

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