

# Mutation position distribution of NHEJ

Sequences: % Total ( % NHEJ, no. )

56.4% (77.7% , 83296)

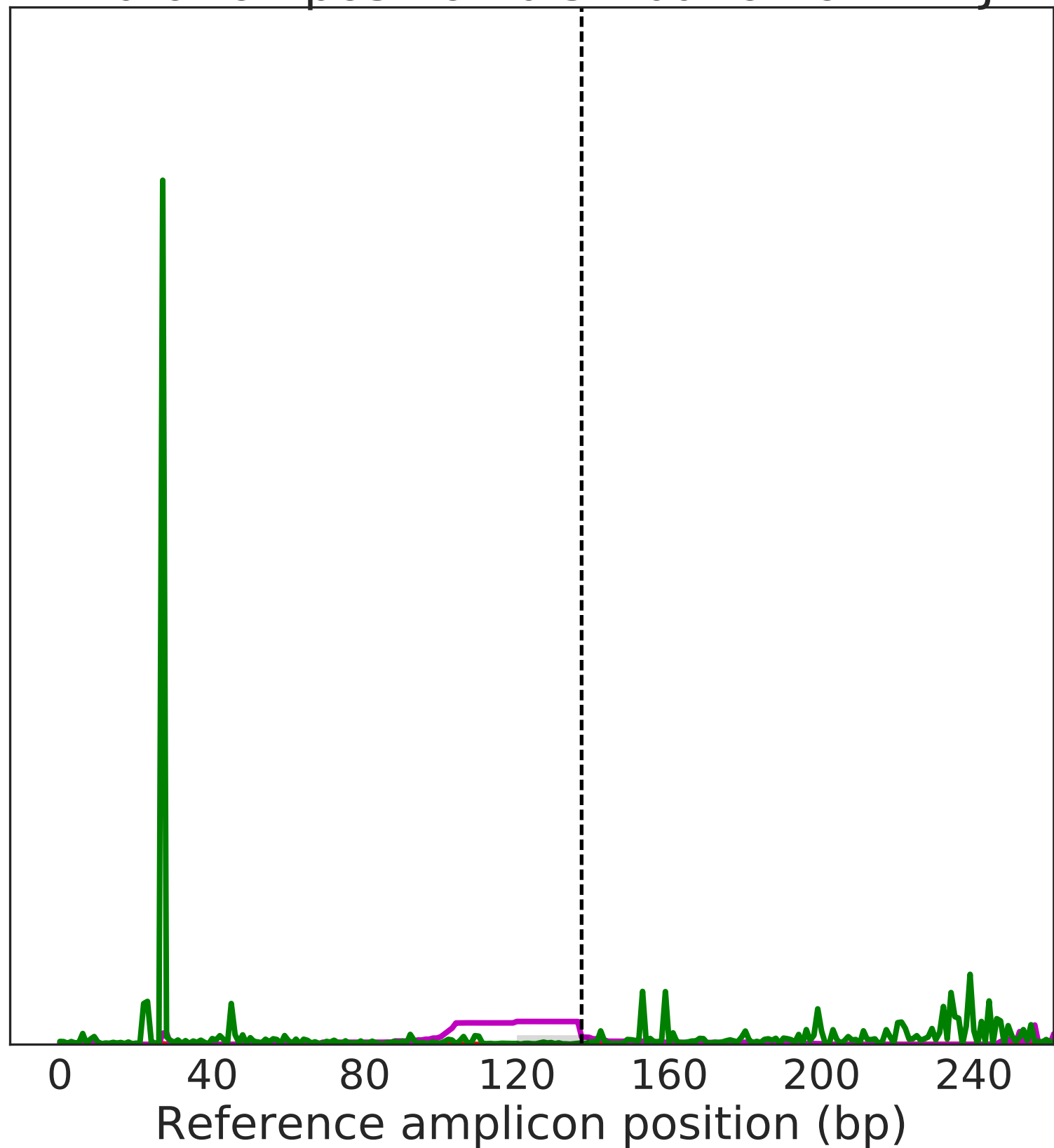
45.1% (62.2% , 66636)

33.8% (46.6% , 49977)

22.5% (31.1% , 33318)

11.3% (15.5% , 16659)

0.0% (0.0% , 0)



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