

# Mutation position distribution of NHEJ

Sequences: % Total ( % NHEJ, no. )

3.3% (19.3% , 15958)

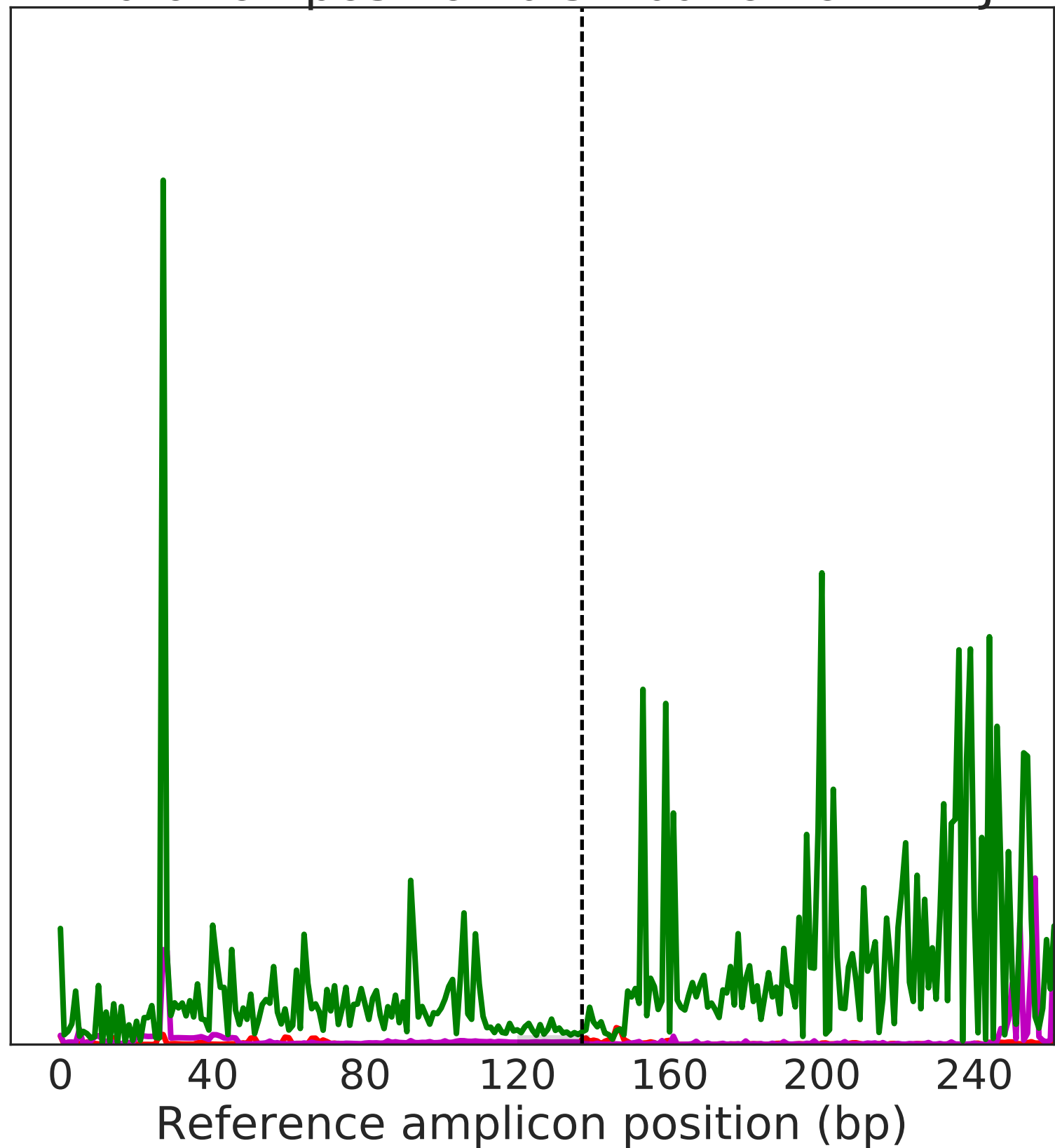
2.7% (15.5% , 12766)

2.0% (11.6% , 9574)

1.3% (7.7% , 6383)

0.7% (3.9% , 3191)

0.0% (0.0% , 0)



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