

C A T T A C C 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 C 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 - 54.20% (13503 reads)
C A T T A C C T G A G G **C** 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 - 0.22% (54 reads)

bold Substitutions

 Insertions

- Deletions