

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