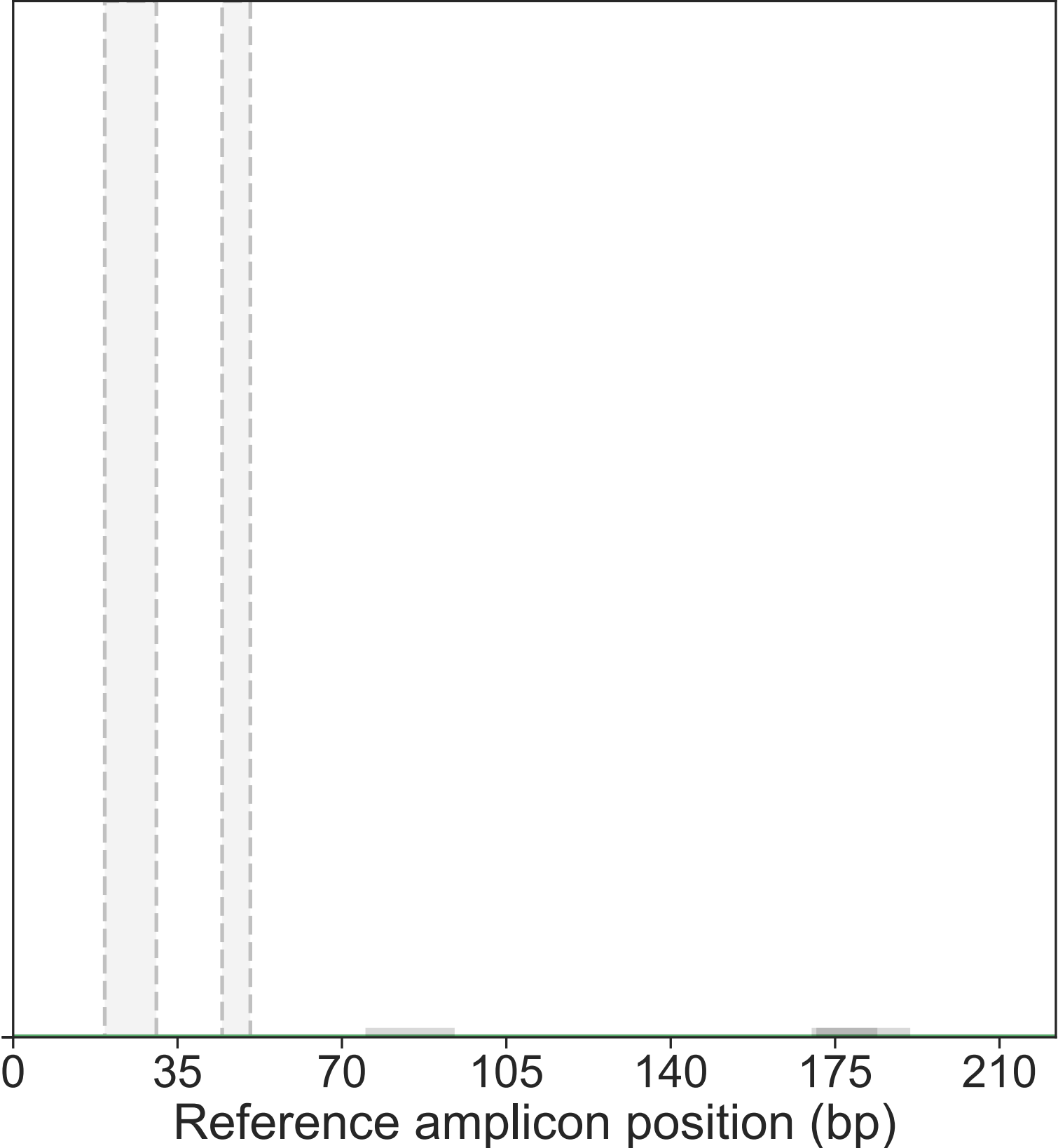


Mutation position distribution: FANC

Sequences: % Total (% FANC, no.)

0.0% (0.0% , 0)



- Insertions
- Deletions
- Substitutions
- sgRNA
- Quantification window