

Mutation position distribution of NHEJ

Sequences: % Total (% NHEJ, no.)

68.0% (86.0% , 185833)

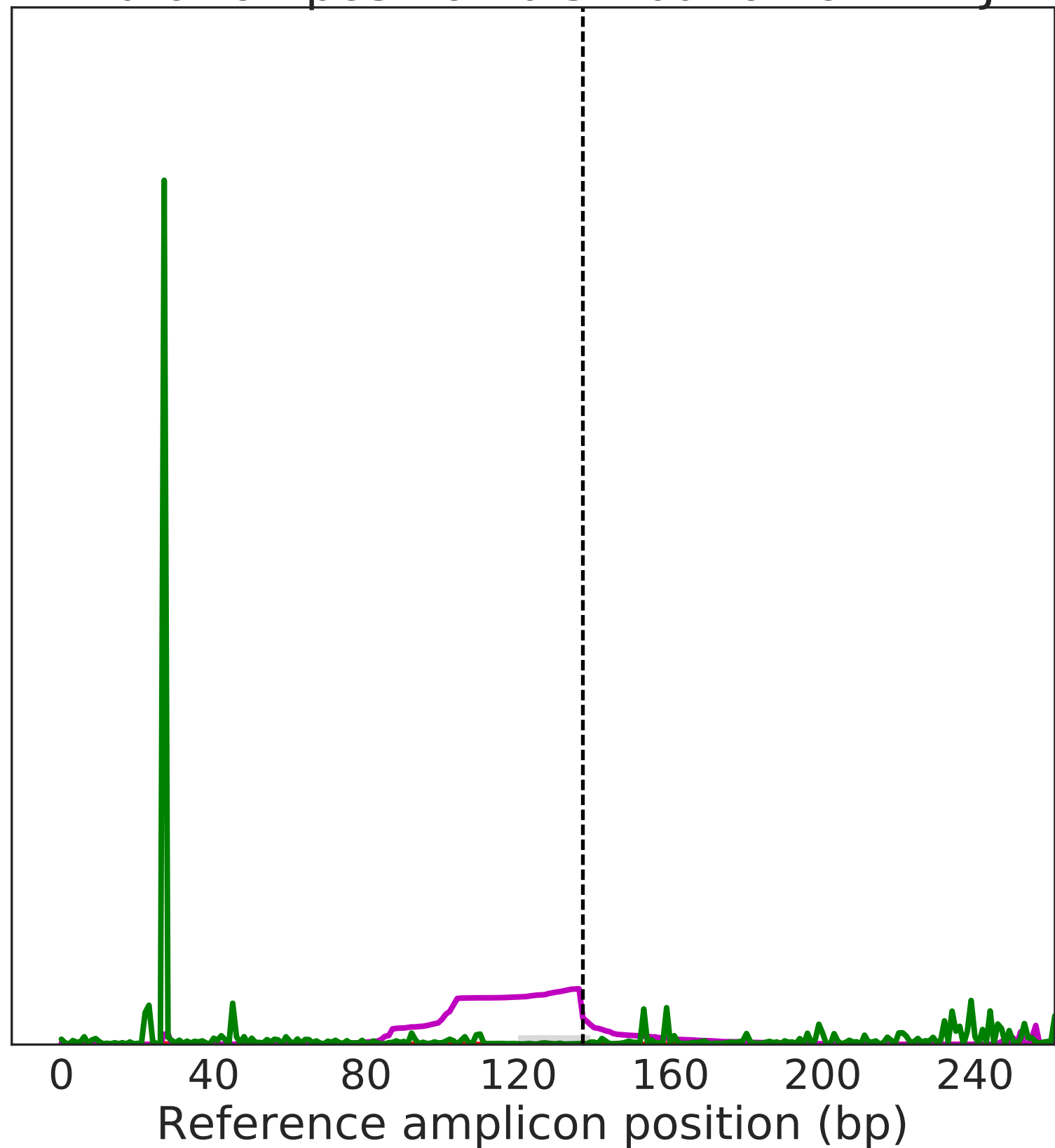
54.4% (68.8% , 148666)

40.8% (51.6% , 111499)

27.2% (34.4% , 74333)

13.6% (17.2% , 37166)

0.0% (0.0% , 0)



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