

Mutation position distribution of NHEJ

Sequences: % Total (% NHEJ, no.)

9.7% (67.7% , 28947)

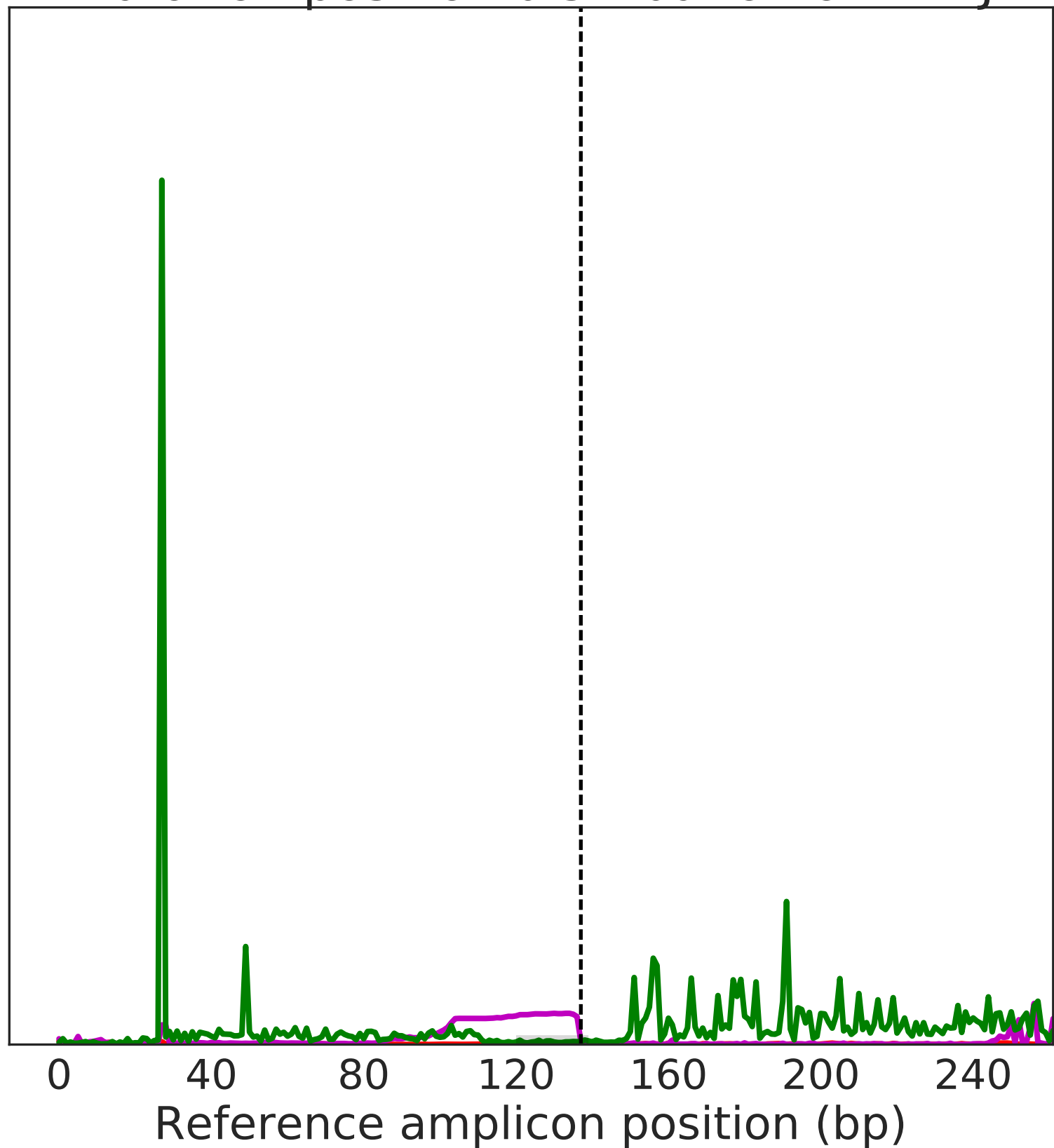
7.7% (54.2% , 23158)

5.8% (40.6% , 17368)

3.9% (27.1% , 11579)

1.9% (13.5% , 5789)

0.0% (0.0% , 0)



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