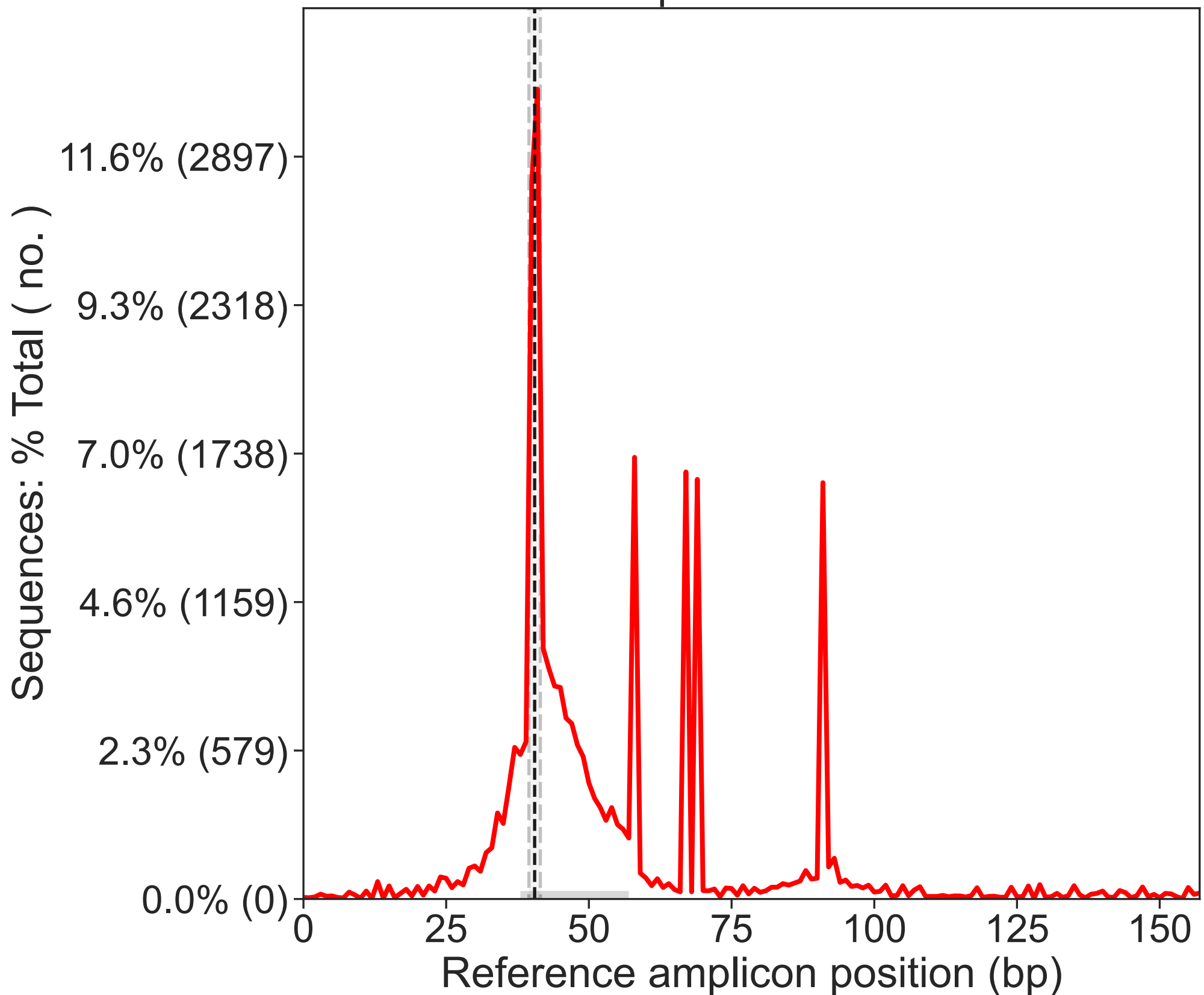


Mutation position distribution



Quantification window



Combined Insertions/Deletions/Substitutions



Predicted cleavage position



sgRNA