


C A T T A C A T G A G G T G T T C G T T G T A A C T C A T A T A A A C T G A G T - Reference

PE Extension

PE spacer sgRNA

C A T T A C A T G A G G T G T T C G T T G **C** A A C T C A T A T A A A C T G A G T - 0.37% (93 reads)
C A T T A C A T G A G G T G T T C G T T G **C** A **C** C **G** - - - - - A C T **C** **G** G T - 0.22% (54 reads)

bold Substitutions
 Insertions
- Deletions