Mutation position distribution of NHEJ 14.9% (100.0%, 7) Sequences: % Total (% NHEJ, 11.9% (80.0%, 5) 8.9% (60.0%, 4) 6.0% (40.0%, 2) 3.0% (20.0%, 1) 0.0% (0.0%, 0) 3 6 9 12 15 18 21 Reference amplicon position (bp) **Insertions Deletions Substitutions**