

Mutation position distribution of NHEJ

Sequences: % Total (% NHEJ, no.)

5.3% (inf% , 0)
4.2% (inf% , 0)
3.2% (inf% , 0)
2.1% (inf% , 0)
1.1% (inf% , 0)
0.0% (nan% , 0)

0 3 6 9 12 15 18

Reference amplicon position (bp)

— Insertions
— Deletions
— Substitutions