

Mutation position distribution of NHEJ

Sequences: % Total (% NHEJ, no.)

3.9% (28.9% , 4513)

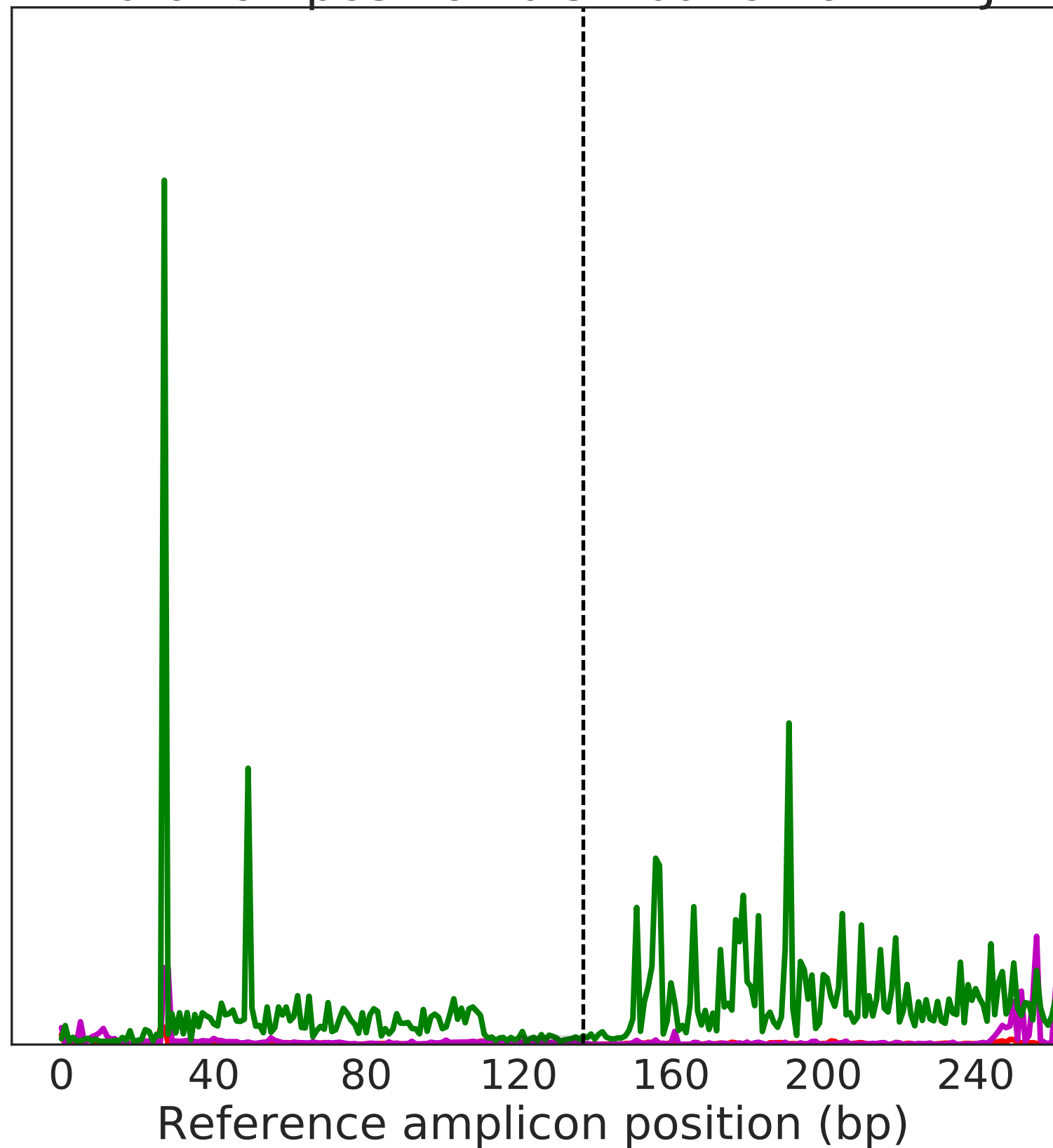
3.1% (23.1% , 3610)

2.3% (17.3% , 2707)

1.6% (11.6% , 1805)

0.8% (5.8% , 902)

0.0% (0.0% , 0)



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