

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**

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Substitutions

Insertions

Deletions

Predicted cleavage position