

# Mutation position distribution of NHEJ

Sequences: % Total ( % NHEJ, no. )

3.8% (21.7% , 32127)

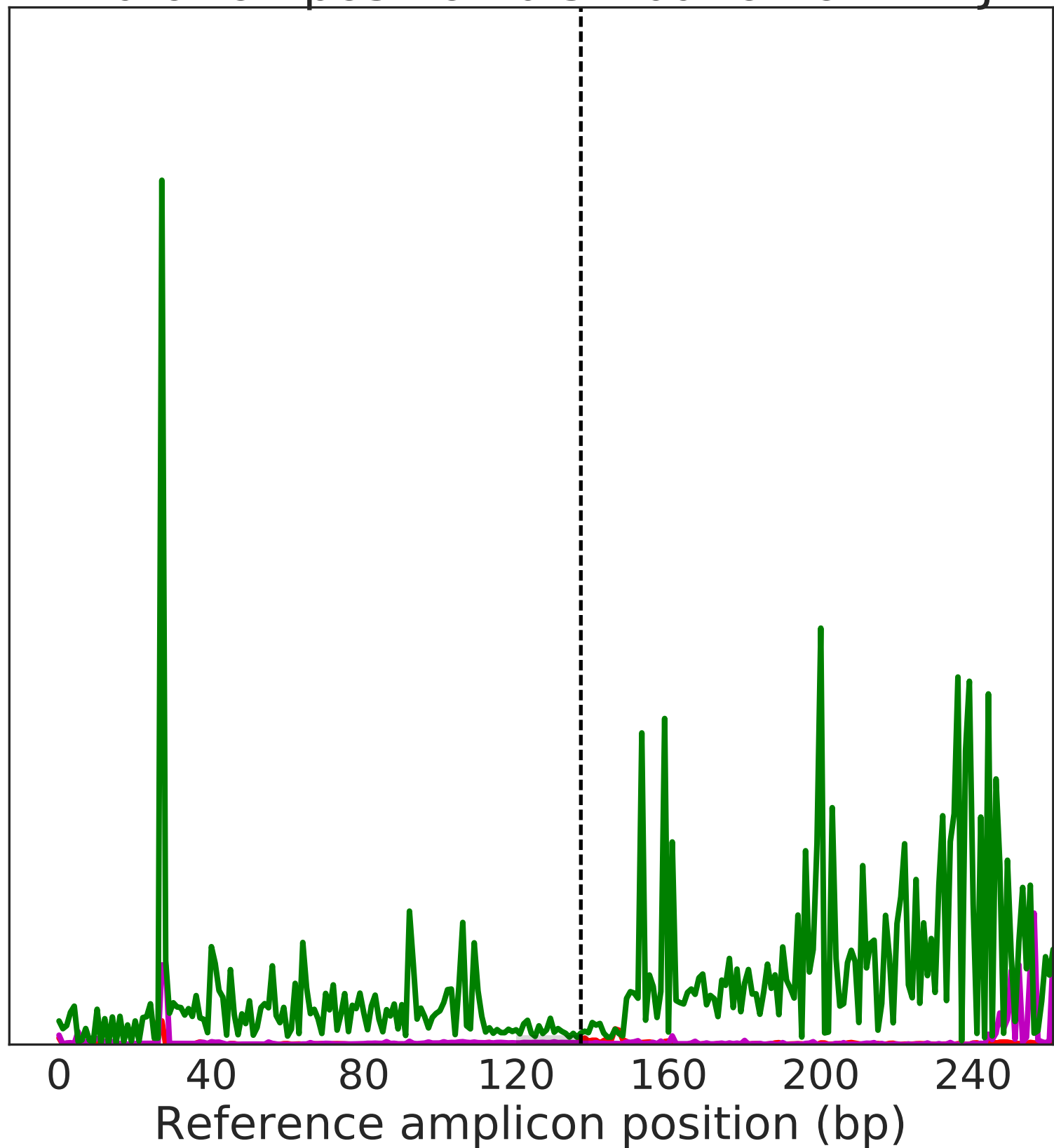
3.1% (17.4% , 25701)

2.3% (13.0% , 19276)

1.5% (8.7% , 12850)

0.8% (4.3% , 6425)

0.0% (0.0% , 0)



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