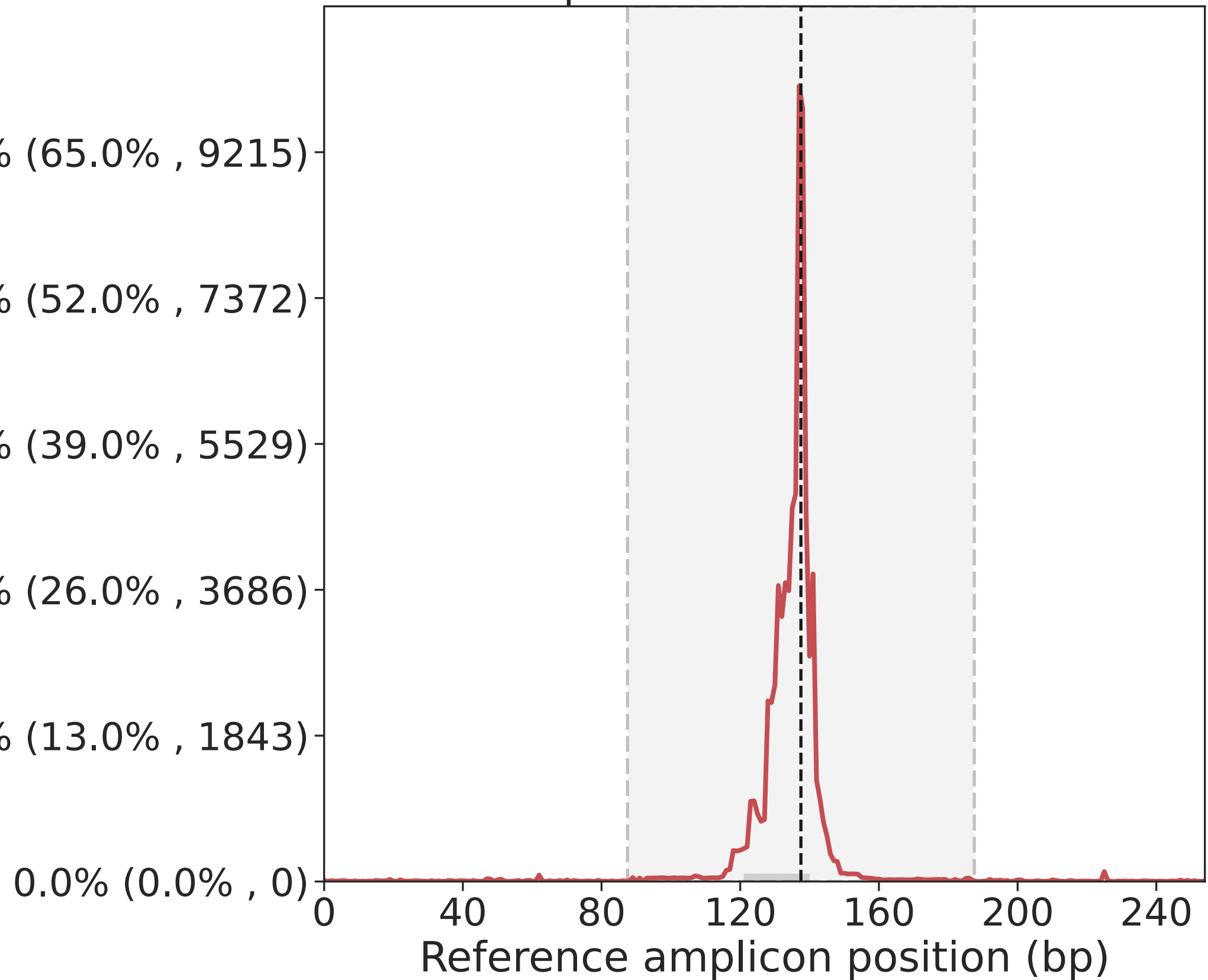


Mutation position distribution: RICTOR

Sequences: % Total (% RICTOR, no.)



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