

Mutation position distribution of NHEJ

Sequences: % Total (% NHEJ, no.)

5.5% (38.5% , 12900)

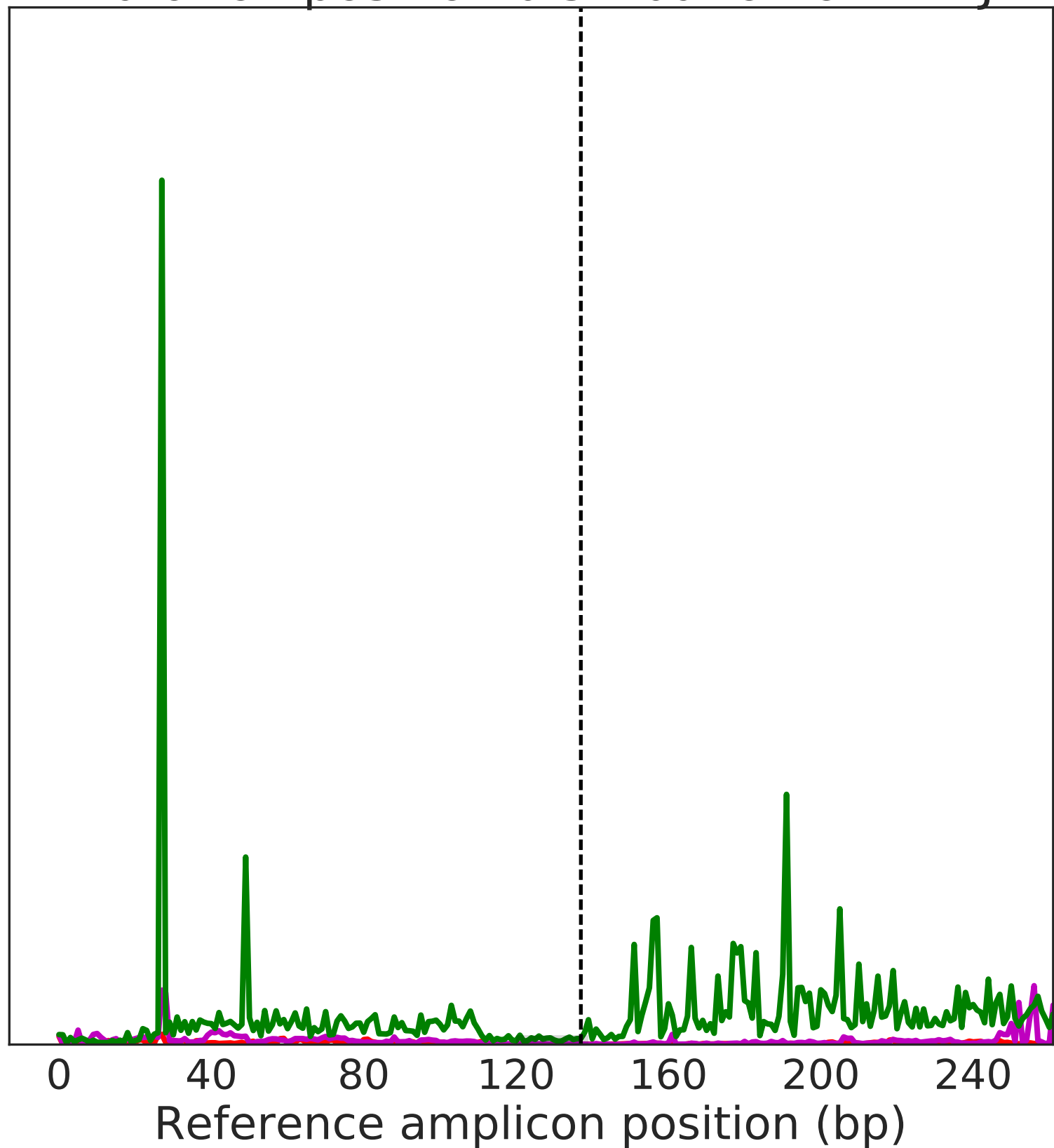
4.4% (30.8% , 10320)

3.3% (23.1% , 7740)

2.2% (15.4% , 5160)

1.1% (7.7% , 2580)

0.0% (0.0% , 0)



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