Mutation position distribution of NHEJ 29.2% (100.0%, 7) % Total (% NHEJ, 23.3% (80.0%, 5) 17.5% (60.0%, 4) Sequences: 11.7% (40.0%, 2) 5.8% (20.0%, 1) 0.0% (0.0%, 0) 3 6 9 12 15 18 21 Reference amplicon position (bp) **Insertions Deletions** Substitutions