

Mutation position distribution of NHEJ

Sequences: % Total (% NHEJ, no.)

5.9% (45.2% , 7130)

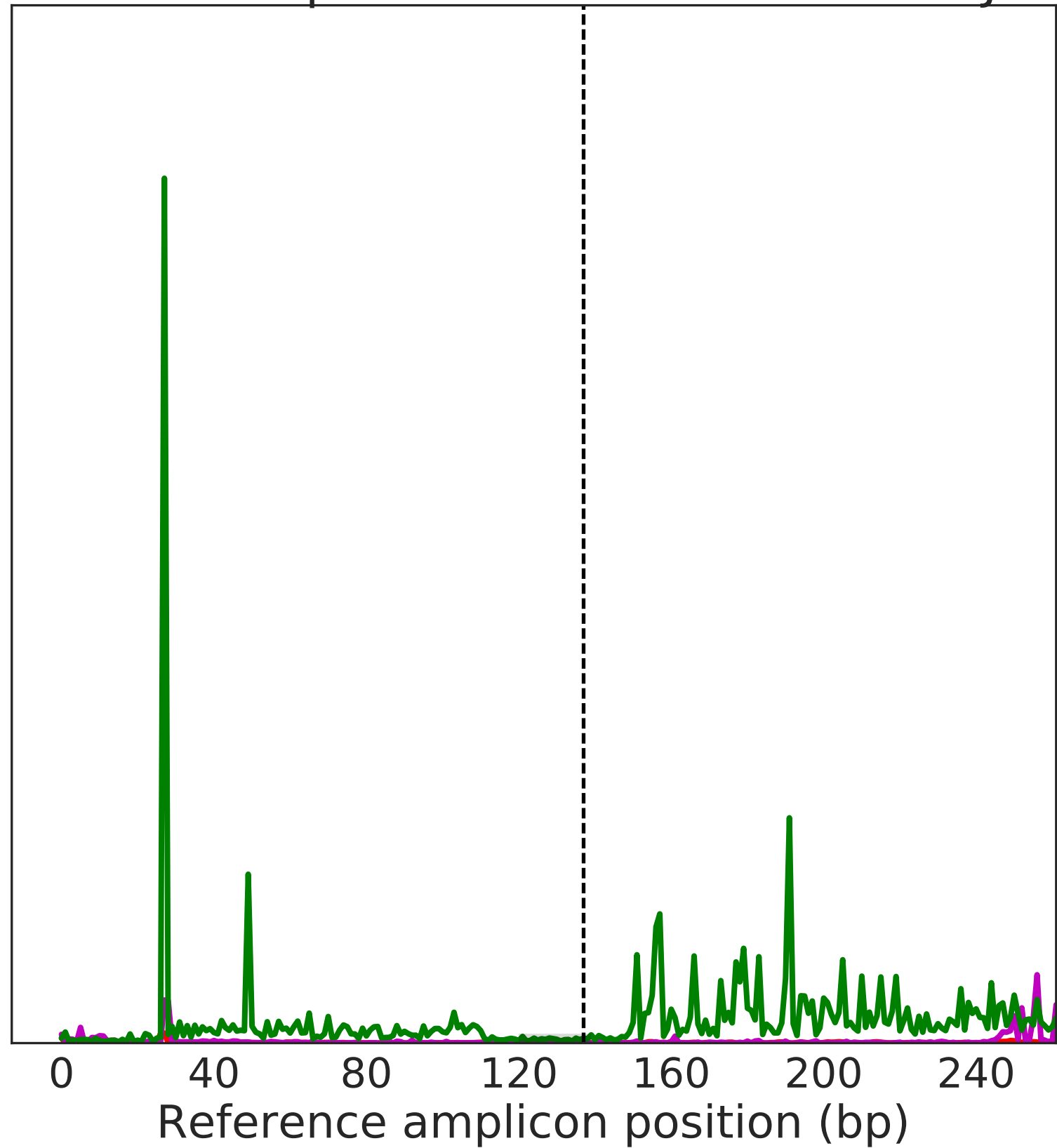
4.7% (36.2% , 5704)

3.5% (27.1% , 4278)

2.3% (18.1% , 2852)

1.2% (9.0% , 1426)

0.0% (0.0% , 0)



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