

Mutation position distribution of NHEJ

Sequences: % Total (% NHEJ, no.)

4.3% (30.0% , 14046)

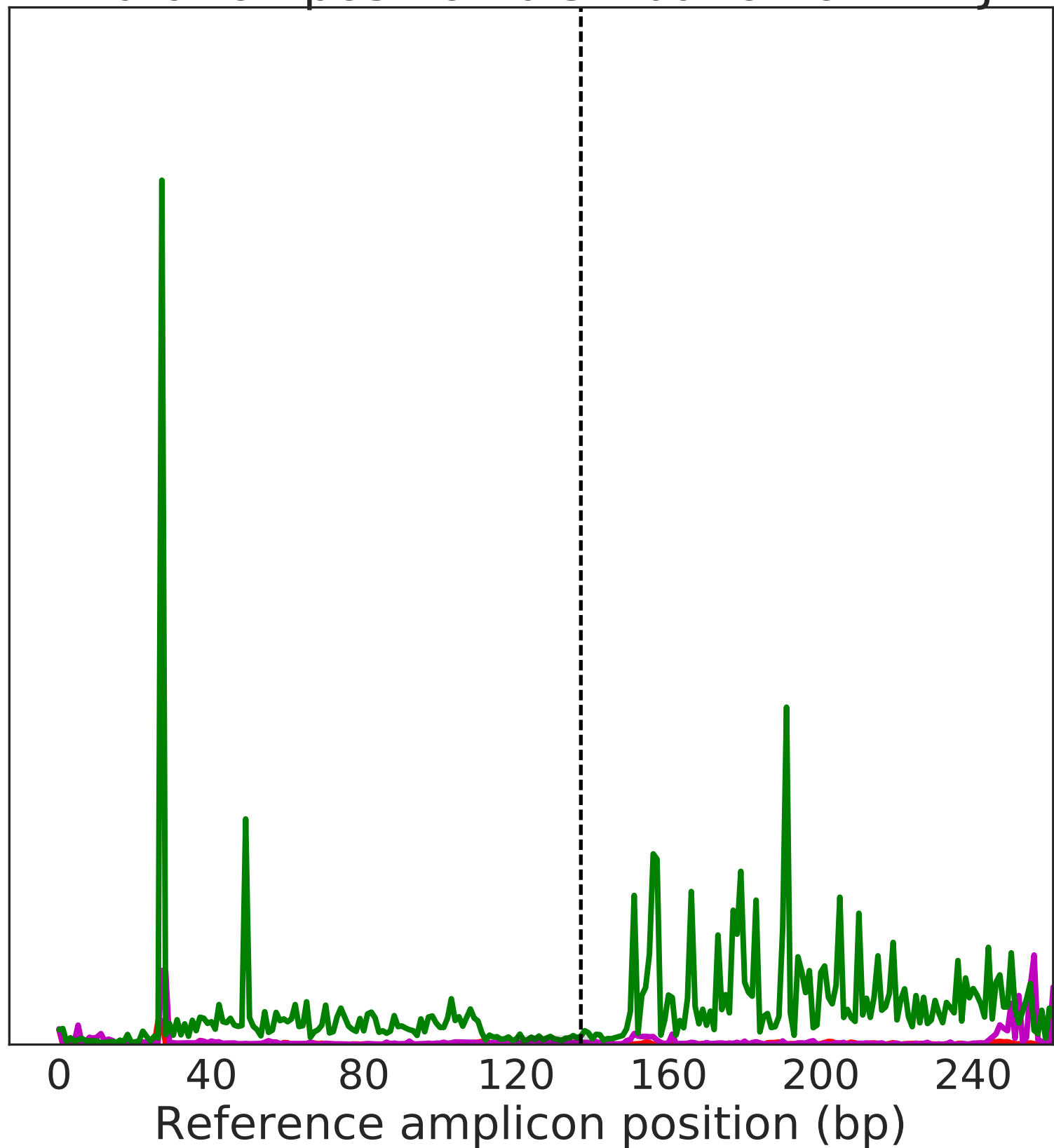
3.4% (24.0% , 11237)

2.6% (18.0% , 8428)

1.7% (12.0% , 5618)

0.9% (6.0% , 2809)

0.0% (0.0% , 0)



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