


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

PE nicking sgRNA

A T A T A A A C T G A G T T C C C A T G T T T T G C T T A A T G G T T G A G T T-54.97% (13694 reads)

- bold** Substitutions
-  Insertions
- Deletions
- Predicted cleavage position