

Mutation position distribution of NHEJ

Sequences: % Total (% NHEJ, no.)

4.5% (29.0% , 22885)

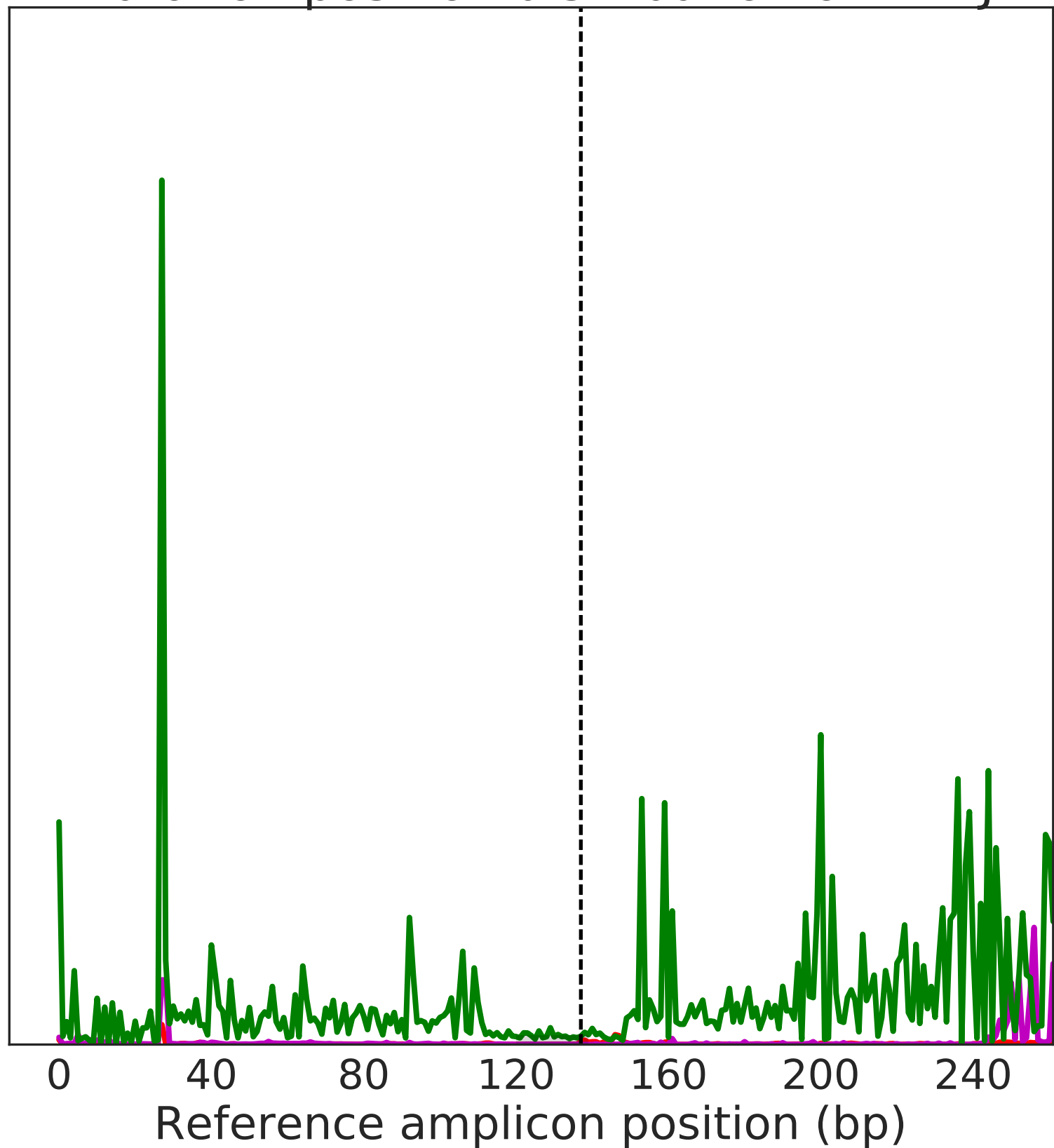
3.6% (23.2% , 18308)

2.7% (17.4% , 13731)

1.8% (11.6% , 9154)

0.9% (5.8% , 4577)

0.0% (0.0% , 0)



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