

Mutation position distribution of NHEJ

Sequences: % Total (% NHEJ, no.)

4.2% (28.2% , 12268)

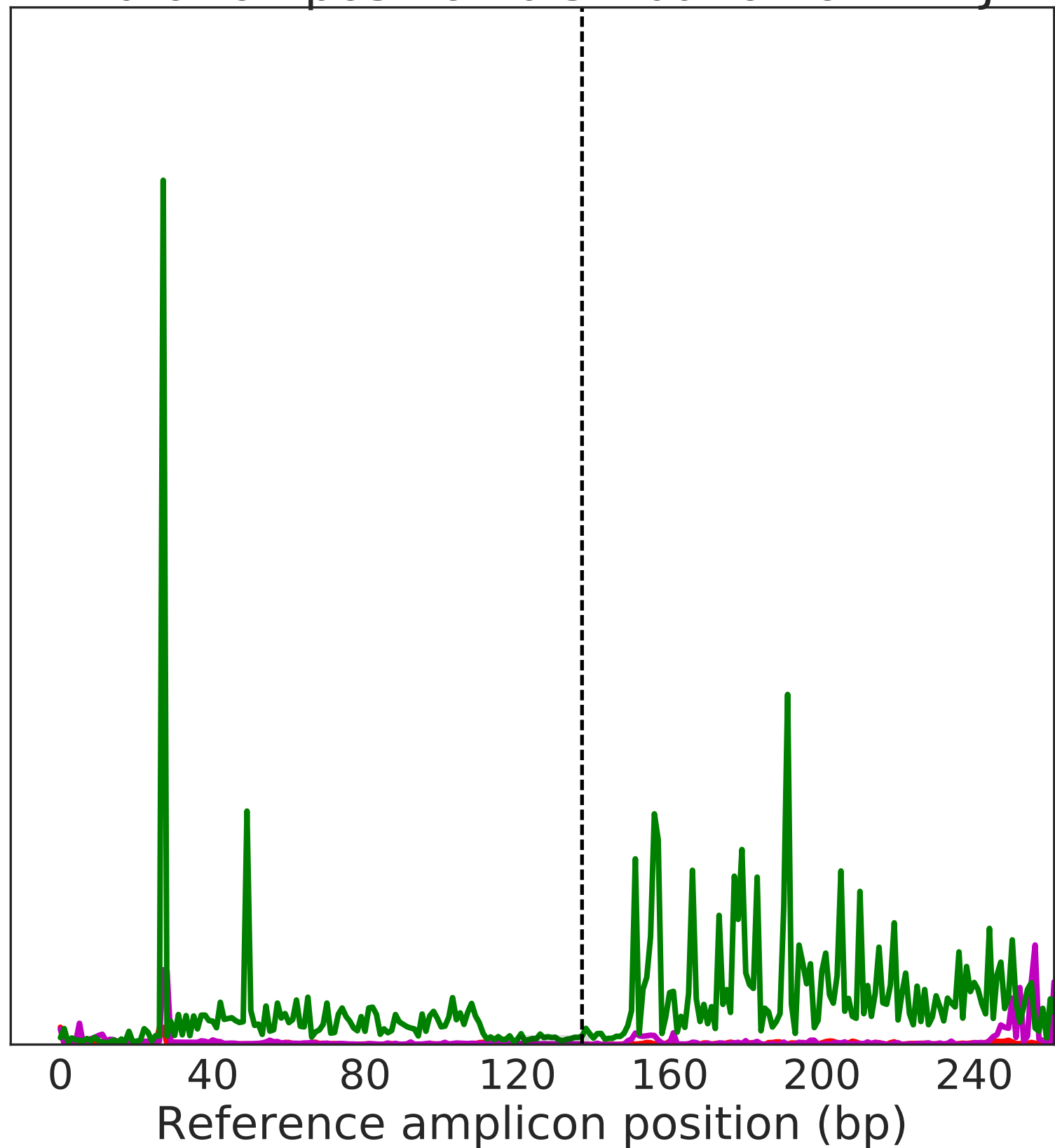
3.3% (22.5% , 9815)

2.5% (16.9% , 7361)

1.7% (11.3% , 4907)

0.8% (5.6% , 2453)

0.0% (0.0% , 0)



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