

# Mutation position distribution of NHEJ

Sequences: % Total ( % NHEJ, no. )

4.0% (28.2% , 12340)

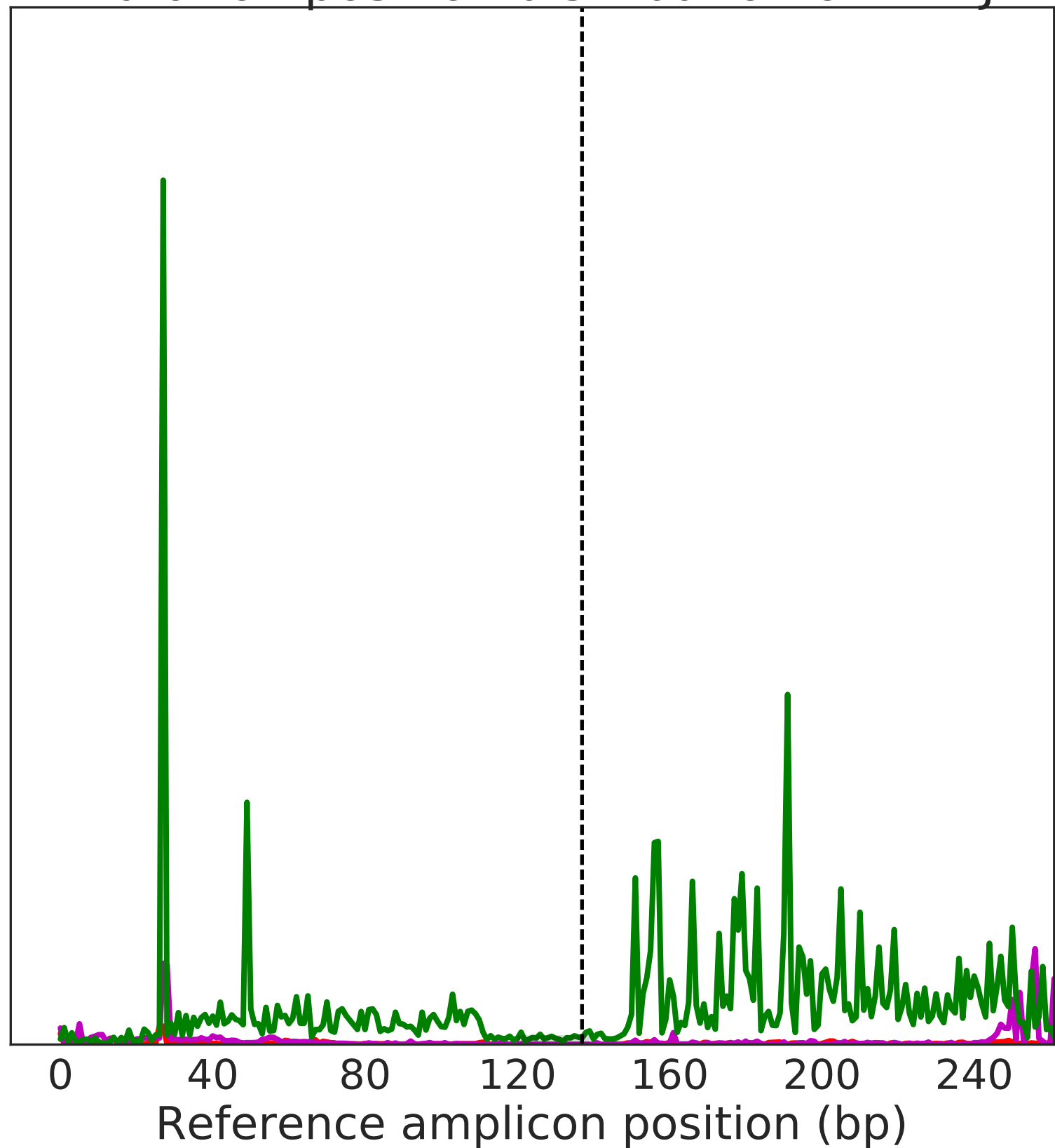
3.2% (22.6% , 9872)

2.4% (16.9% , 7404)

1.6% (11.3% , 4936)

0.8% (5.6% , 2468)

0.0% (0.0% , 0)



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