

Mutation position distribution of NHEJ

Sequences: % Total (% NHEJ, no.)

66.3% (76.1% , 31656)

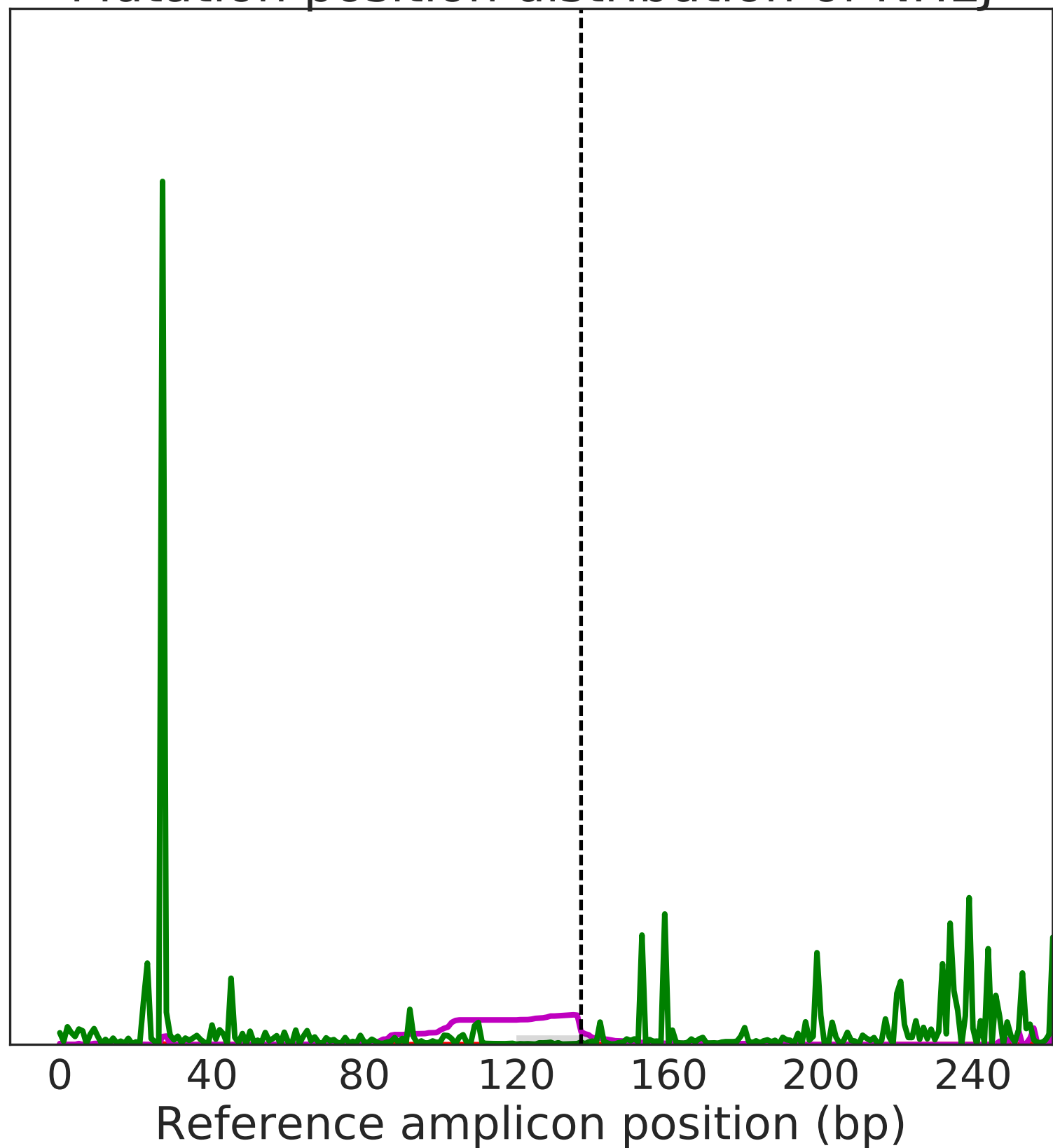
53.1% (60.8% , 25324)

39.8% (45.6% , 18993)

26.5% (30.4% , 12662)

13.3% (15.2% , 6331)

0.0% (0.0% , 0)



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