

# Mutation position distribution of NHEJ

Sequences: % Total ( % NHEJ, no. )

3.9% (26.0% , 11562)

3.1% (20.8% , 9249)

2.3% (15.6% , 6937)

1.5% (10.4% , 4624)

0.8% (5.2% , 2312)

0.0% (0.0% , 0)

0 40 80 120 160 200 240

Reference amplicon position (bp)

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

