

Mutation position distribution of NHEJ

Sequences: % Total (% NHEJ, no.)

4.5% (inf% , 0)
3.6% (inf% , 0)
2.7% (inf% , 0)
1.8% (inf% , 0)
0.9% (inf% , 0)
0.0% (nan% , 0)

0 3 6 9 12 15 18 21

Reference amplicon position (bp)

- Insertions
- Deletions
- Substitutions