Mutation position distribution of NHEJ ¿ 50.0% (100.0%, 0) % Total ( % NHEJ, 40.0% (80.0%, 0) 30.0% (60.0%, 0) Sequences: 20.0% (40.0%, 0) 10.0% (20.0%, 0) 0.0% (0.0%, 0) 3 6 9 12 15 18 21 Reference amplicon position (bp) **Insertions Deletions** Substitutions