

Mutation position distribution of NHEJ

Sequences: % Total (% NHEJ, no.)

3.9% (23.1% , 26074)

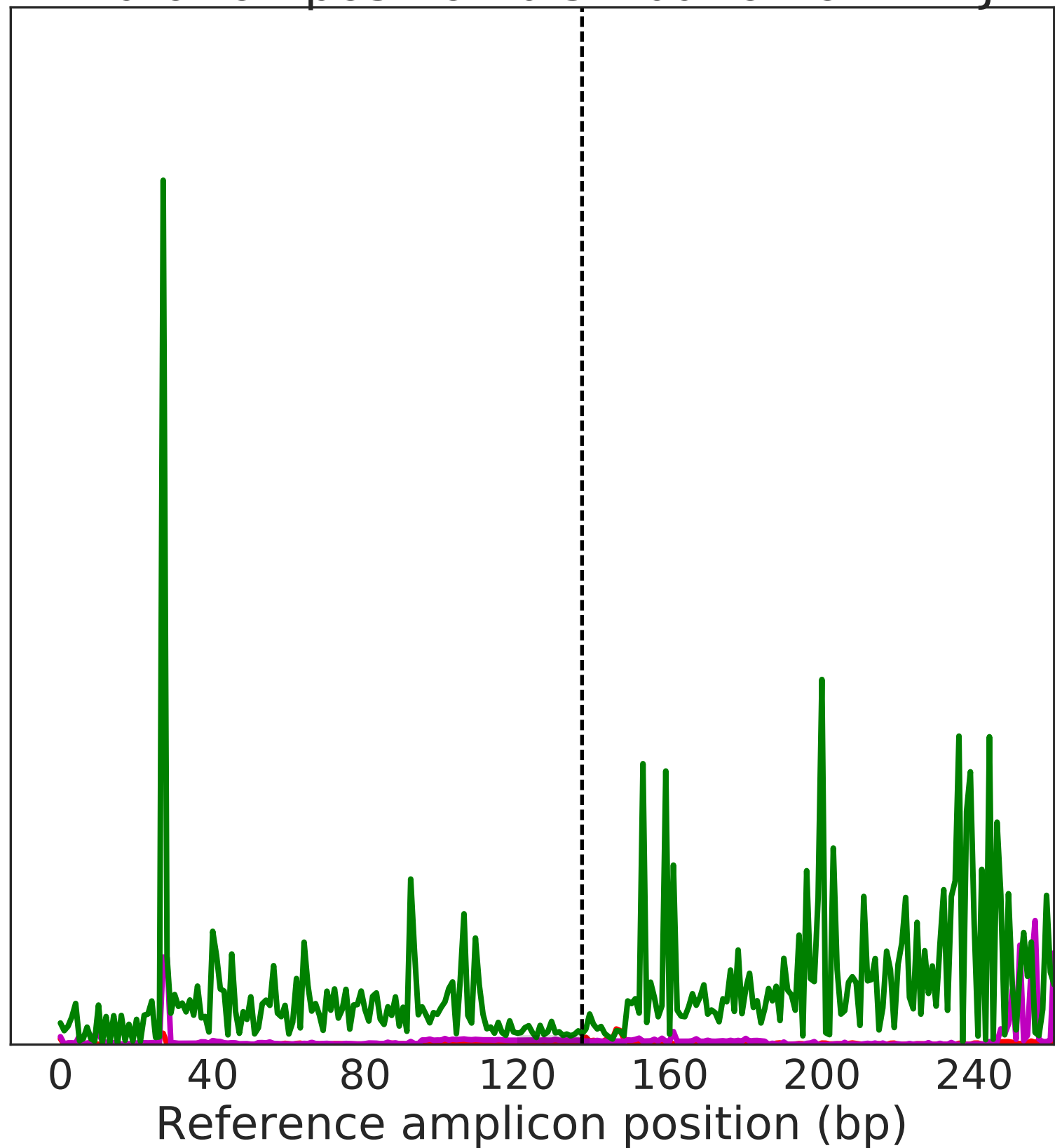
3.1% (18.5% , 20859)

2.3% (13.9% , 15644)

1.6% (9.3% , 10429)

0.8% (4.6% , 5214)

0.0% (0.0% , 0)



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