

# Mutation position distribution of NHEJ

Sequences: % Total ( % NHEJ, no. )

5.8% (43.7% , 19204)

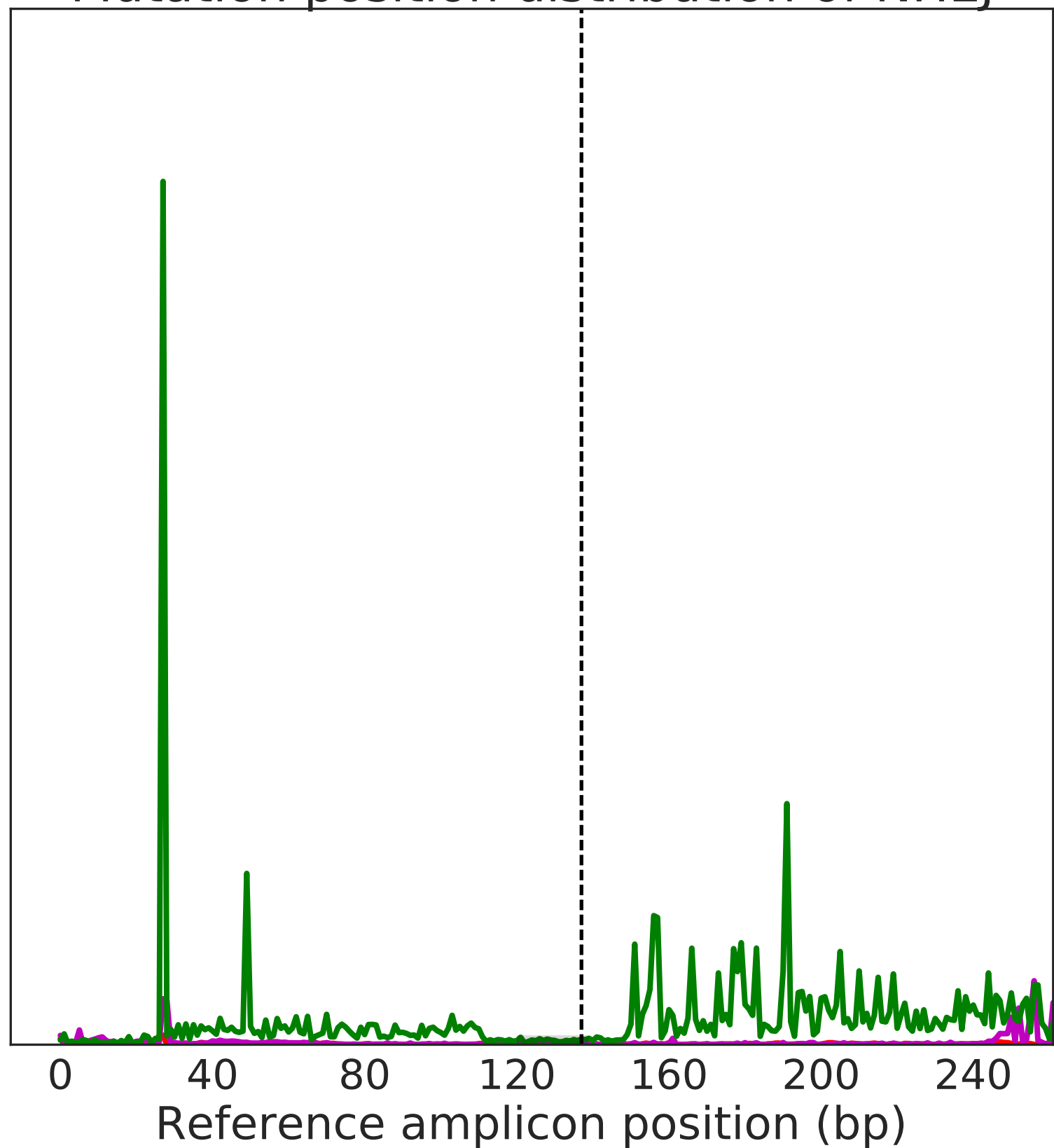
4.6% (35.0% , 15363)

3.5% (26.2% , 11522)

2.3% (17.5% , 7681)

1.2% (8.7% , 3840)

0.0% (0.0% , 0)



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