

Mutation position distribution of NHEJ

Sequences: % Total (% NHEJ, no.)

5.6% (40.7% , 5834)

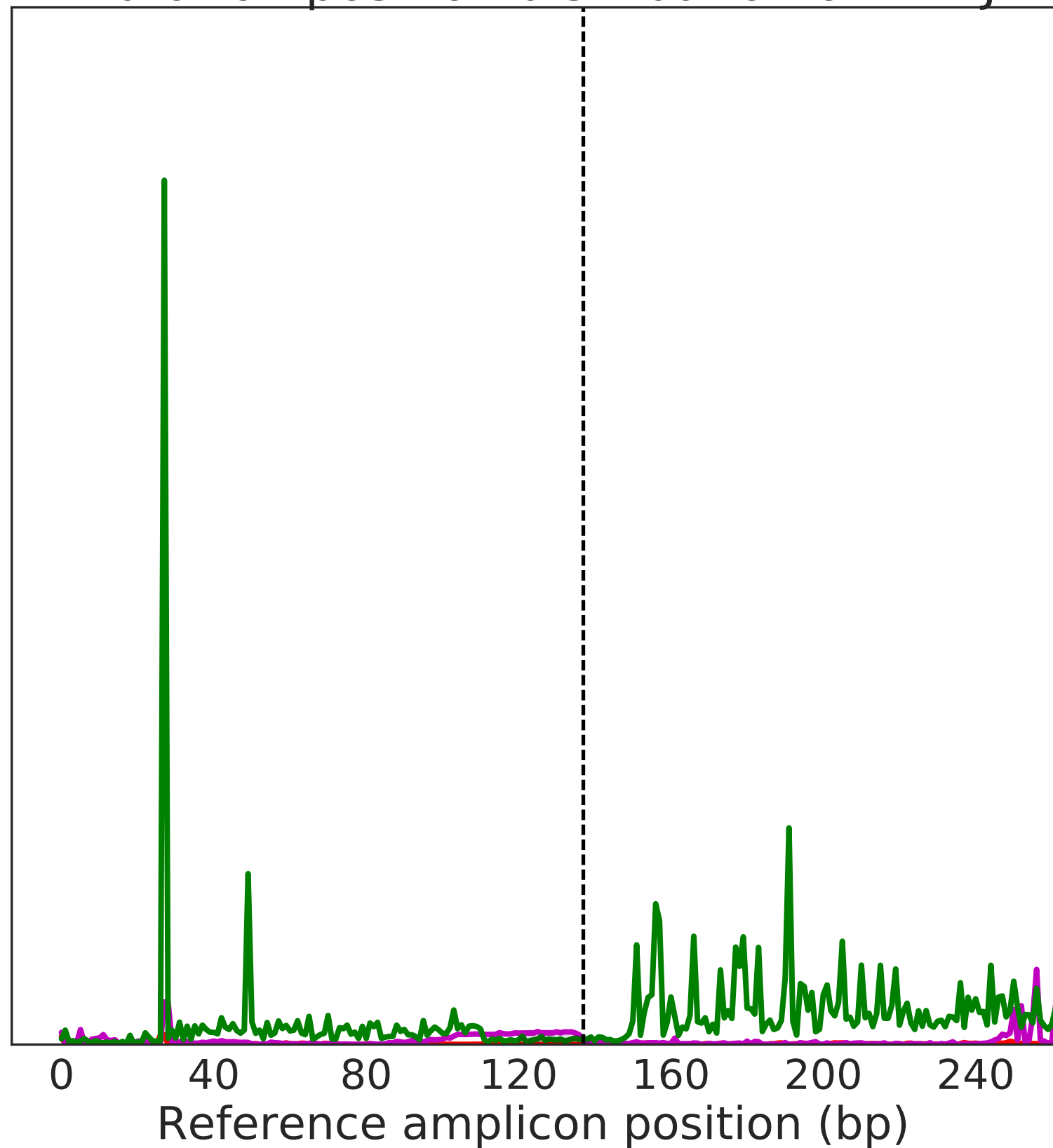
4.5% (32.6% , 4667)

3.3% (24.4% , 3500)

2.2% (16.3% , 2333)

1.1% (8.1% , 1166)

0.0% (0.0% , 0)



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