

Mutation position distribution of NHEJ

Sequences: % Total (% NHEJ, no.)

67.7% (86.5% , 146970)

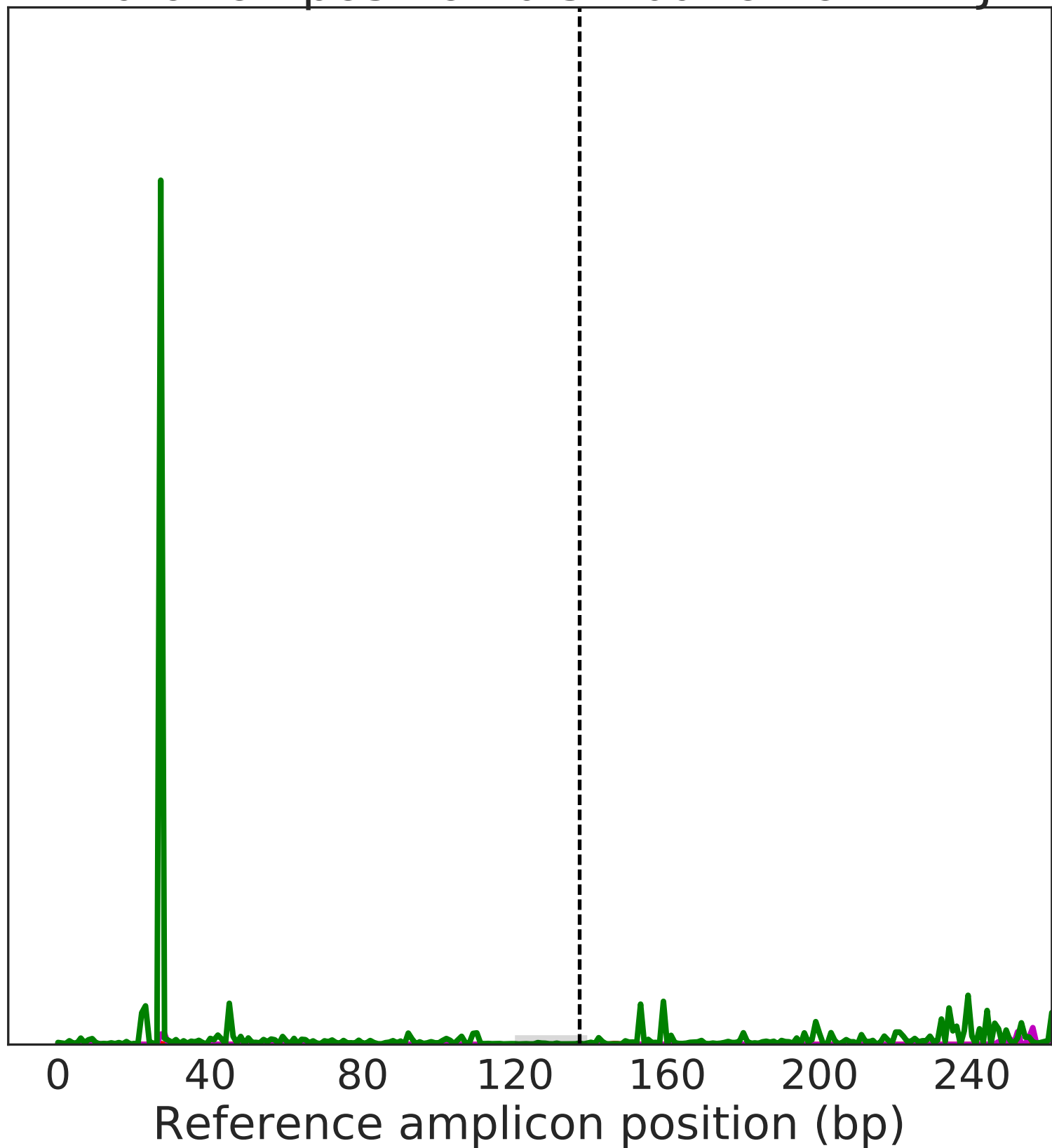
54.1% (69.2% , 117576)

40.6% (51.9% , 88182)

27.1% (34.6% , 58788)

13.5% (17.3% , 29394)

0.0% (0.0% , 0)



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