

# Mutation position distribution of NHEJ

Sequences: % Total ( % NHEJ, no. )

62.9% (77.5% , 114079)

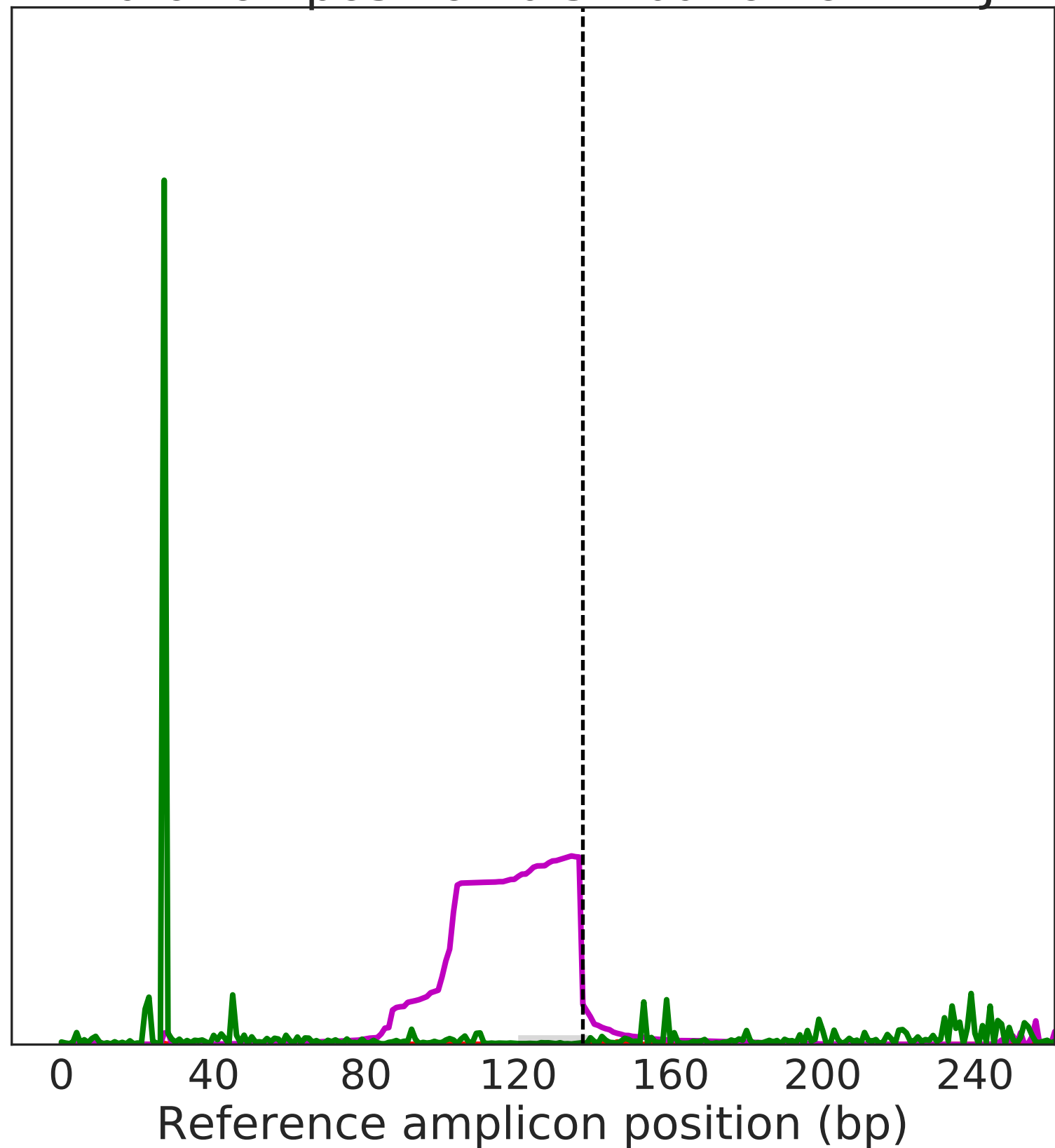
50.3% (62.0% , 91263)

37.7% (46.5% , 68447)

25.1% (31.0% , 45631)

12.6% (15.5% , 22815)

0.0% (0.0% , 0)



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