

# Mutation position distribution of NHEJ

Sequences: % Total ( % NHEJ, no. )

4.0% (29.8% , 5311)

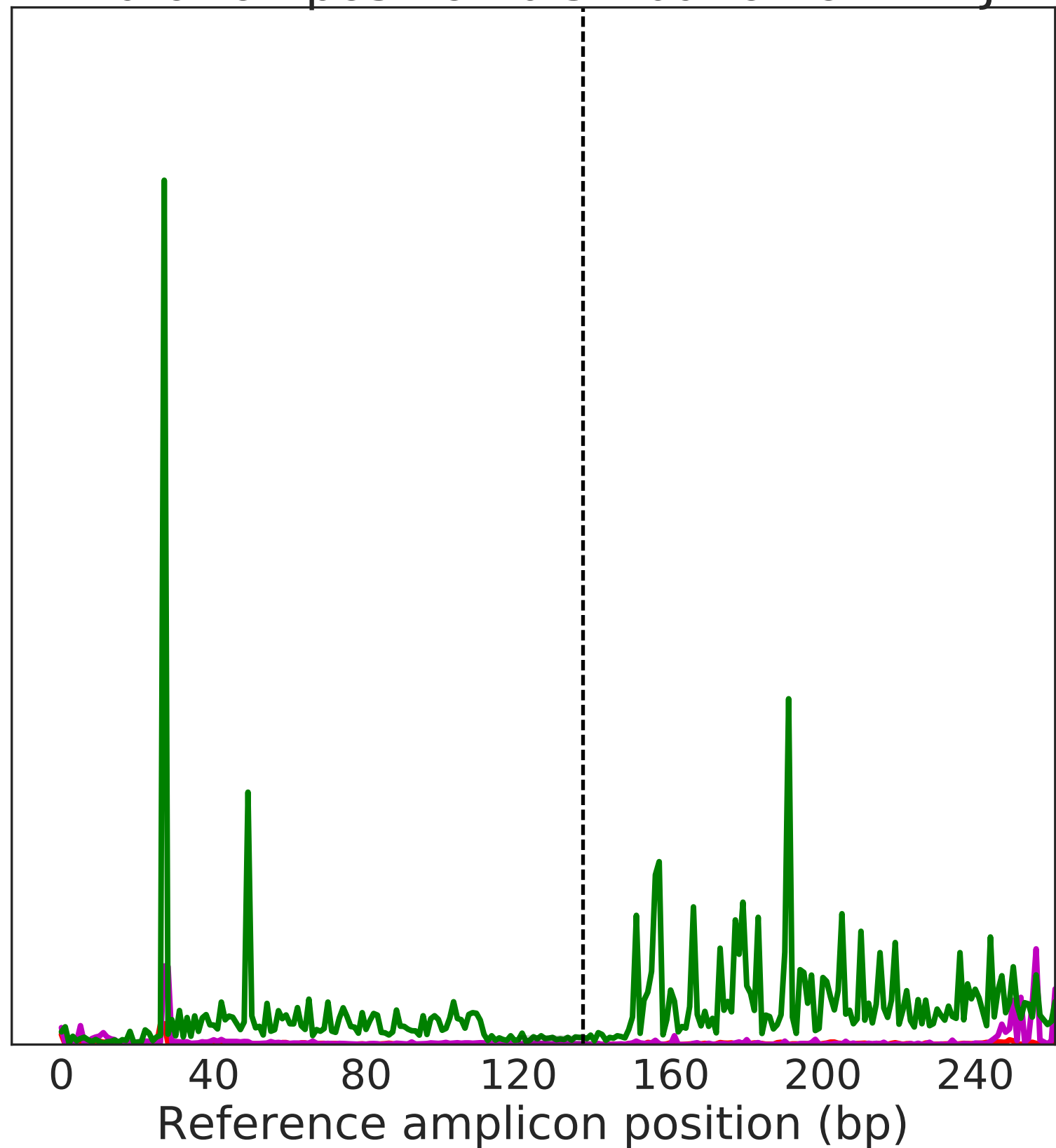
3.2% (23.8% , 4249)

2.4% (17.9% , 3187)

1.6% (11.9% , 2124)

0.8% (6.0% , 1062)

0.0% (0.0% , 0)



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