

Mutation position distribution of NHEJ

Sequences: % Total (% NHEJ, no.)

99.0% (99.0% , 102)

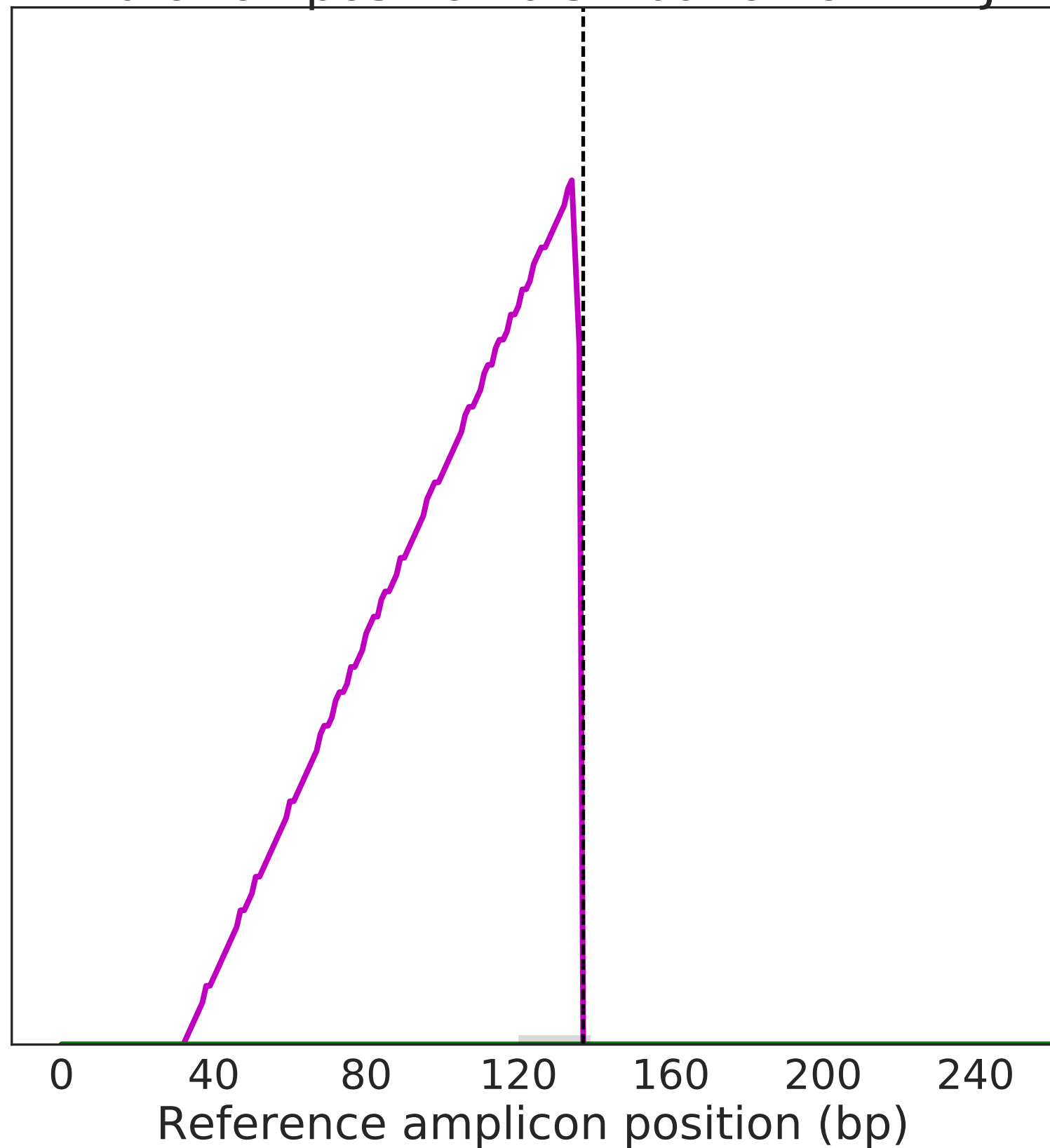
79.2% (79.2% , 82)

59.4% (59.4% , 61)

39.6% (39.6% , 41)

19.8% (19.8% , 20)

0.0% (0.0% , 0)



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