

# Mutation position distribution of NHEJ

Sequences: % Total ( % NHEJ, no. )

66.5% (85.5% , 89940)

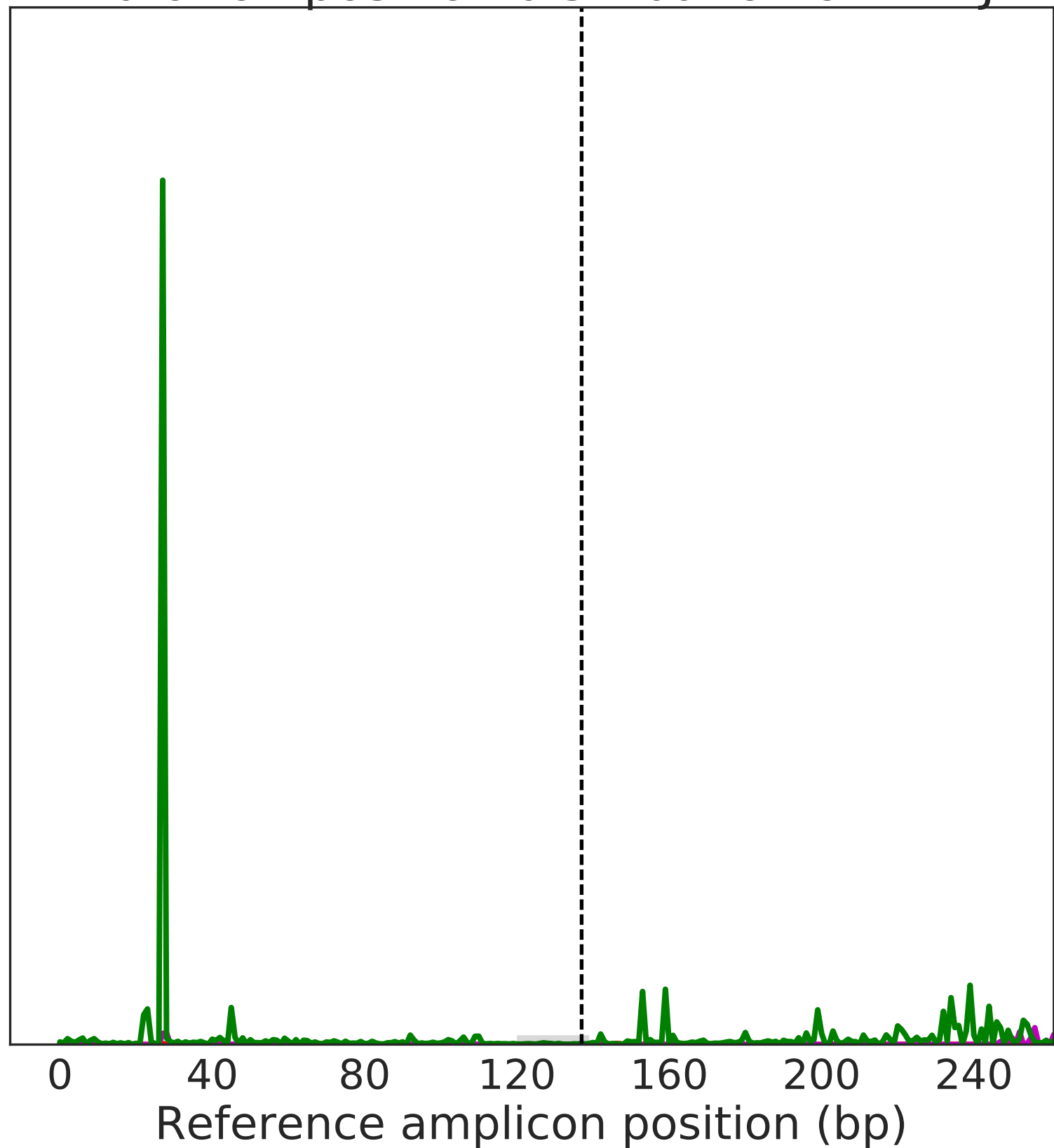
53.2% (68.4% , 71952)

39.9% (51.3% , 53964)

26.6% (34.2% , 35976)

13.3% (17.1% , 17988)

0.0% (0.0% , 0)



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