

Mutation position distribution of NHEJ

Sequences: % Total (% NHEJ, no.)

2.4% (16.9% , 8014)

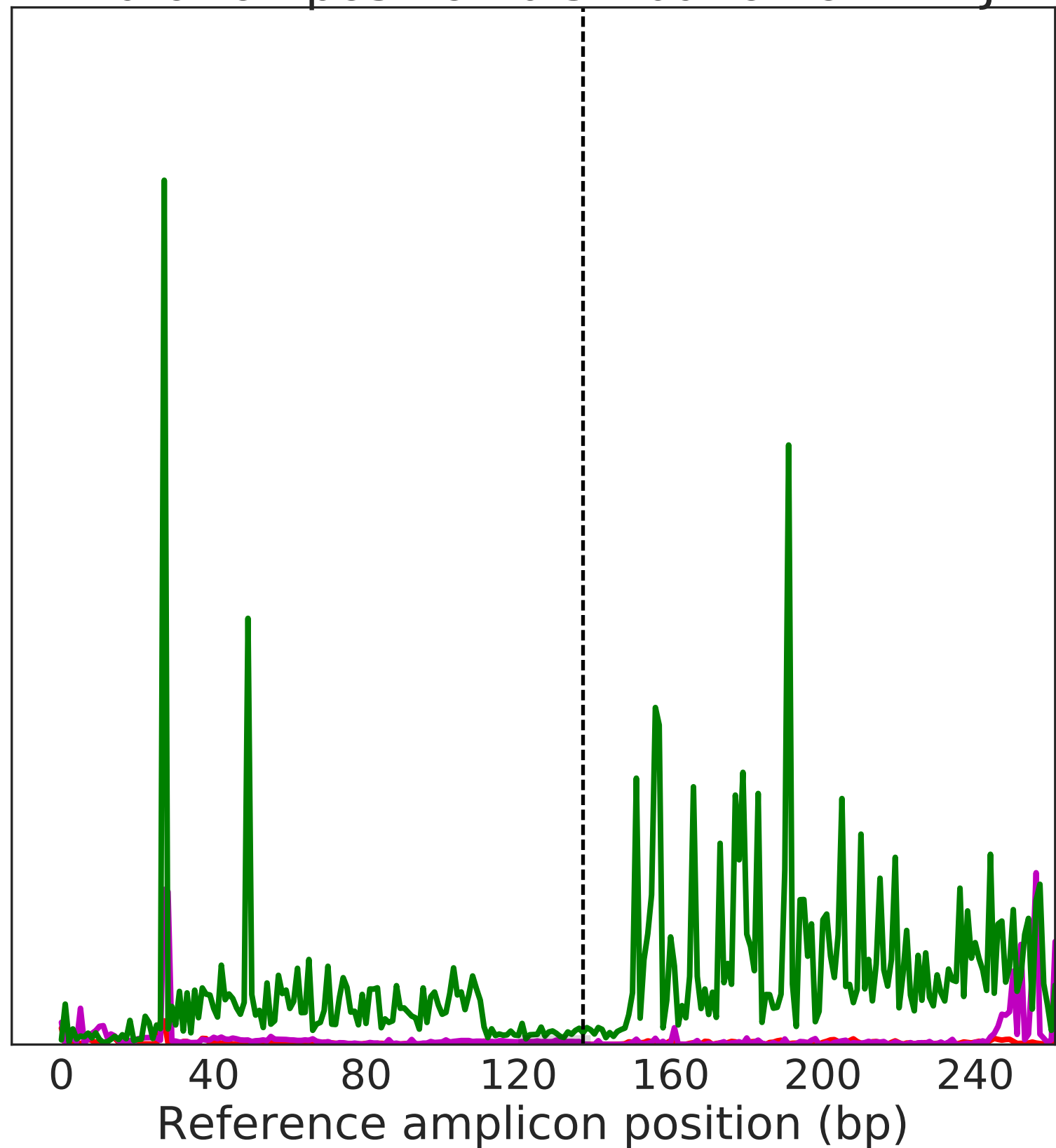
1.9% (13.5% , 6411)

1.4% (10.2% , 4808)

0.9% (6.8% , 3205)

0.5% (3.4% , 1602)

0.0% (0.0% , 0)



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