

Mutation position distribution of NHEJ

Sequences: % Total (% NHEJ, no.)

10.2% (57.5% , 59190)

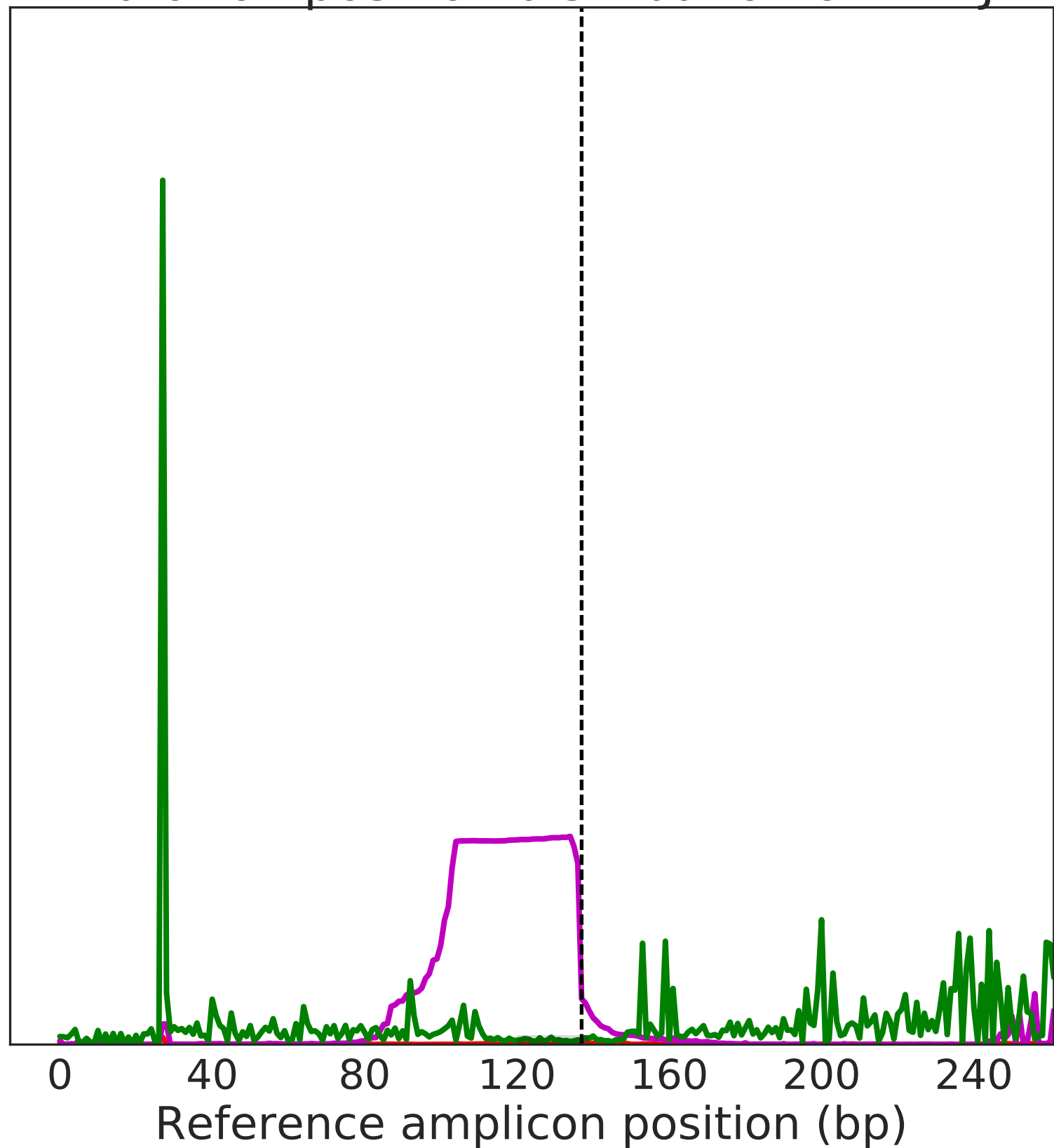
8.2% (46.0% , 47352)

6.1% (34.5% , 35514)

4.1% (23.0% , 23676)

2.0% (11.5% , 11838)

0.0% (0.0% , 0)



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