

# Mutation position distribution of NHEJ

Sequences: % Total ( % NHEJ, no. )

6.2% (45.1% , 20872)

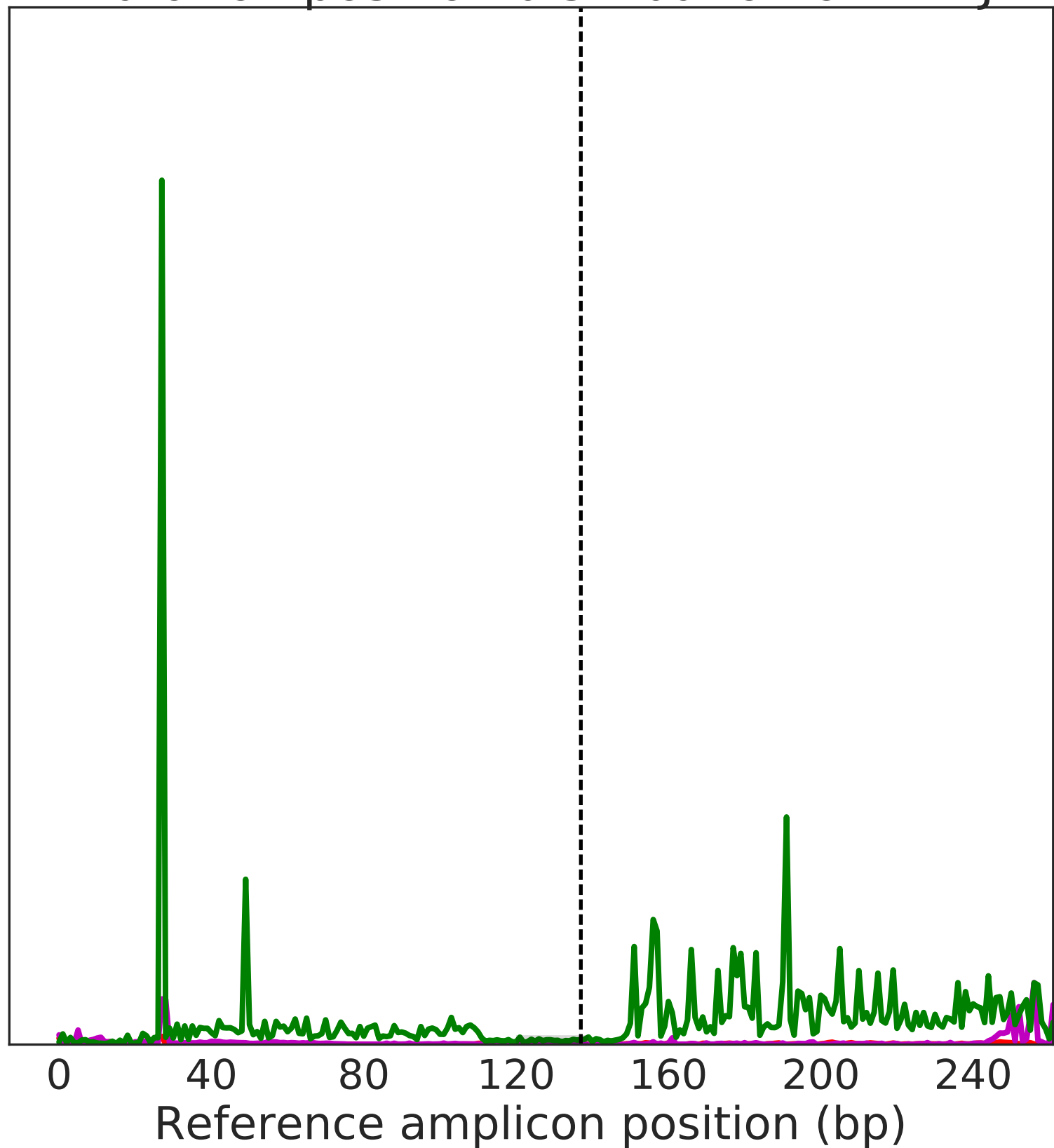
5.0% (36.0% , 16698)

3.7% (27.0% , 12523)

2.5% (18.0% , 8349)

1.2% (9.0% , 4174)

0.0% (0.0% , 0)



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