CTAGTTTTCTCGCACTTAAGCTCTGGTTCAGACAGAAACG-Reference

sgRNA

CTAGTTTTCTCGCACTTAAGCTCTGGTTCAGACAGAAACG-83.99% (299 reads)

CTAGTTTTCTCGCACCTAAGCTCTGGTTCAGACAGAAACG-1.12% (4 reads)

CTAGTTTTCTCGCACTTAAGCCTCTGGTTCAGACAGAAAC-0.84% (3 reads)

CTAGTTTTCTCGCACTTAAGCTCTGGCTCAGACAGAAACG-0.84% (3 reads)

CTAGTATTCTCGCACTTAAGCTCTGGTTCAGACAGAAACG-0.56% (2 reads)

CTAGTTTTCTCGCGCTTAAGCTCTGGTTCAGACAGAAACG-0.56% (2 reads)

CTAGTTTTCTCGCACTTAAGCTCTGTTTCAGACAGAAACG-0.56% (2 reads)

CTAGTTTTCTCGCACTTAAGCTCTGGTTCATACAGAAACG-0.56% (2 reads)

CTAGTTTTCTCGCACTTAAGCTCTGGTTCAGACAGAAGCG-0.56% (2 reads)

CTAGTCTTCTCGCACTTAAGCTCTGGTTCAGACAGAAACG-0.56% (2 reads)

CTAGTTTTCTCGCACTTAAGCTCTGGTTCAGACAGAACCG-0.56% (2 reads)

CTAGTTGTCTCGCACTTAAGCTCTGGTTCAGACAGAAACG-0.56% (2 reads)

CTAGTTTTCTCGCACTCAAGCTCTGGTTCAGACAGAAACG-0.28% (1 reads)

CTAGTTTTCTCGCACTTAA-CTCTGGTTCAGACAGAAACG-0.28% (1 reads)

CTTGTTTTCTCGCACTTAAGCTCTGGTTCATACAGAAACG-0.28% (1 reads)

CTTGTTTTCTCGCACTTAAGCTCTGGTTCAGACAGAAACG-0.28% (1 reads)

CTAGTTTTTTCGCACTTAAGCTCTGGTTCAGACAGAAACG-0.28% (1 reads)

CTAGTTTTTTCGCACTTAAGCTATGGTTCAGACAGAAACG-0.28% (1 reads)

CTAGTTTTCGCACTTAAGCTCTGGTTCAGACAGAAACG-0.28% (1 reads)

CTAGTTTTCTCGCACTTTAAGCTCTGGTTCAGACAGAAACG-0.28% (1 reads)

CTAGTTTTCTCGCACTTCAGCTCTGGTTCAGACAGAATACG-0.28% (1 reads)

CTAGTTTTCTCGCACTTCAGCTCTGGTTCAGACAGAAACG-0.28% (1 reads)

CTAGTTTTCTCGCACTTCAGCACTGGTTCAGACAGAAACG-0.28% (1 reads)

CTAGTTTTCTCGCACTTAAGTCTGGTTCAGACAGAAACG-0.28% (1 reads)

CTAGTTTTCTCGCACTTAAGGCTCTGGTTCAGACAGAAACG-0.28% (1 reads)

CTAGTTTTCTCGCACTTAAGCTCTTGTTCAGACAGAAACG-0.28% (1 reads)

CTAGTTTCCTCGCACTTAAGCTCTGGTTCAGACAGAAACG-0.28% (1 reads)

CTAGTTTTCTCGCACTTAAGCTCTGGTTCTGACAGAAACG-0.28% (1 reads)

CTAGTTTTATCGCACTTAAGCTCTGGTTCAGACAGAAACG-0.28% (1 reads)

CTAGTTTTCTCGCACTTAAGCTCTGGTTCAGAATAGAAACG-0.28% (1 reads)

CTAGTTTTCTCGCACTTAAGCTCTGGTTCAGACTGAAACG-0.28% (1 reads)

CTAGTTTTCTCGCACTTAAGCTCTGGTTCAGACAGTAACG-0.28% (1 reads)

CTAGTTTTCACGCACTTAAGCTCTGGTTCAGACAGAAACG-0.28% (1 reads)

CTAGTTTTCTACGCACTTAAGCTCTGGTTCAGACAGAAACG-0.28% (1 reads)

CTAGTTTTCTCGCACTTAAGCTCTGGTTCAGACAGAAACT-0.28% (1 reads)

CTAGTTTTCTCCCACTTAAGCTCTGGTTCAGACAGAAACG-0.28% (1 reads)

CTAGTTTTCTCGCACTTAAGCTCTGGTTCAGACAGAAAAG-0.28% (1 reads)

CTAGTTTTCTCGCACGTAAGCTCTGGTTCAGACAGAAACG-0.28% (1 reads)

CTAGTTTTCTCGCACTTAAGCTCCGGTTCAGACAGAAACG-0.28% (1 reads)

CTAGTTTTCTCGCACTAAAGCTCTGGTTCAGACAGAAACG-0.28% (1 reads)

CTAGTTTTCTCGCACTTAAGCCCTGGTTCAGACAGAAACG-0.28% (1 reads)

CTAGTTTTCTCGCACTTAAGACTCTGGTTCAGGGCAGAAAC-0.28% (1 reads)

GTAGTTTTCTCGCACTTAAGCTCTGGTTCTGACAGAAACG-0.28% (1 reads)

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