

Mutation position distribution of NHEJ

Sequences: % Total (% NHEJ, no.)

4.4% (30.5% , 14681)

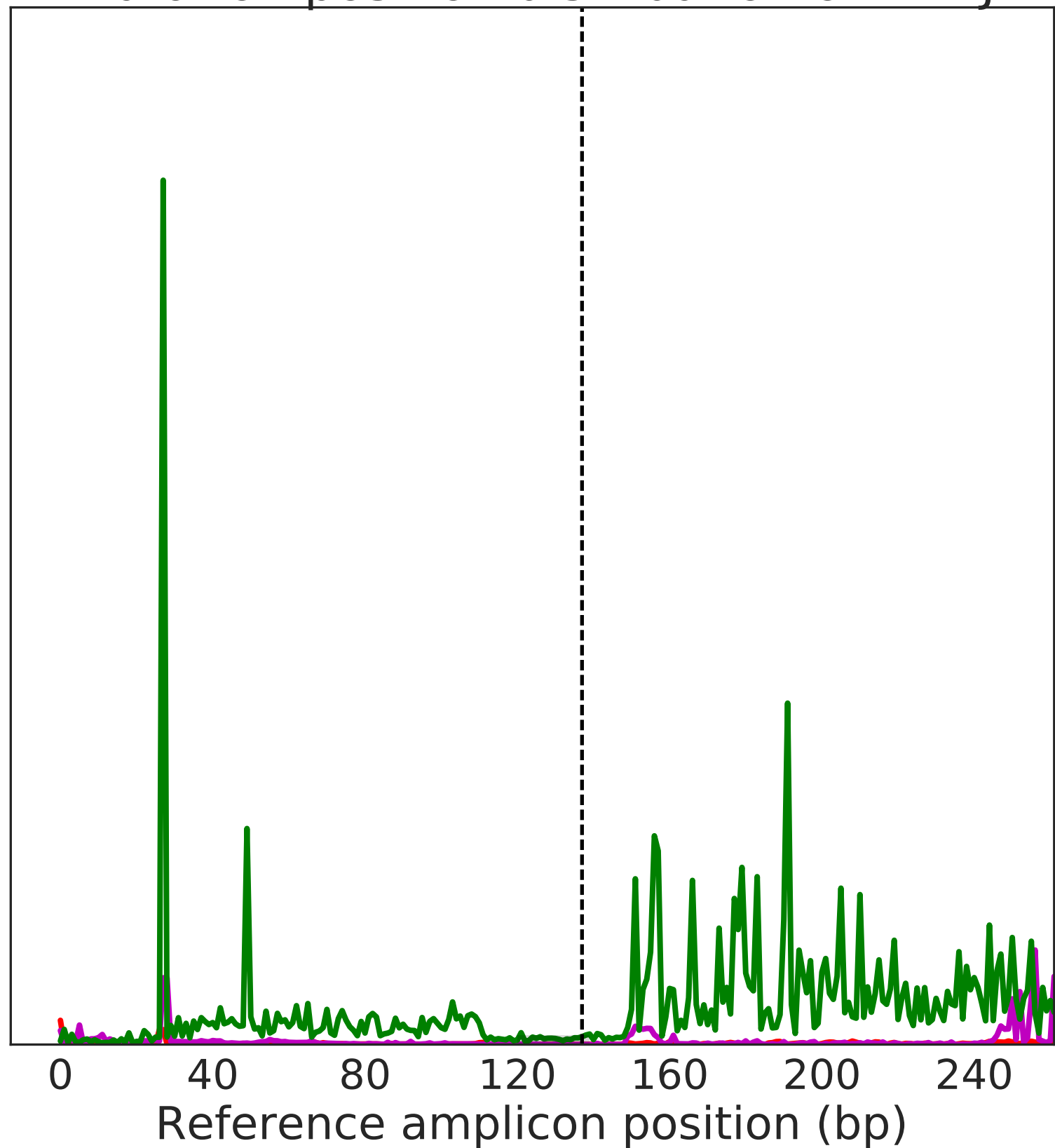
3.5% (24.4% , 11744)

2.6% (18.3% , 8808)

1.7% (12.2% , 5872)

0.9% (6.1% , 2936)

0.0% (0.0% , 0)



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