

# Mutation position distribution of NHEJ

Sequences: % Total ( % NHEJ, no. )

76.1% (87.1% , 168683)

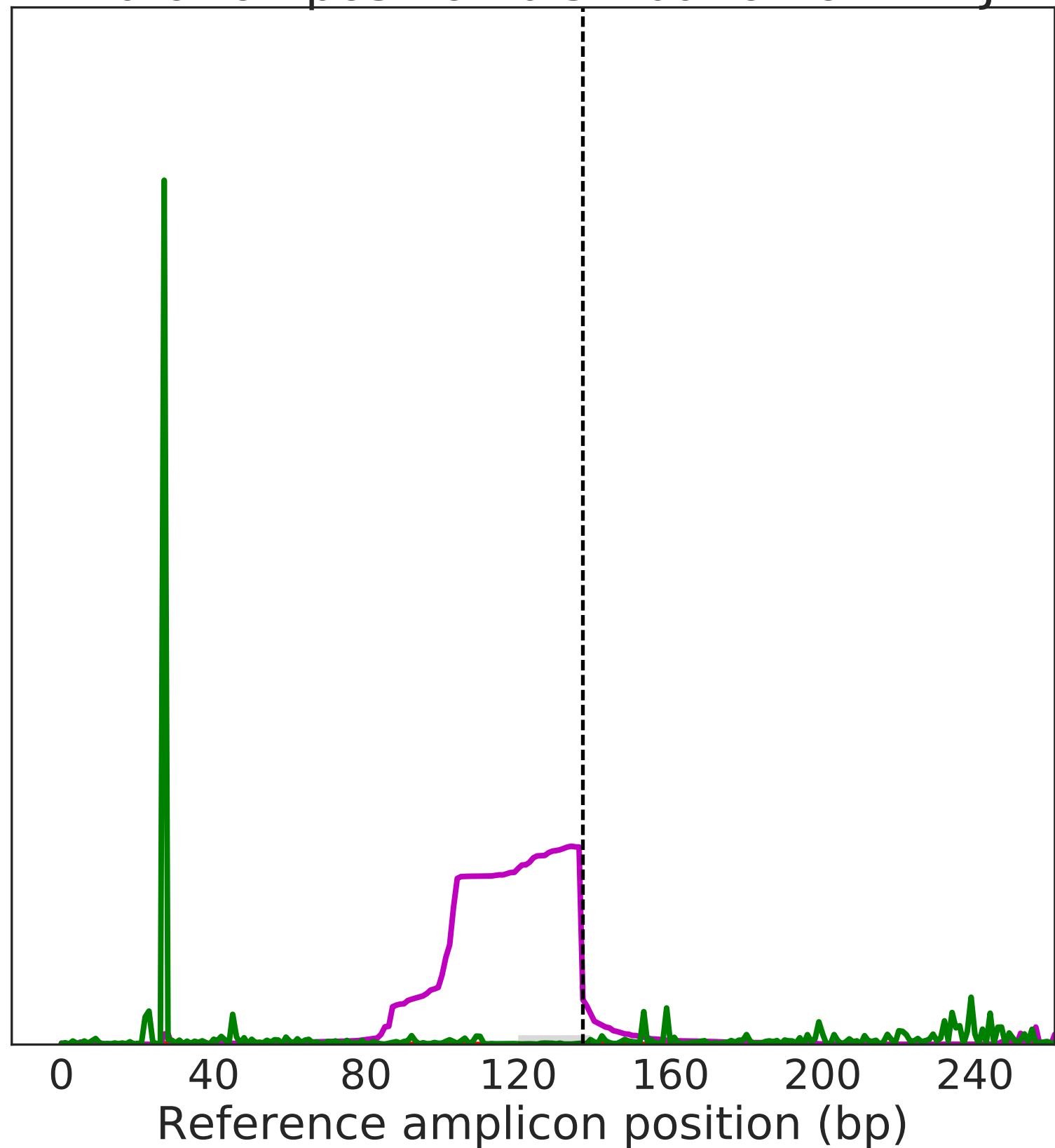
60.9% (69.7% , 134947)

45.6% (52.3% , 101210)

30.4% (34.8% , 67473)

15.2% (17.4% , 33736)

0.0% (0.0% , 0)



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