

Mutation position distribution of NHEJ

Sequences: % Total (% NHEJ, no.)

4.0% (28.7% , 12206)

3.2% (22.9% , 9764)

2.4% (17.2% , 7323)

1.6% (11.5% , 4882)

0.8% (5.7% , 2441)

0.0% (0.0% , 0)

0 40 80 120 160 200 240

Reference amplicon position (bp)

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

