

Mutation position distribution of NHEJ

Sequences: % Total (% NHEJ, no.)

3.1% (18.9% , 20030)

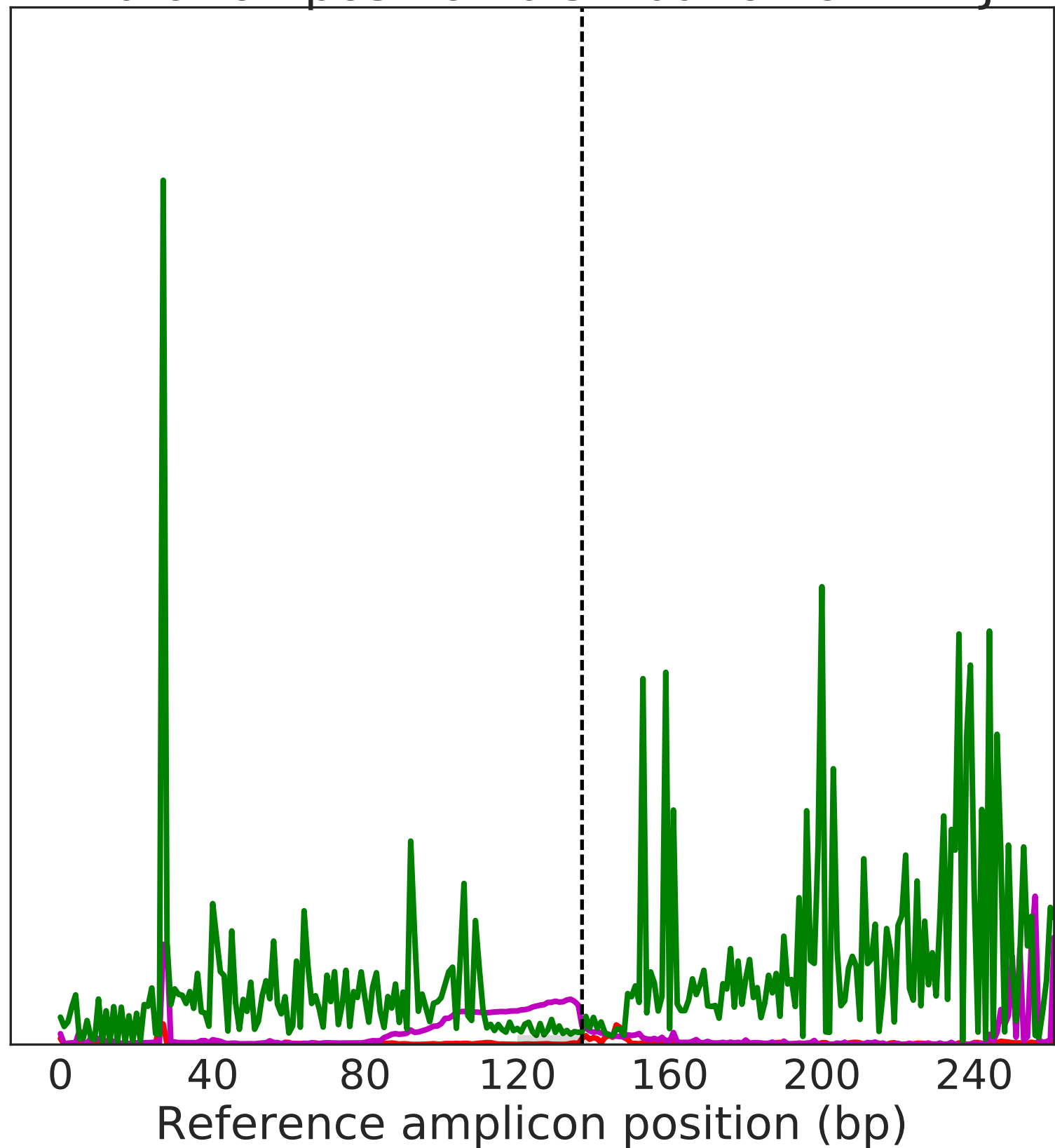
2.5% (15.1% , 16024)

1.8% (11.3% , 12018)

1.2% (7.5% , 8012)

0.6% (3.8% , 4006)

0.0% (0.0% , 0)



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