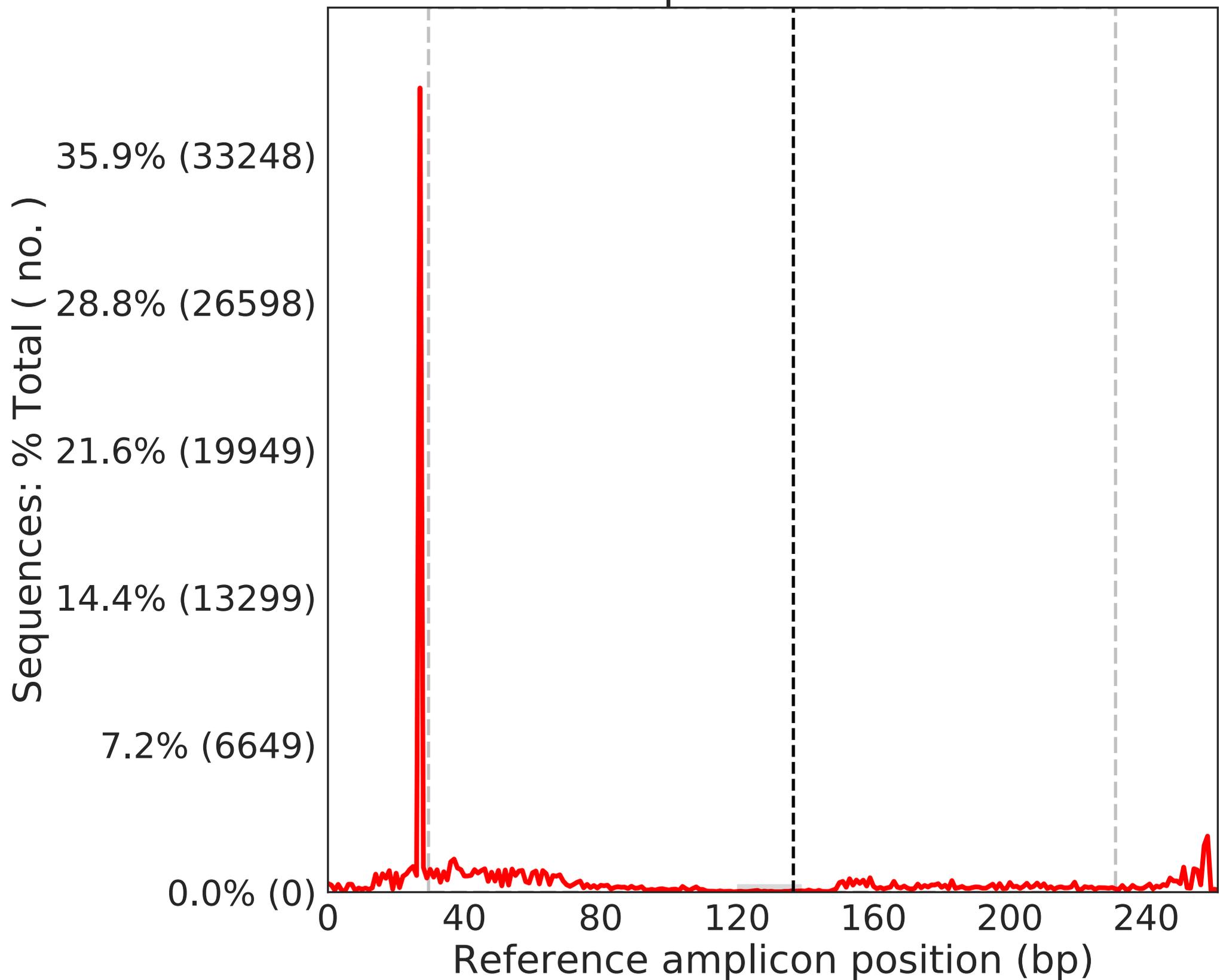


# Mutation position distribution



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