

Mutation position distribution of NHEJ

Sequences: % Total (% NHEJ, no.)

4.3% (29.2% , 18700)

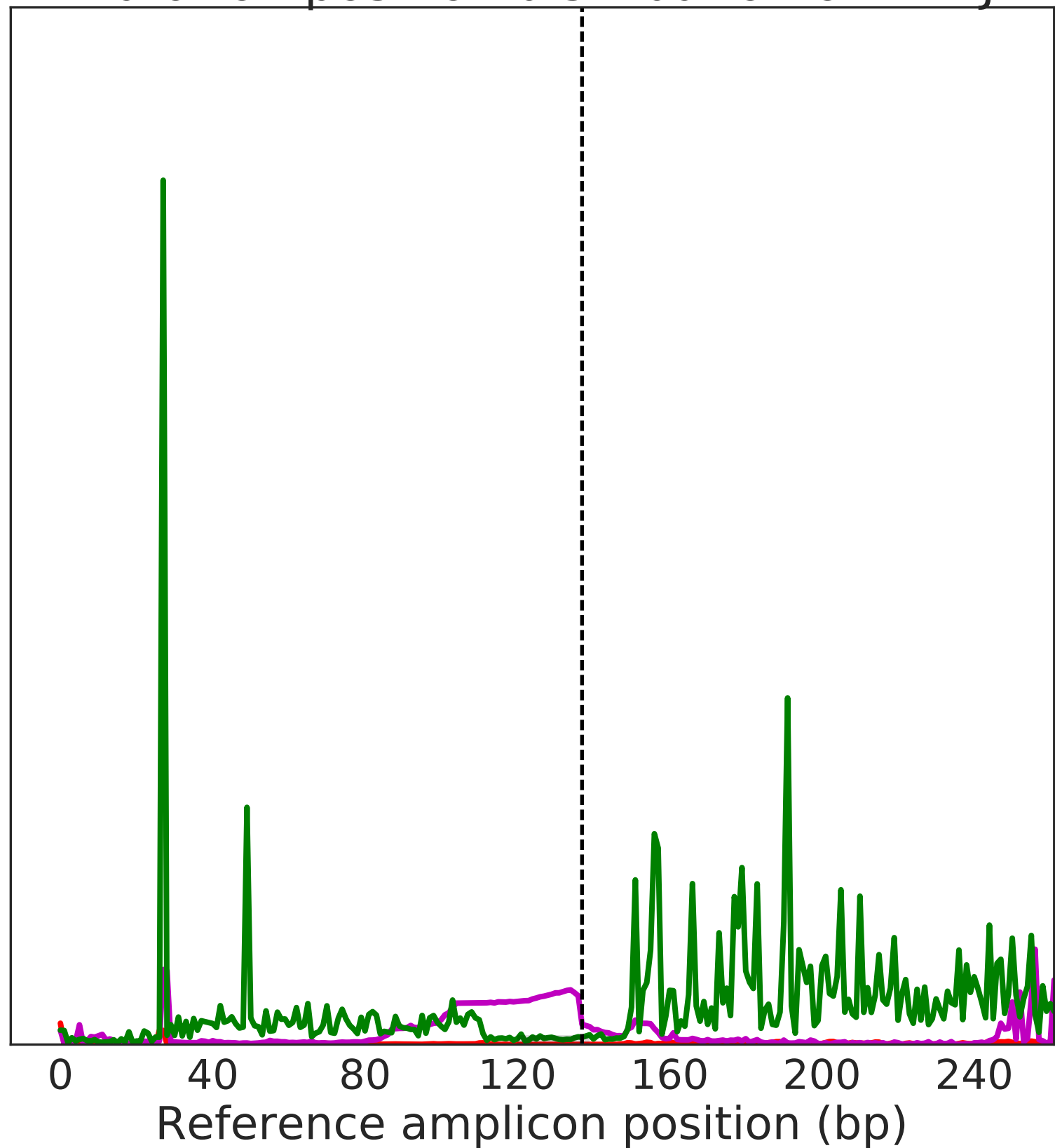
3.4% (23.4% , 14960)

2.6% (17.5% , 11220)

1.7% (11.7% , 7480)

0.9% (5.8% , 3740)

0.0% (0.0% , 0)



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