

Mutation position distribution of NHEJ

Sequences: % Total (% NHEJ, no.)

3.3% (inf% , 0)

2.7% (inf% , 0)

2.0% (inf% , 0)

1.3% (inf% , 0)

0.7% (inf% , 0)

0.0% (nan% , 0)

0

3

6

9

12

15

18

21

Reference amplicon position (bp)

- Insertions
- Deletions
- Substitutions