

# Mutation position distribution of NHEJ

Sequences: % Total ( % NHEJ, no. )

4.2% (27.0% , 13058)

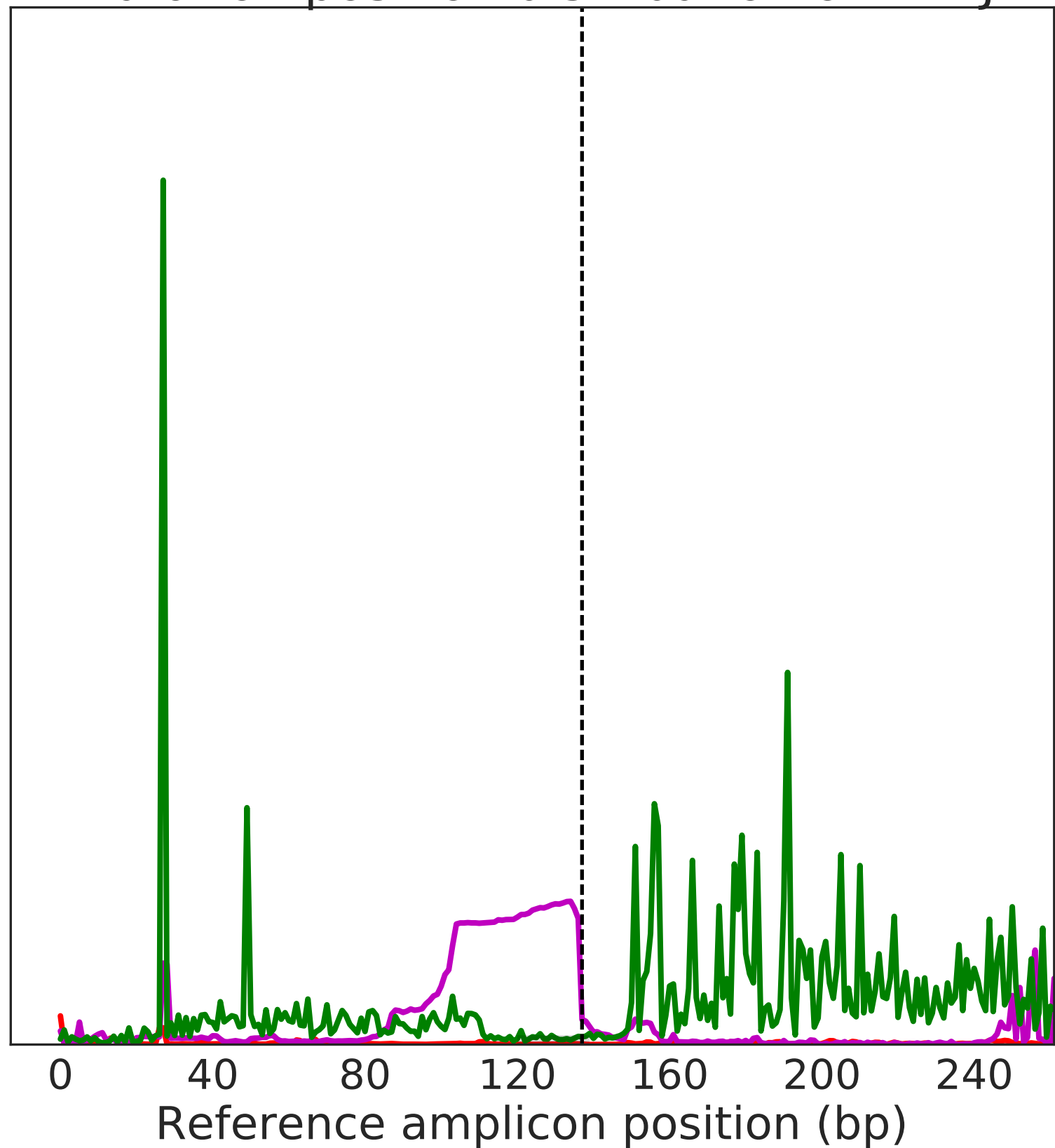
3.4% (21.6% , 10446)

2.5% (16.2% , 7834)

1.7% (10.8% , 5223)

0.8% (5.4% , 2611)

0.0% (0.0% , 0)



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