

Mutation position distribution of NHEJ

Sequences: % Total (% NHEJ, no.)

5.0% (35.2% , 15611)

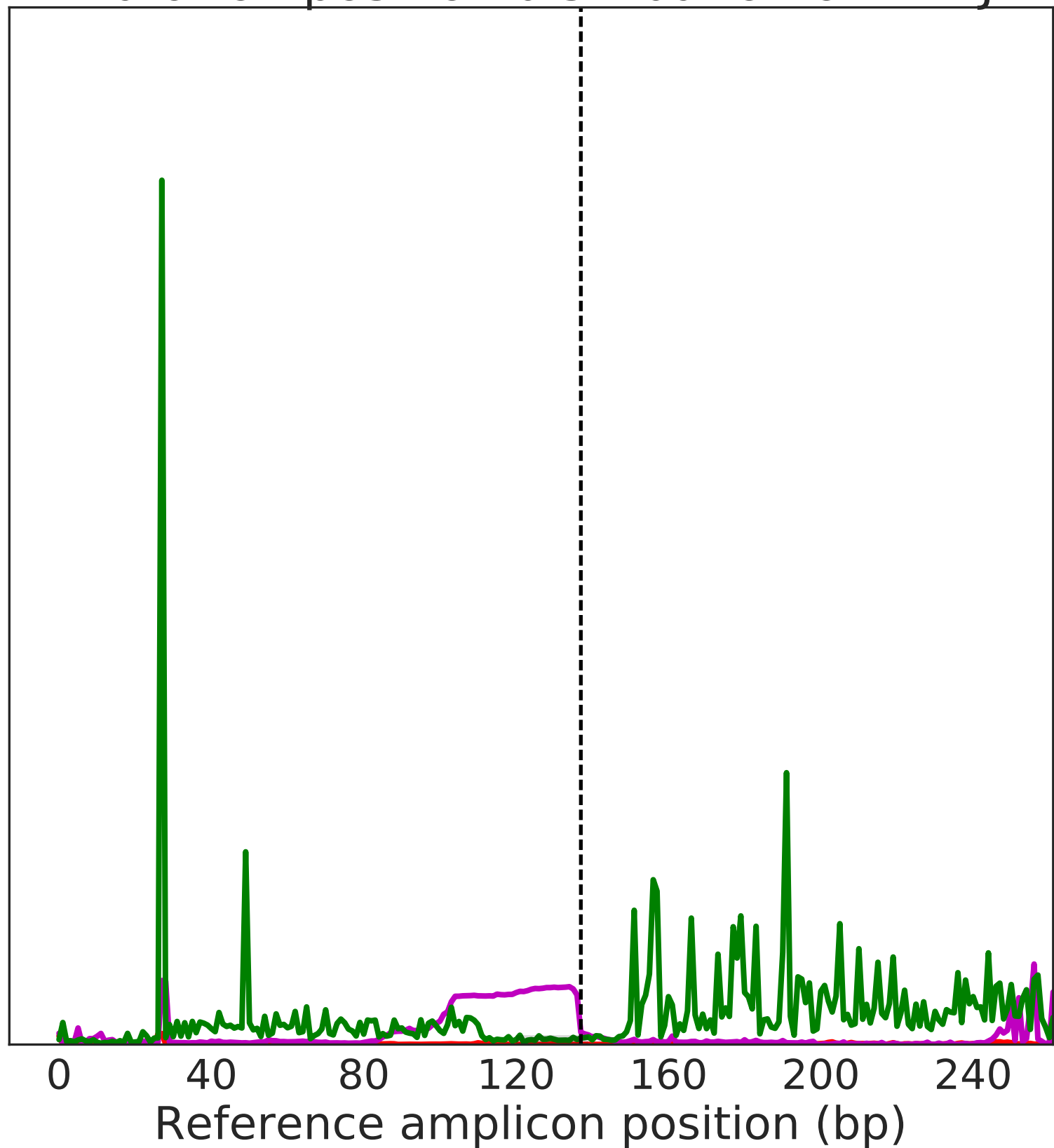
4.0% (28.2% , 12488)

3.0% (21.1% , 9366)

2.0% (14.1% , 6244)

1.0% (7.0% , 3122)

0.0% (0.0% , 0)



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