

# Mutation position distribution of NHEJ

Sequences: % Total ( % NHEJ, no. )

6.1% (39.2% , 20949)

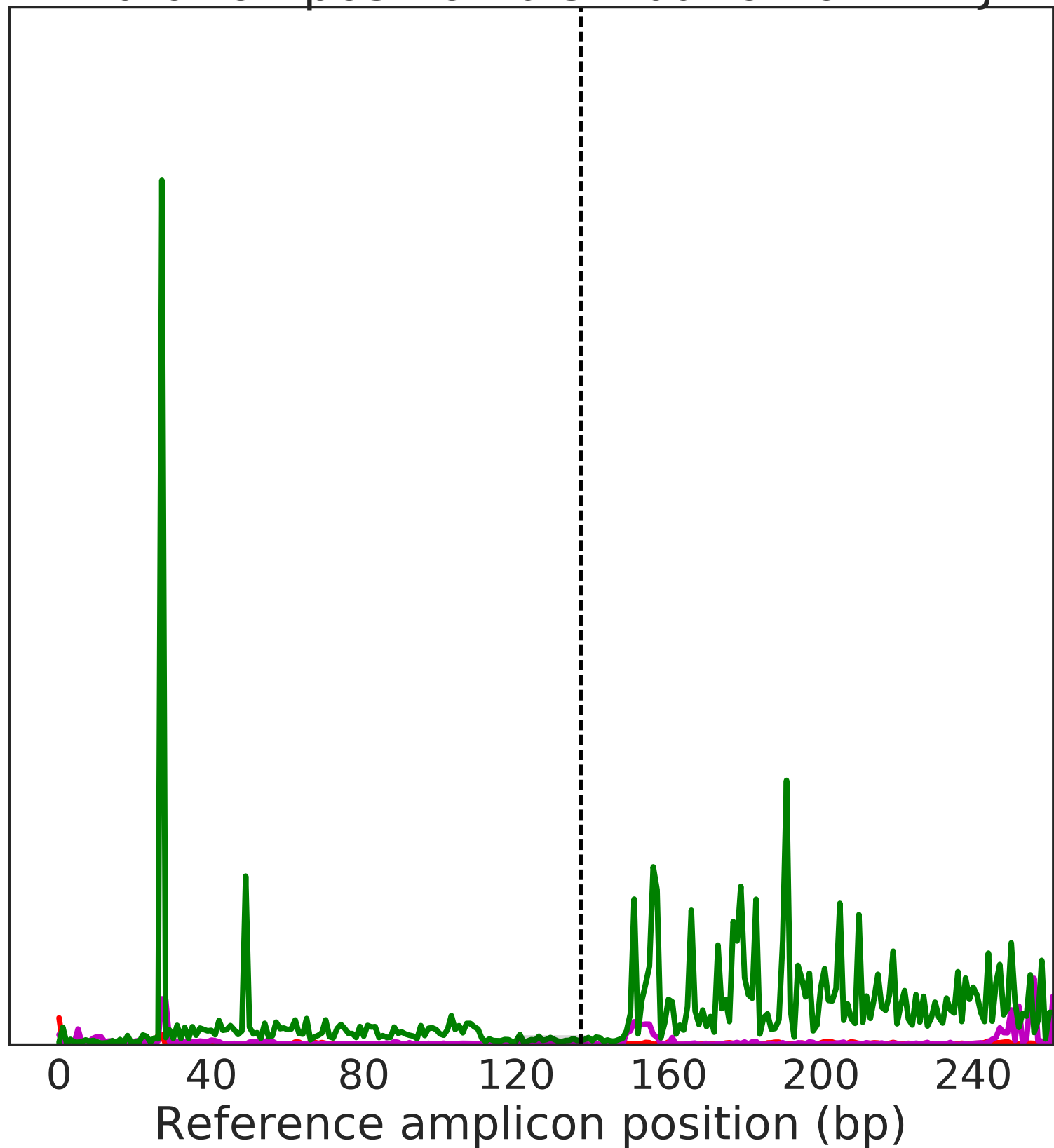
4.9% (31.4% , 16759)

3.7% (23.5% , 12569)

2.4% (15.7% , 8379)

1.2% (7.8% , 4189)

0.0% (0.0% , 0)



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