

# Mutation position distribution of HDR

Sequences: % Total ( % HDR, no. )

23.3% (100.0% , 73017)

18.6% (80.0% , 58413)

14.0% (60.0% , 43810)

9.3% (40.0% , 29206)

4.7% (20.0% , 14603)

0.0% (0.0% , 0)

0 40 80 120 160 200 240

Reference amplicon position (bp)

- Insertions
- Deletions
- Substitutions
- Predicted cleavage position
- sgRNA

