

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

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Deletions

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Predicted cleavage position