

Mutation position distribution of NHEJ

Sequences: % Total (% NHEJ, no.)

3.0% (18.7% , 25912)

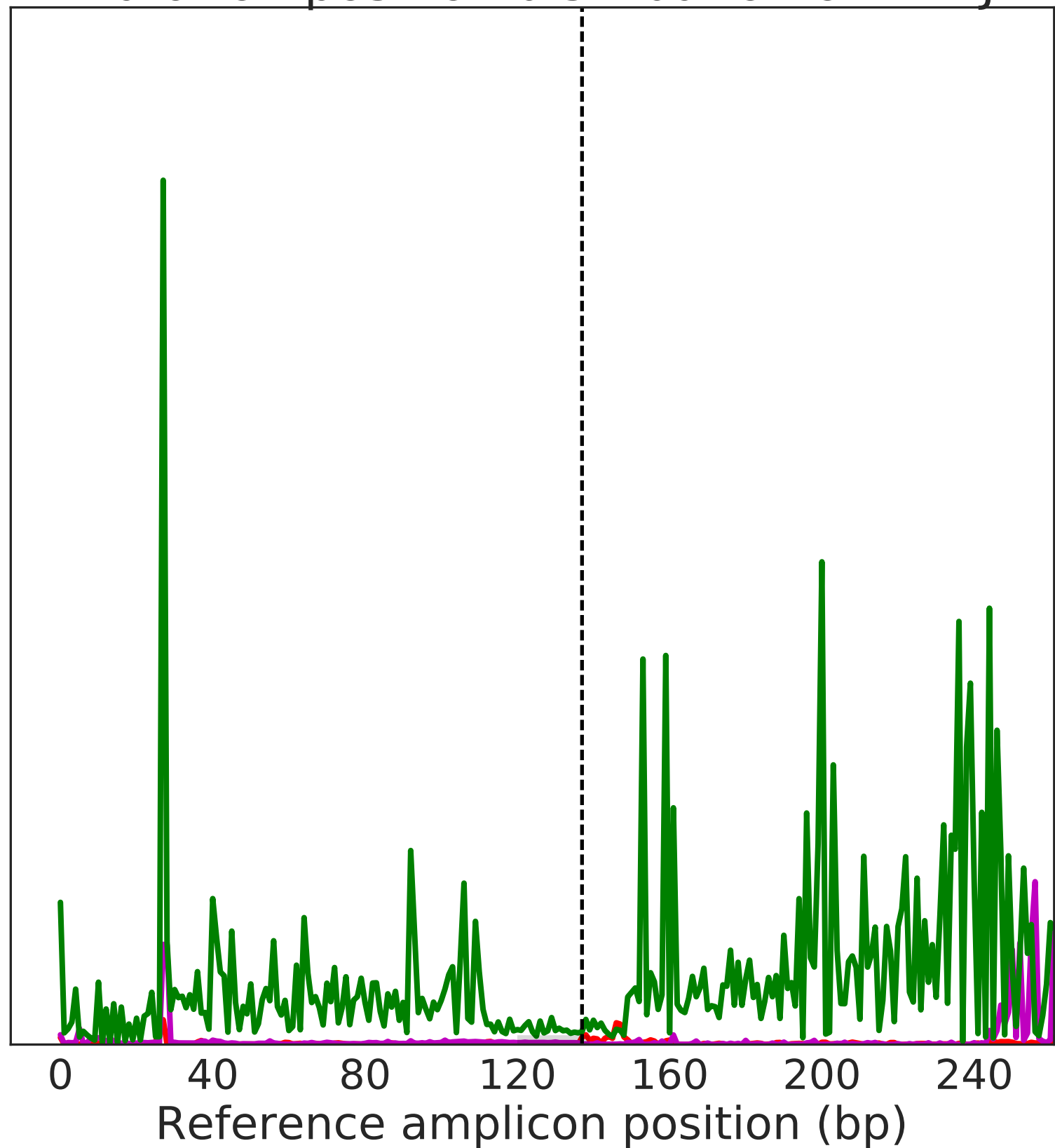
2.4% (14.9% , 20730)

1.8% (11.2% , 15547)

1.2% (7.5% , 10365)

0.6% (3.7% , 5182)

0.0% (0.0% , 0)



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