

Mutation position distribution of NHEJ

Sequences: % Total (% NHEJ, no.)

4.1% (29.8% , 8663)

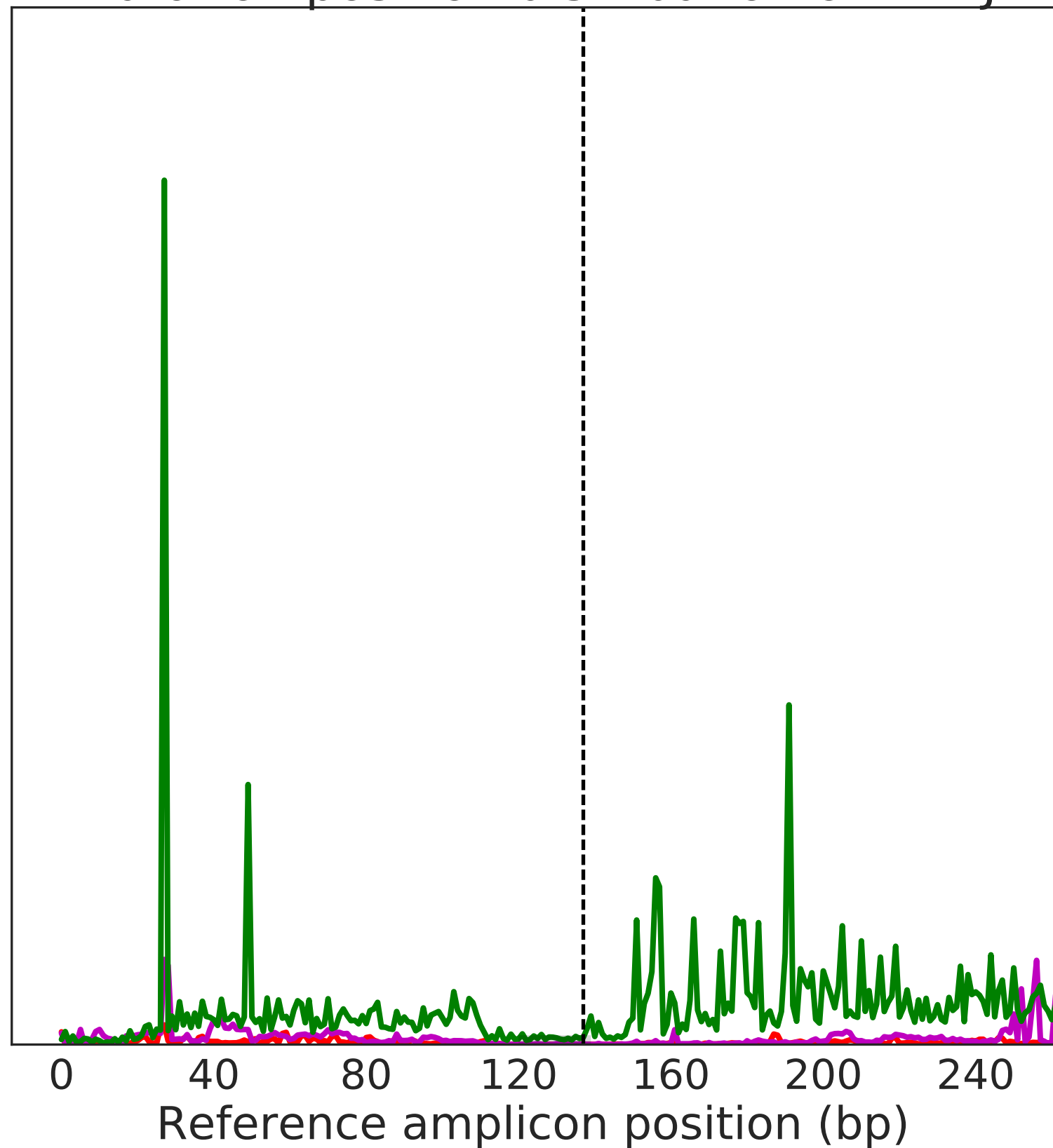
3.3% (23.9% , 6930)

2.4% (17.9% , 5197)

1.6% (11.9% , 3465)

0.8% (6.0% , 1732)

0.0% (0.0% , 0)



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