

Mutation position distribution of NHEJ

Sequences: % Total (% NHEJ, no.)

33.6% (74.3% , 12168)

26.9% (59.5% , 9734)

20.2% (44.6% , 7300)

13.4% (29.7% , 4867)

6.7% (14.9% , 2433)

0.0% (0.0% , 0)

0

55

110

165

220

275

330

Reference amplicon position (bp)



Insertions



Deletions



Substitutions



Predicted cleavage position



sgRNA

