

Mutation position distribution of NHEJ

Sequences: % Total (% NHEJ, no.)

3.4% (18.5% , 21211)

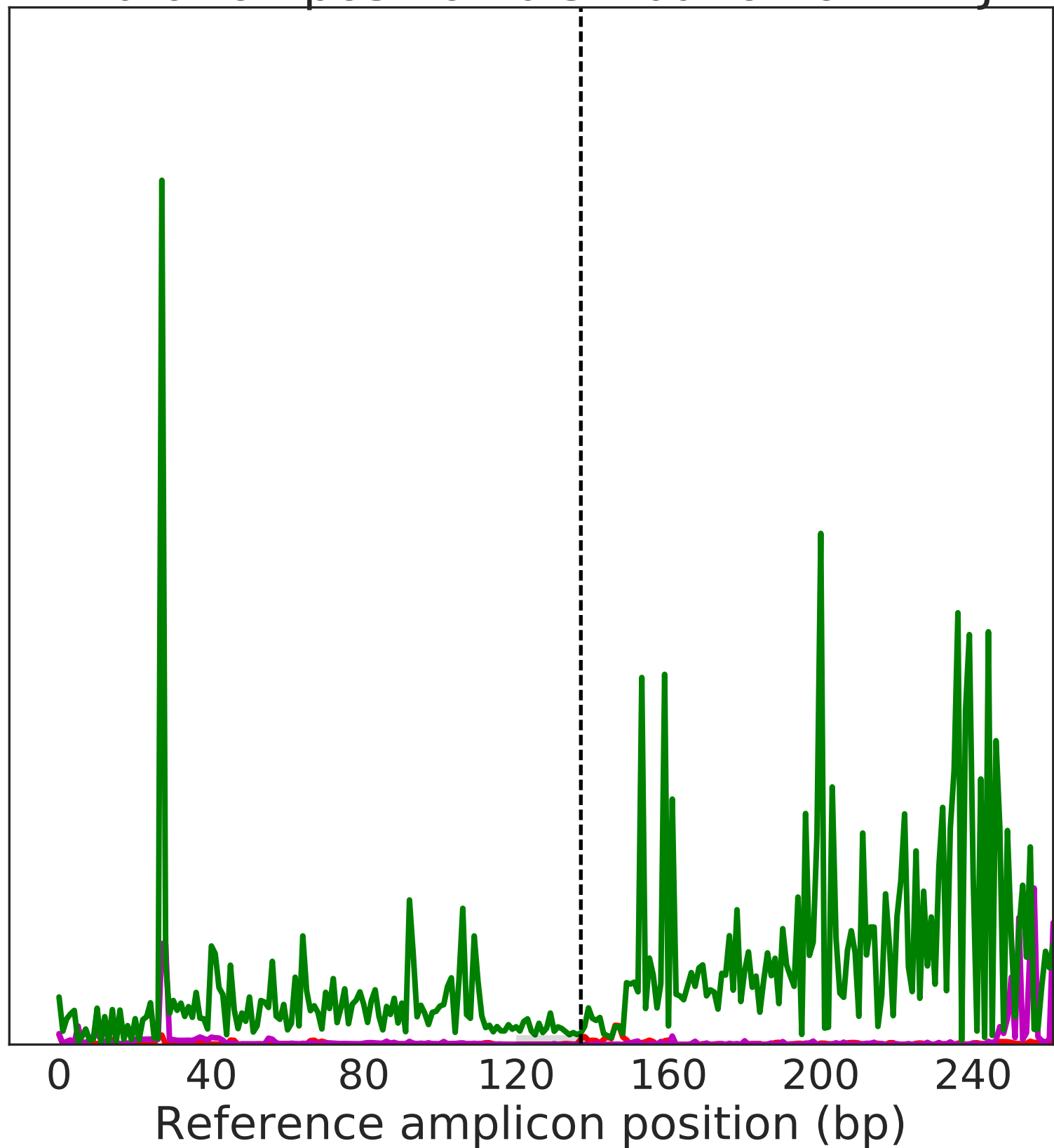
2.7% (14.8% , 16968)

2.0% (11.1% , 12726)

1.3% (7.4% , 8484)

0.7% (3.7% , 4242)

0.0% (0.0% , 0)



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