

# Mutation position distribution of NHEJ

Sequences: % Total ( % NHEJ, no. )

3.6% (23.0% , 26776)

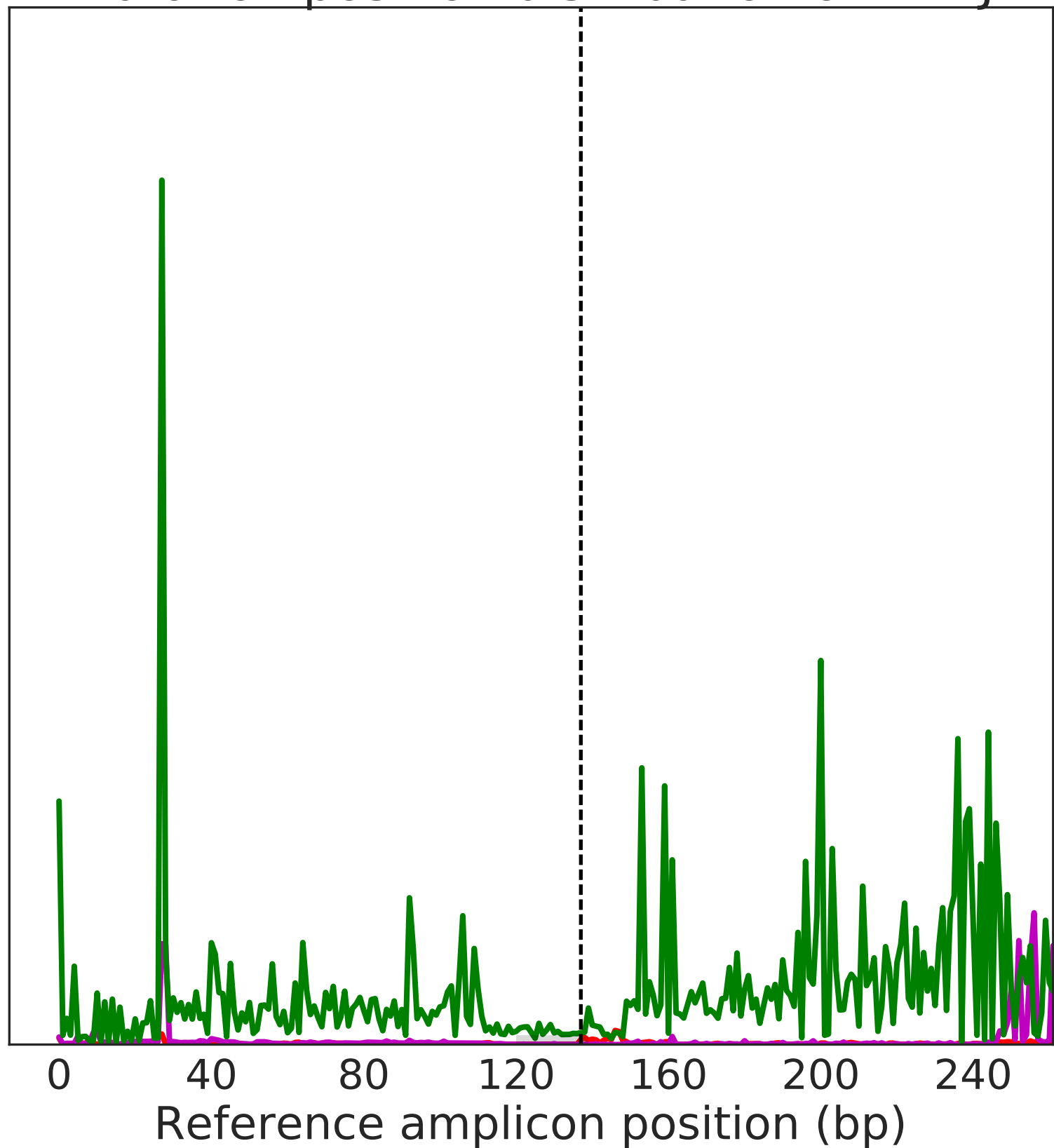
2.9% (18.4% , 21420)

2.2% (13.8% , 16065)

1.5% (9.2% , 10710)

0.7% (4.6% , 5355)

0.0% (0.0% , 0)



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