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

sgRNA

C C A G T T T T C T C G C A C T T A A G C T C T G G T T C G G A C A G A A A T A - 87.10% (17603 reads)
C C A G T T T T C T C G C A C T T A A G C T C T G G T T **T** G G A **T** C G A A A **G** A - 1.70% (343 reads)
C C A G T T **G** T C T C G C A C T T A A G C T C T G G T T C G G A C A G A A A T A - 0.38% (76 reads)
C C A G T **G** T T C T C G C A C T T A A G C T C T G G T T C G G A C A G A A A T A - 0.32% (65 reads)
C C A G T T T T C T C **T** C A C T T A A G C T C T G G T T C G G A C A G A A A T A - 0.25% (50 reads)
C C A G T T T T C T C G C A C T T A A G C T C T G G T T C **T** G A C A G A A A T A - 0.25% (50 reads)
A C A G T T T T C T C G C A C T T A A G C T C T G G T T C G G A C A G A A A T A - 0.24% (49 reads)
C C A G T **C** T C T C G C A C T T A A G C T C T G G T T C G G A C A G A A A T A - 0.23% (46 reads)
C C A G T T T T C T C G C A C T T A **G** G C T C T G G T T C G G A C A G A A A T A - 0.22% (44 reads)
C C A G **A** T T T C T C G C A C T T A A G C T C T G G T T C G G A C A G A A A T A - 0.21% (42 reads)

bold   Substitutions

☐   Insertions

-    Deletions

------   Predicted cleavage position