A G T A G C C C C G A C T C T T G T A C G G T C G G C A T C T G A G A C C A G T   Reference

sgRNA ▬▬▬▬▬▬▬▬▬▬▬▬▬▬▬▬▬▬▬▬

A G T A G C C C C G A C T C T T G T A C G G T C G G C A T C T G A G A C C A G T   97.67% (210 reads)
A G **G** A G C C C **A** G A **A G** - - T G T A **A** G G T **A** G G C A T C T G A G A C C A G **G**   0.47% (1 reads)
A G T A G C C C C G A C T - - T G T A C G G T C G G C A T C T G A G A C C A G T   0.47% (1 reads)
A G T A G C C C C G A C T C T T G **C** A C G G T C G G **A** C A T C T G A G A C C A G T   0.47% (1 reads)
A G T A G C C C C G A C T C T T G T **G** C G G T C G G C A T C T G A G A C C A G T   0.47% (1 reads)
A G T A G C C C C **T** A C T C T T G T A C G G T C G G C A T C T G A G A C C A G T   0.47% (1 reads)

bold   Substitutions

☐   Insertions

-    Deletions

- - - - -   Predicted cleavage position