

Mutation position distribution of NHEJ

Sequences: % Total (% NHEJ, no.)

3.1% (18.0% , 18411)

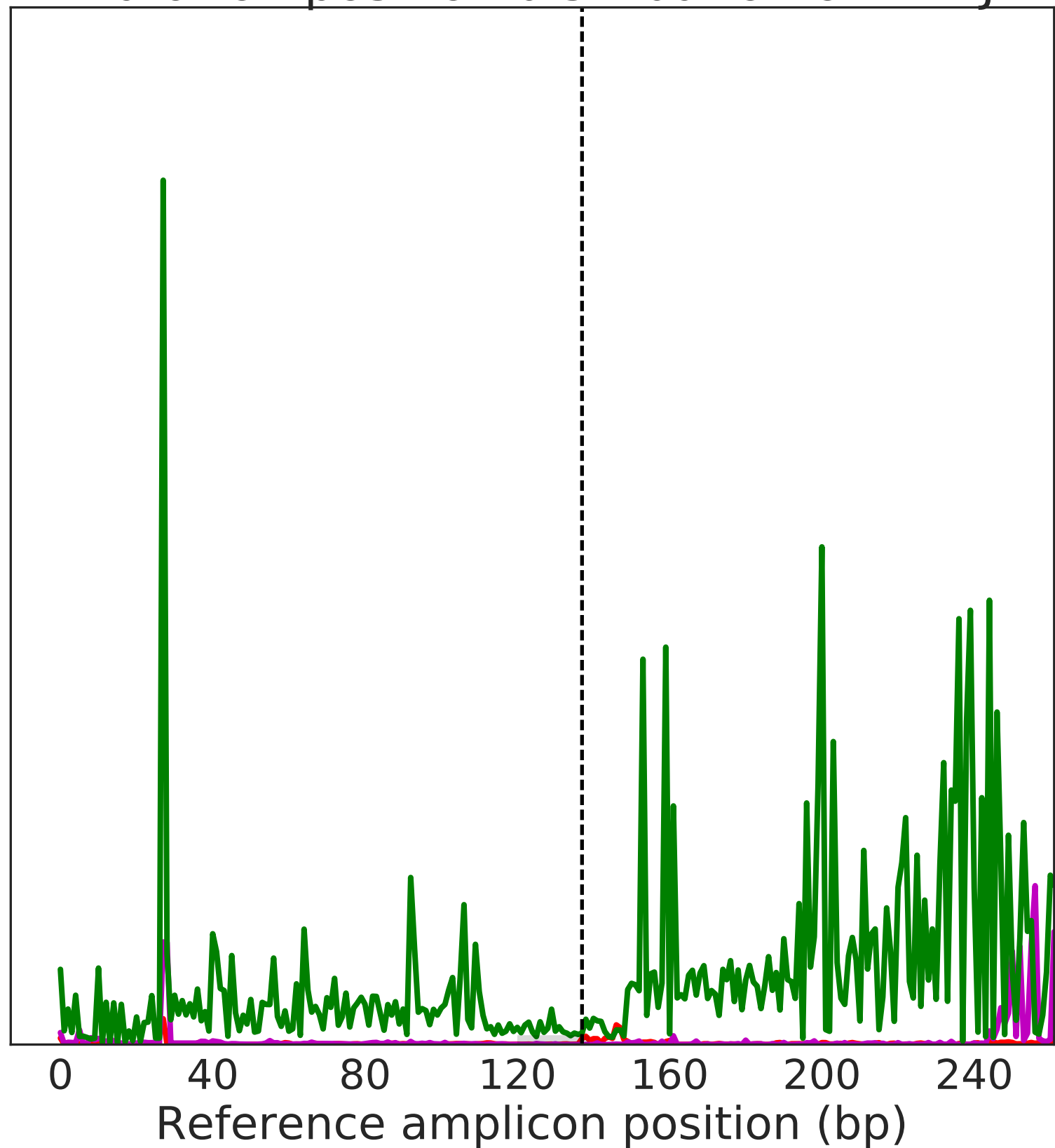
2.5% (14.4% , 14728)

1.9% (10.8% , 11046)

1.3% (7.2% , 7364)

0.6% (3.6% , 3682)

0.0% (0.0% , 0)



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