

Mutation position distribution of NHEJ

Sequences: % Total (% NHEJ, no.)

2.2% (100.0% , 13)

1.7% (80.0% , 10)

1.3% (60.0% , 7)

0.9% (40.0% , 5)

0.4% (20.0% , 2)

0.0% (0.0% , 0)

0 20 40 60 80 100 120

Reference amplicon position (bp)

