

Mutation position distribution of NHEJ

Sequences: % Total (% NHEJ, no.)

69.8% (82.6% , 152392)

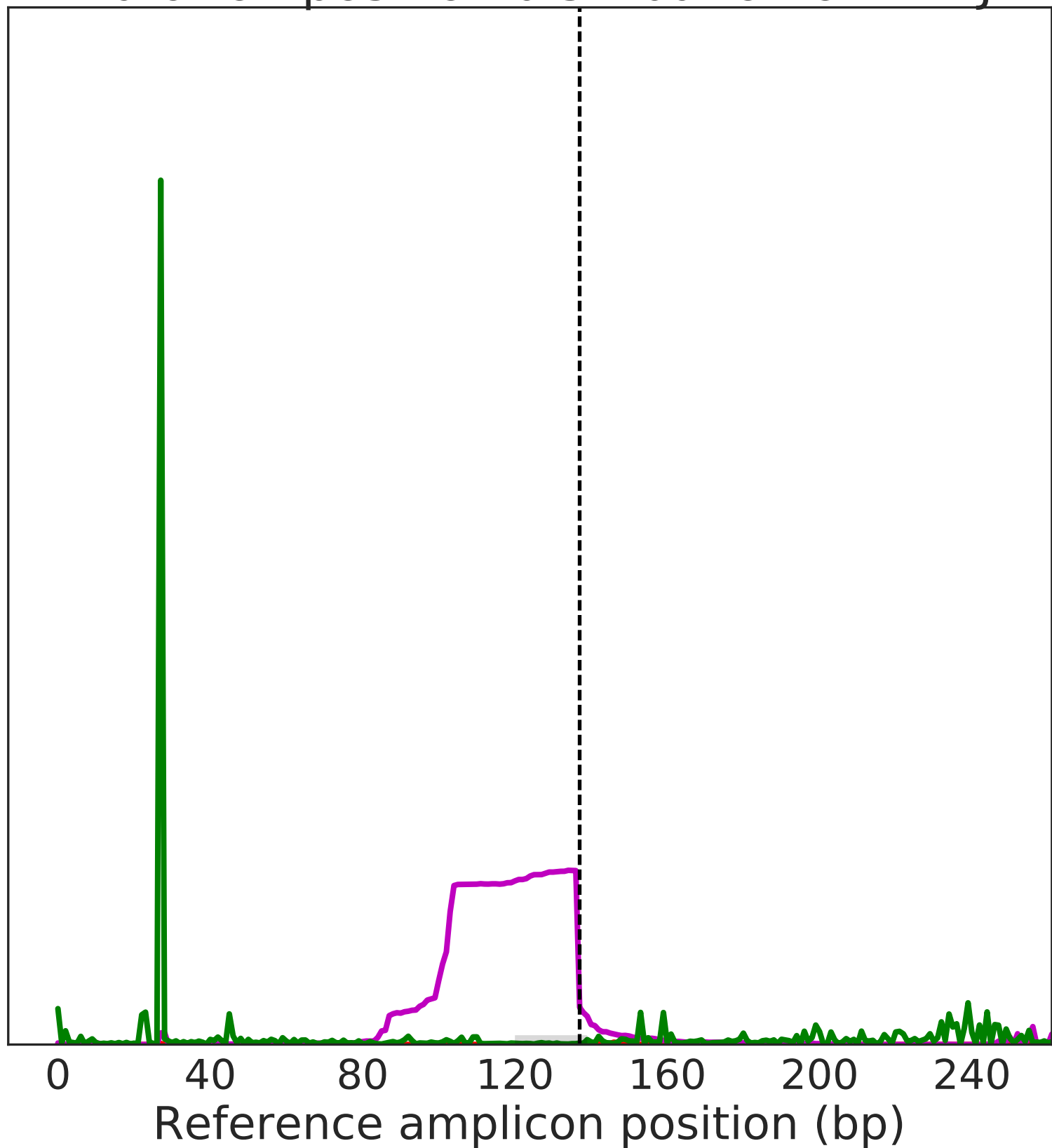
55.8% (66.1% , 121913)

41.9% (49.6% , 91435)

27.9% (33.0% , 60956)

14.0% (16.5% , 30478)

0.0% (0.0% , 0)



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