

Mutation position distribution of NHEJ

Sequences: % Total (% NHEJ, no.)

6.2% (inf% , 0)
5.0% (inf% , 0)
3.8% (inf% , 0)
2.5% (inf% , 0)
1.2% (inf% , 0)
0.0% (nan% , 0)

0 3 6 9 12 15 18 21

Reference amplicon position (bp)

- Insertions
- Deletions
- Substitutions