

Mutation position distribution of NHEJ

Sequences: % Total (% NHEJ, no.)

5.0% (35.3% , 11445)

4.0% (28.3% , 9156)

3.0% (21.2% , 6867)

2.0% (14.1% , 4578)

1.0% (7.1% , 2289)

0.0% (0.0% , 0)

0 40 80 120 160 200 240

Reference amplicon position (bp)

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

