

Mutation position distribution of NHEJ

Sequences: % Total (% NHEJ, no.)

4.3% (21.3% , 22197)

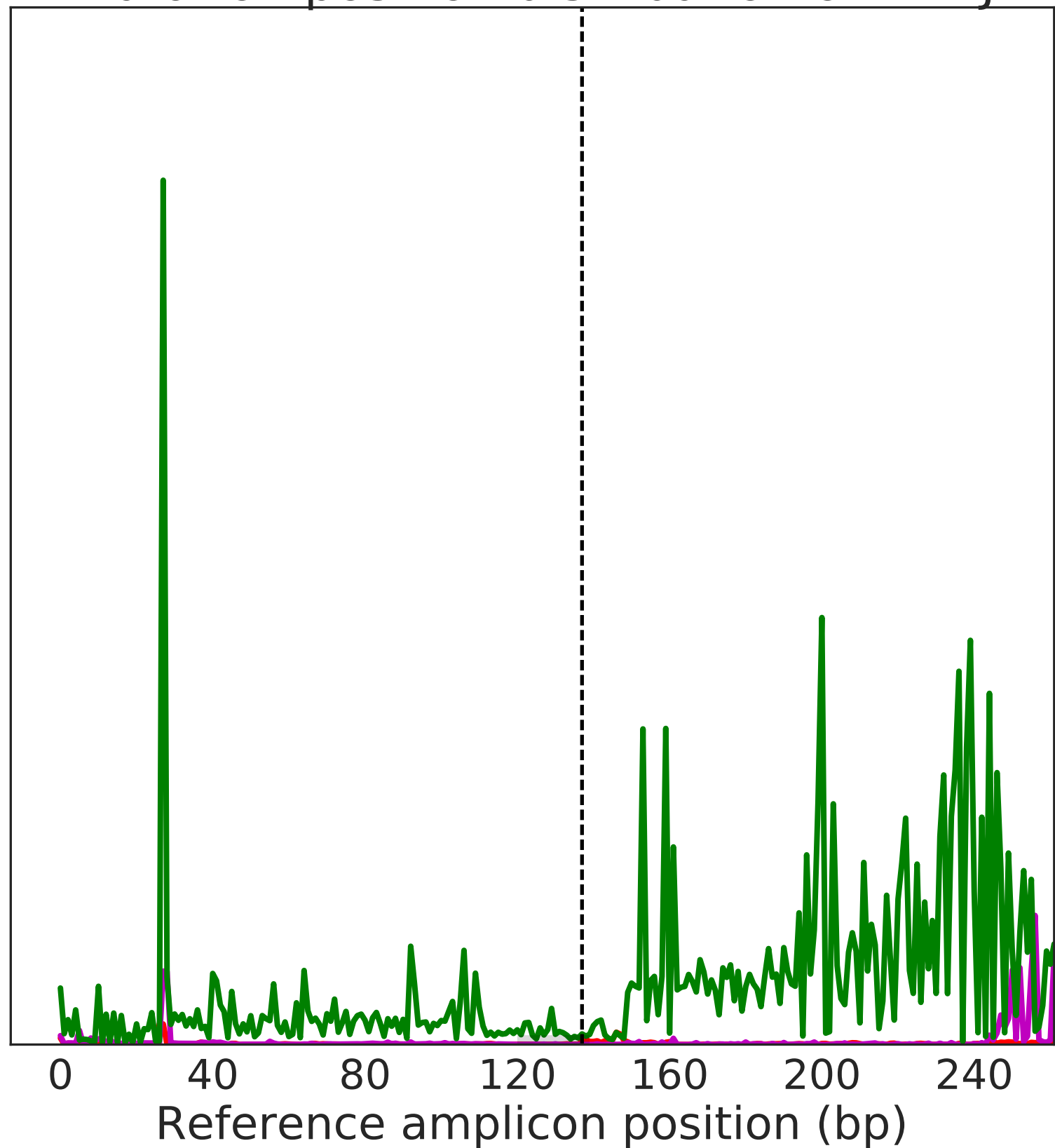
3.4% (17.0% , 17757)

2.6% (12.8% , 13318)

1.7% (8.5% , 8878)

0.9% (4.3% , 4439)

0.0% (0.0% , 0)



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