

Mutation position distribution of NHEJ

Sequences: % Total (% NHEJ, no.)

3.8% (28.6% , 4755)

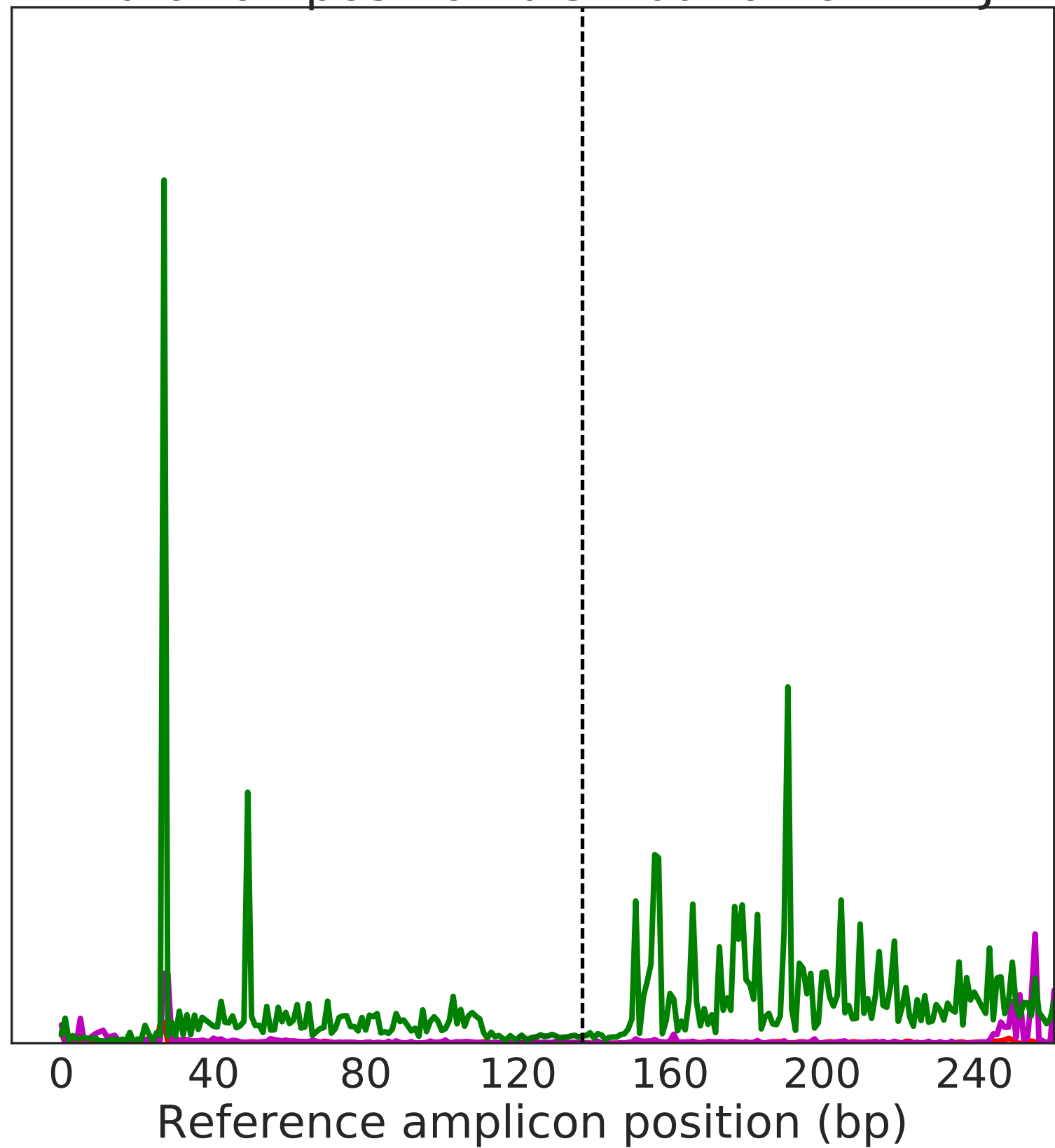
3.0% (22.9% , 3804)

2.3% (17.1% , 2853)

1.5% (11.4% , 1902)

0.8% (5.7% , 951)

0.0% (0.0% , 0)



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