

Mutation position distribution of NHEJ

Sequences: % Total (% NHEJ, no.)

11.1% (inf% , 0)

8.9% (inf% , 0)

6.7% (inf% , 0)

4.4% (inf% , 0)

2.2% (inf% , 0)

0.0% (nan% , 0)

0

3

6

9

12

15

18

21

Reference amplicon position (bp)

- Insertions
- Deletions
- Substitutions