

Mutation position distribution of NHEJ

Sequences: % Total (% NHEJ, no.)

8.3% (60.9% , 31082)

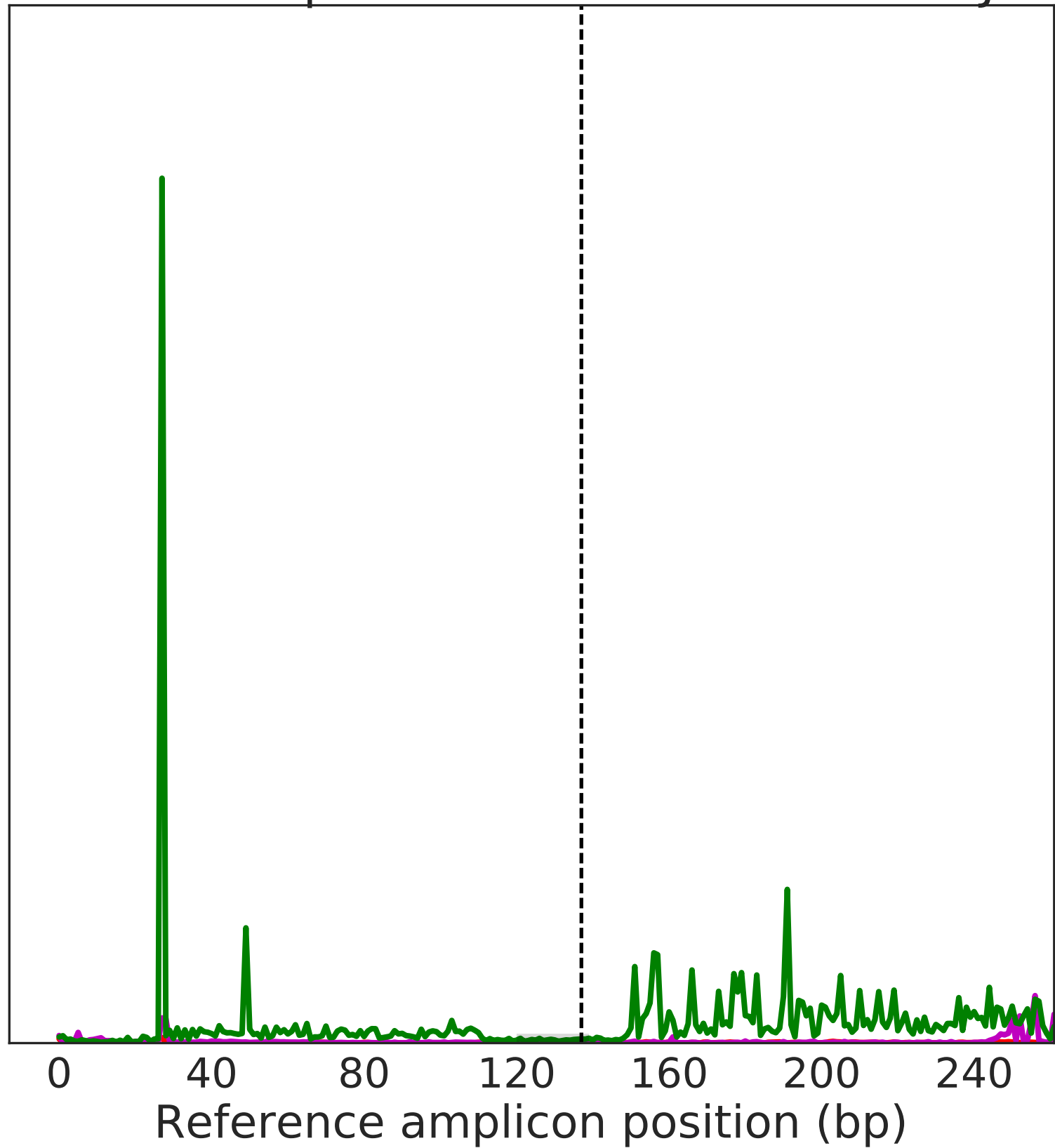
6.6% (48.7% , 24865)

5.0% (36.5% , 18649)

3.3% (24.4% , 12432)

1.7% (12.2% , 6216)

0.0% (0.0% , 0)



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