

Mutation position distribution of NHEJ

Sequences: % Total (% NHEJ, no.)

4.3% (29.1% , 14643)

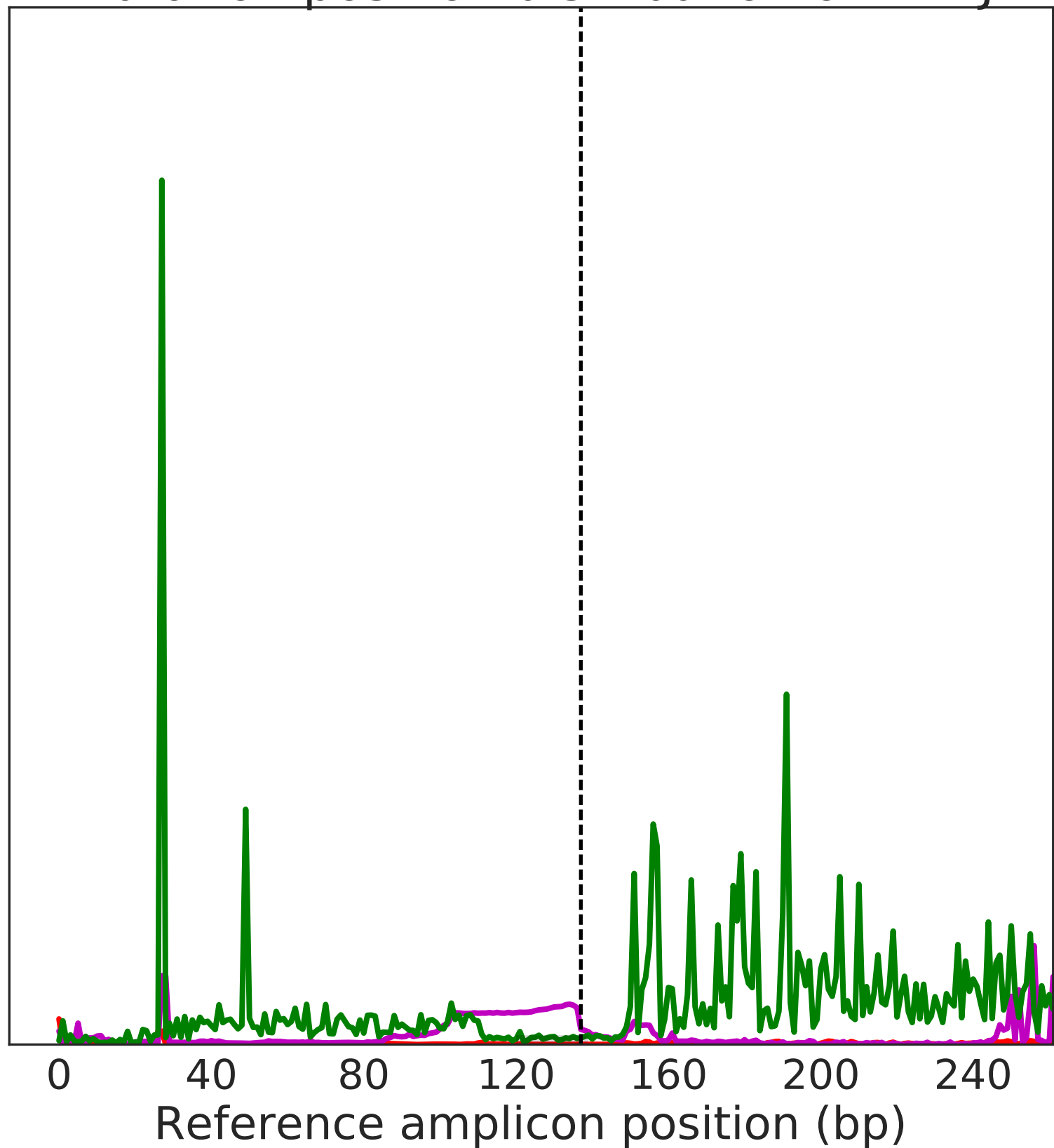
3.5% (23.3% , 11715)

2.6% (17.5% , 8786)

1.7% (11.6% , 5857)

0.9% (5.8% , 2928)

0.0% (0.0% , 0)



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