Mutation position distribution of NHEJ e 6.7% (100.0% , 1) Sequences: % Total (% NHEJ, 5.3% (80.0%, 1) 4.0% (60.0%, 1) 2.7% (40.0%, 0) 1.3% (20.0%, 0) 0.0% (0.0%, 0) 3 6 9 12 15 18 Reference amplicon position (bp) **Insertions Deletions** Substitutions