

# Mutation position distribution of NHEJ

Sequences: % Total ( % NHEJ, no. )

4.3% (28.5% , 14163)

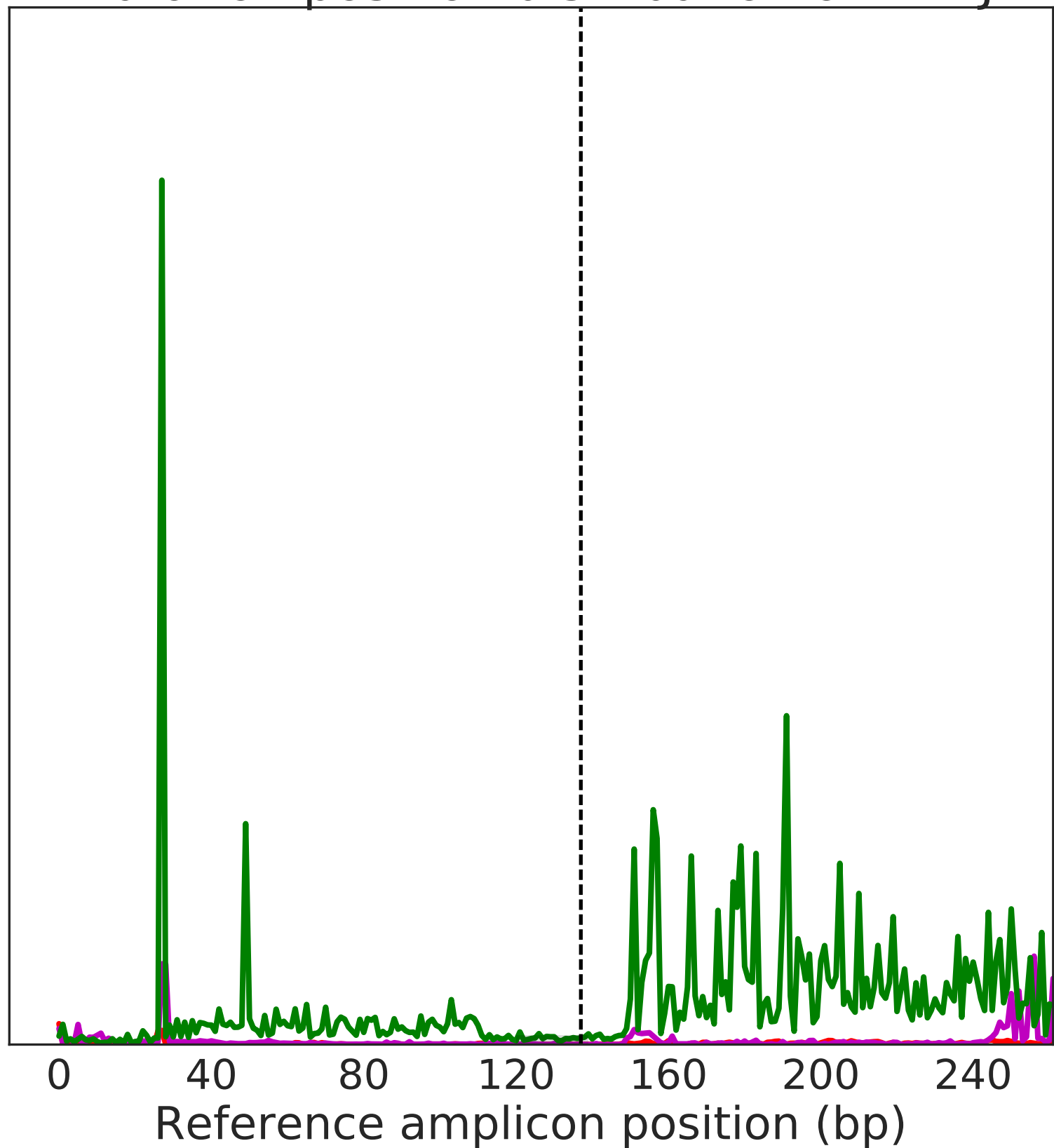
3.4% (22.8% , 11330)

2.6% (17.1% , 8497)

1.7% (11.4% , 5665)

0.9% (5.7% , 2832)

0.0% (0.0% , 0)



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