Mutation position distribution of NHEJ 16.7% (100.0%, 1) Sequences: % Total (% NHEJ, 13.3% (80.0%, 1) 10.0% (60.0%, 1) 6.7% (40.0%, 0) 3.3% (20.0%, 0) 0.0% (0.0%, 0) 3 6 9 12 21 15 18 Reference amplicon position (bp) **Insertions Deletions Substitutions**