

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 - 1.28% (320 reads)

**bold**

Substitutions



Insertions

-

Deletions

-----

Predicted cleavage position