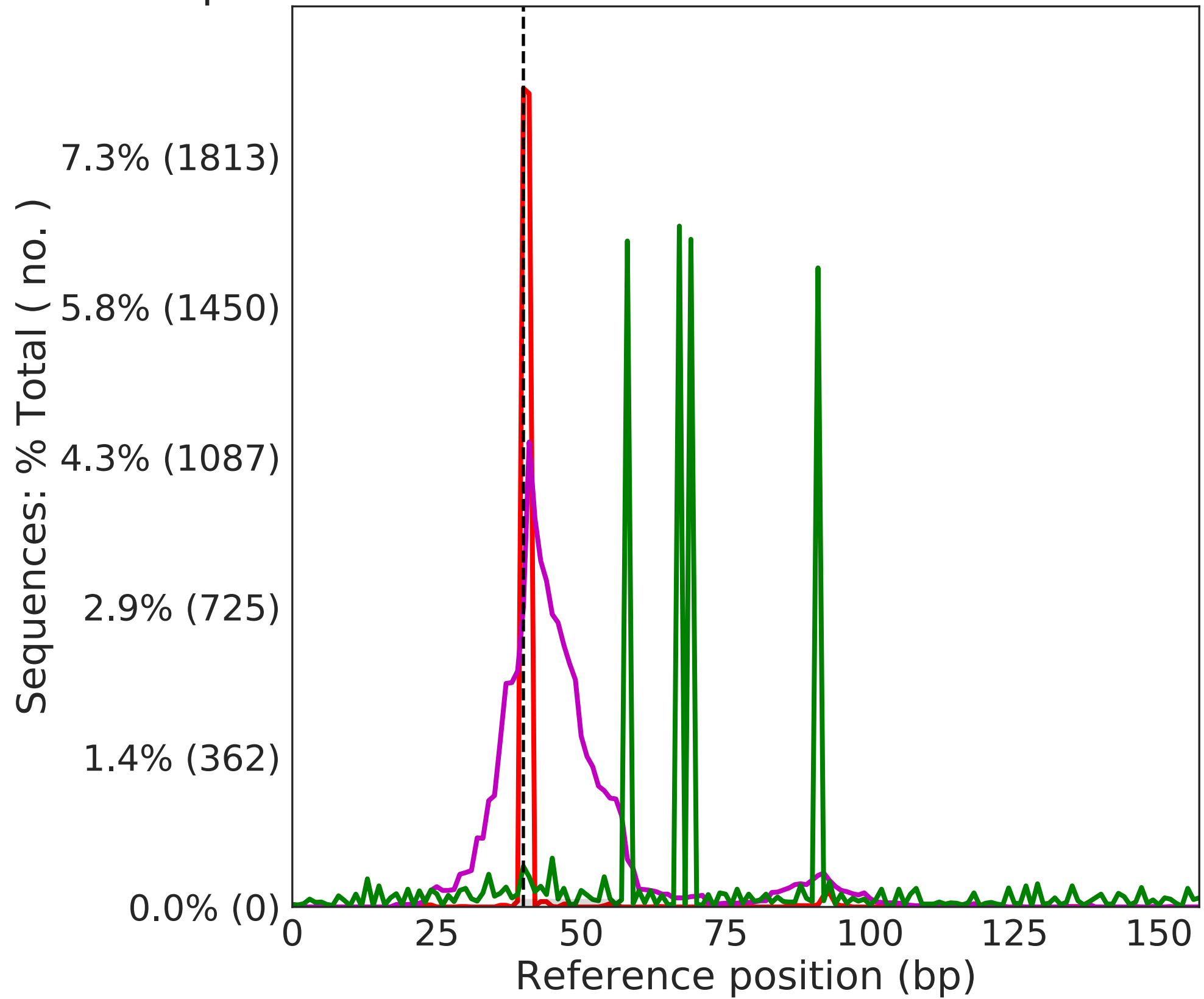


Mutation position distribution in all reads with reference to Reference



- Insertions
- Deletions
- Substitutions
- - - Predicted cleavage position
- sgRNA