```
CAAAGTACTTCCAGTCCTGCCAGGGGATGGGGTA-Reference
sgRNA
```

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C A A A G T A C T T C T T C C A G T C C T G C C A G G G G A T G G G G T A - 56.61% (1174 reads)****

C A A A G T A C T T C T T C C A G T C C T G C C A G G - - A T G G G G T A - 9.02% (187 reads)

C A A A G T A C T T C T T C C A G T C C T G C C A G G - - - - - T A - 4.00% (83 reads)
CAAAGTACTTCTTCCAGTCCTGCCAGGG-ATGGGGTA-3.23% (67 reads)
CAAAGTACTTCTTCCAGTCCTGCCA-----TGGGGTA-3.13% (65 reads)
CAAAGTACTTCTTCCAGTCCTGCCAGGGGGATGGGGT-1.49% (31 reads
CAAAGTACTTCTTCCAGTCCTGCCAGGG-----TA-1.40% (29 reads)
CAAA-TACTTCTTCCAGTCCTGCCAGG--ATGGGGTA-0.87% (18 reads)
CAAAGTACTTCTTCCAGTCCTGCCAGG - - - TAGGGGGT-0.82% (17 reads)
CAAAGTACTTCTTCCAGTCCTGCCAGG--ATGGAGTA-0.77%
CAAAGTACTTCTTCCAGTCCTGCCAGGGG-----TA-0.68% (14 reads)
```

□ Insertions
□ Deletions
------ Predicted cleavage position