

Mutation position distribution of NHEJ

Sequences: % Total (% NHEJ, no.)

3.9% (27.9% , 17062)

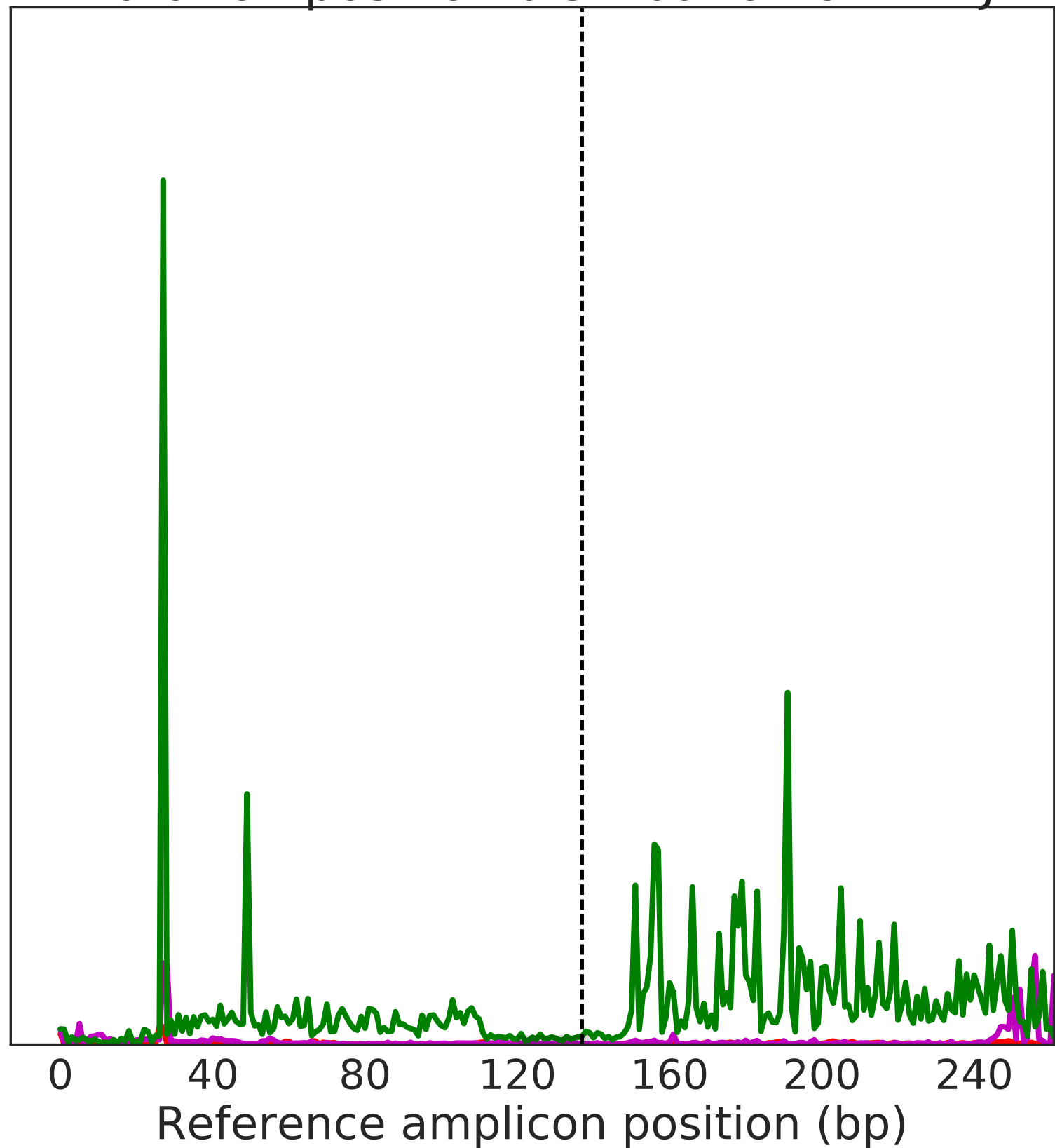
3.1% (22.3% , 13649)

2.3% (16.7% , 10237)

1.6% (11.1% , 6824)

0.8% (5.6% , 3412)

0.0% (0.0% , 0)



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