


C G T C G A C A A C G G C T C C G G C A T G T G C A A G G C C G G C T T C G C G - Reference
PE Extension PE nicking sgRNA

C G T C G A C A A C G G C T C C G G C A T G T G C A A G G C C G G C T T C G C G - 40.00% (2 reads)

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