

Mutation position distribution of NHEJ

Sequences: % Total (% NHEJ, no.)

2.8% (18.7% , 20760)

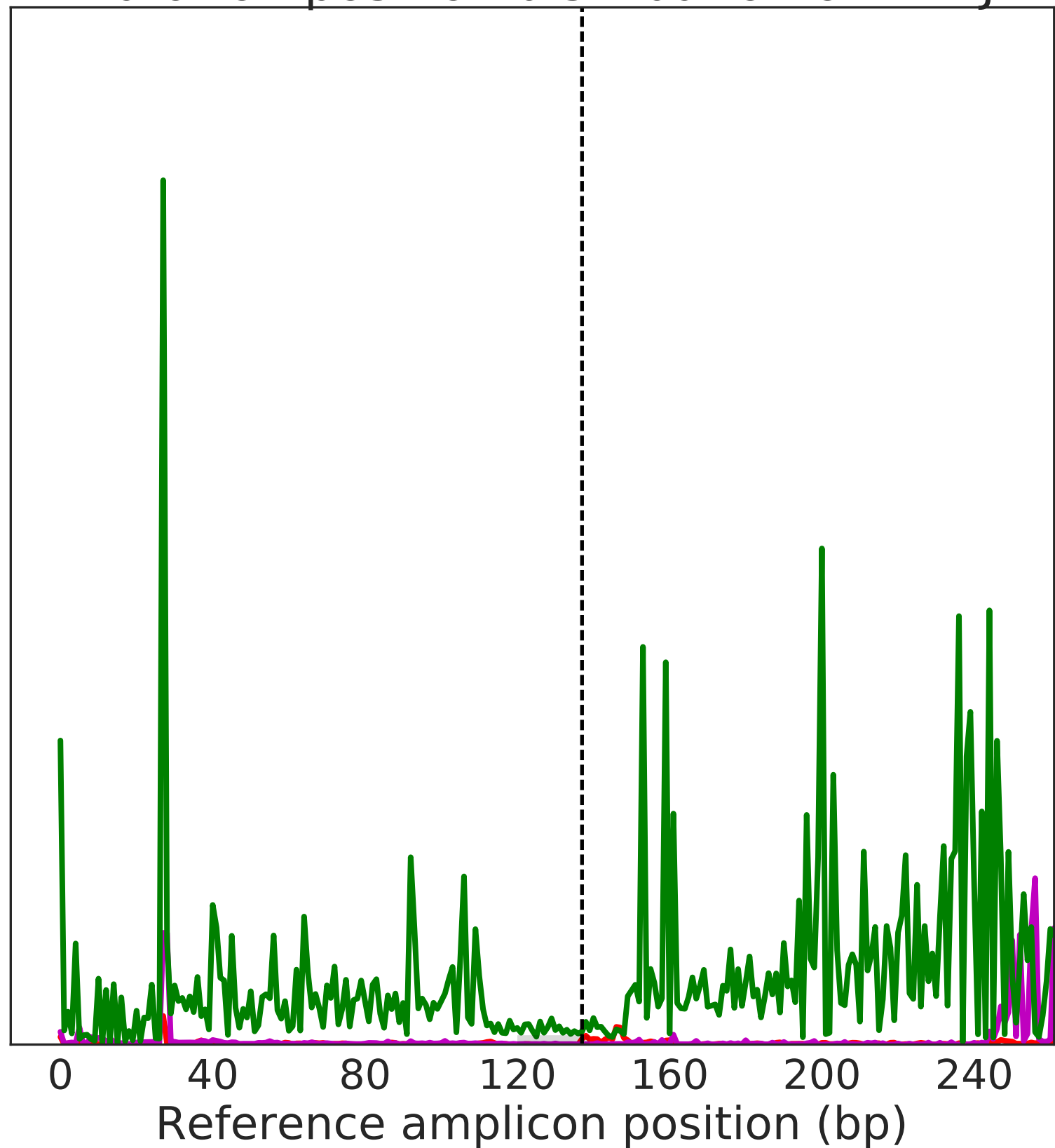
2.3% (15.0% , 16608)

1.7% (11.2% , 12456)

1.1% (7.5% , 8304)

0.6% (3.7% , 4152)

0.0% (0.0% , 0)



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