

# Mutation position distribution of NHEJ

Sequences: % Total ( % NHEJ, no. )

4.5% (29.3% , 11926)

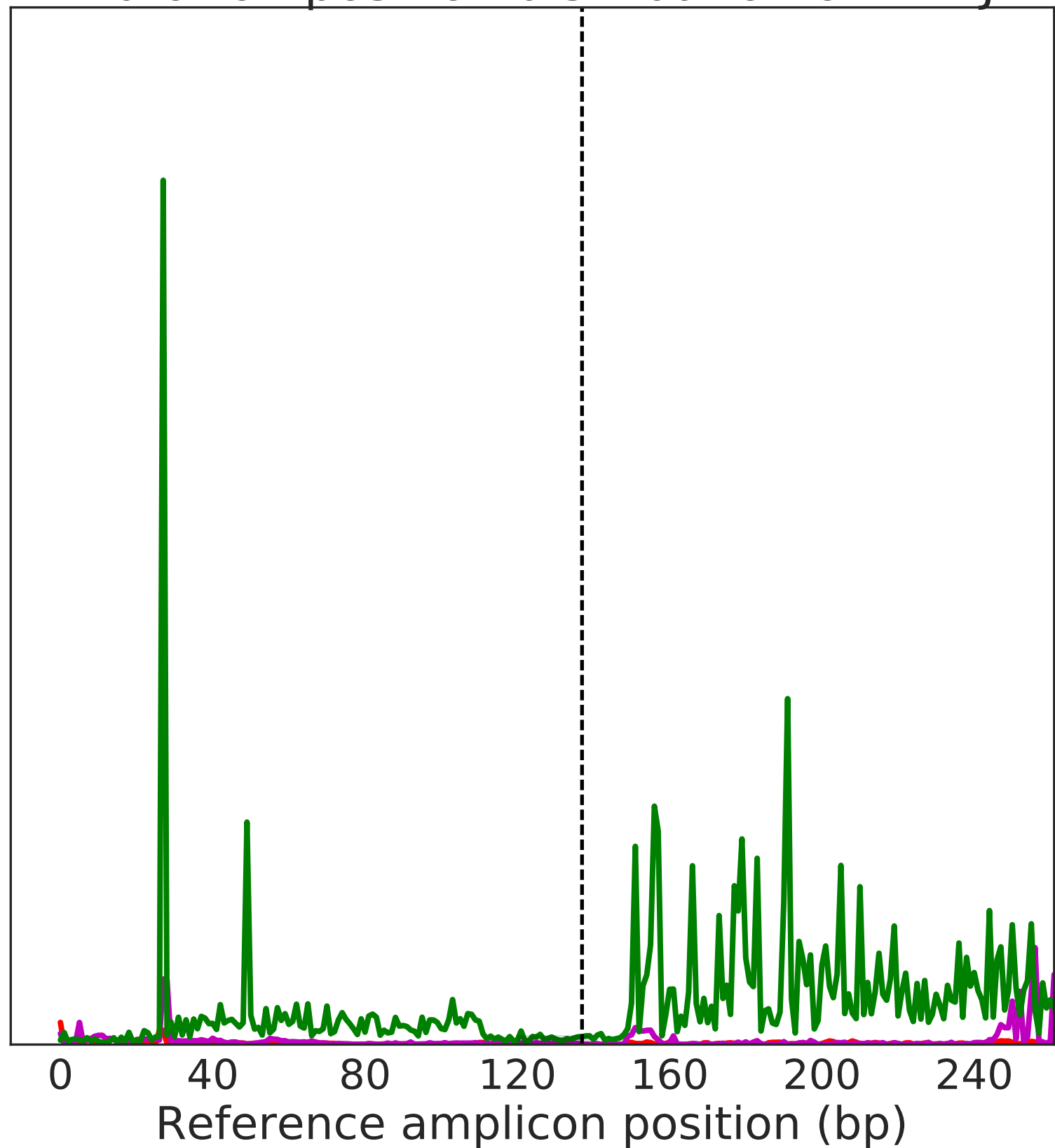
3.6% (23.4% , 9540)

2.7% (17.6% , 7155)

1.8% (11.7% , 4770)

0.9% (5.9% , 2385)

0.0% (0.0% , 0)



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