

Mutation position distribution of NHEJ

Sequences: % Total (% NHEJ, no.)

4.3% (29.3% , 13956)

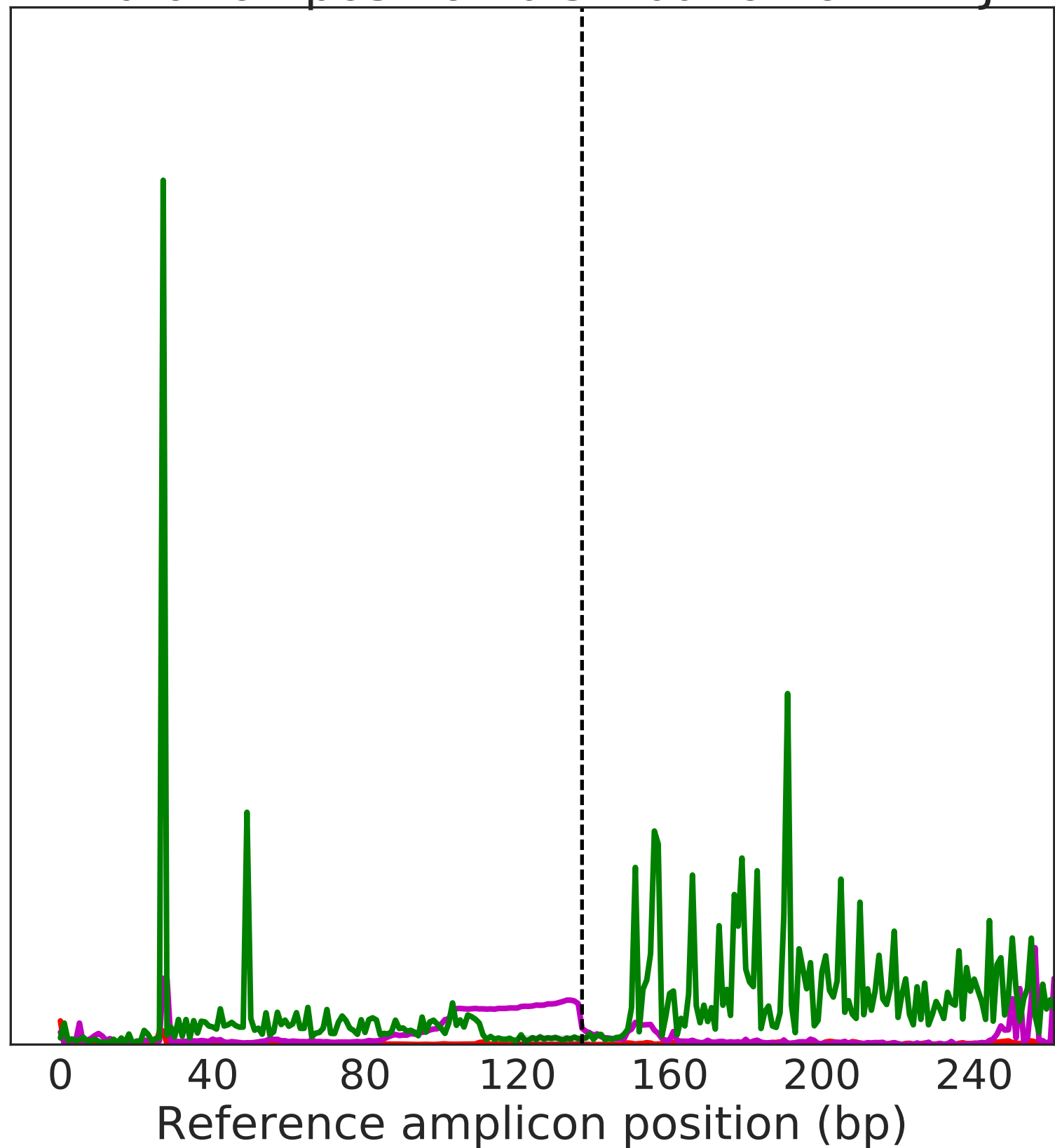
3.4% (23.4% , 11165)

2.6% (17.6% , 8374)

1.7% (11.7% , 5582)

0.9% (5.9% , 2791)

0.0% (0.0% , 0)



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