

# Mutation position distribution of NHEJ

Sequences: % Total ( % NHEJ, no. )

2.9% (19.6% , 3363)

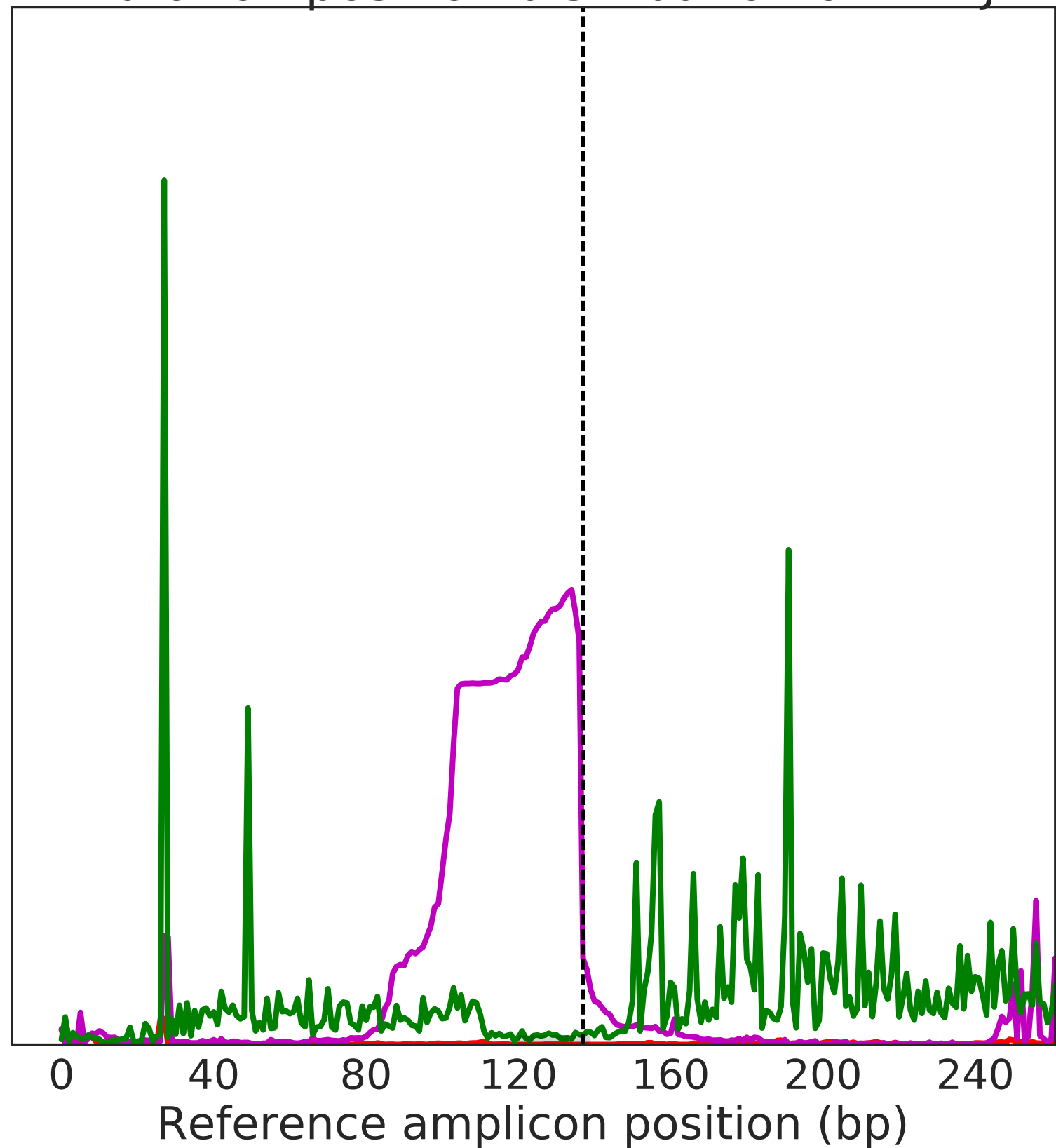
2.3% (15.6% , 2690)

1.7% (11.7% , 2017)

1.2% (7.8% , 1345)

0.6% (3.9% , 672)

0.0% (0.0% , 0)



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