CTAGTTTTCTCGCACTTAAGCTCTGGTTCAGACAGAAACG-Reference sgRNA

CTAGTTTTCTCGCACTTAAGCTTCTGGTTCAGACAGAAACG-85.70% (33426 reads) CTAGTTGTCTCGCACTTAAGCTTCTGGTTCAGACAGAAACG-0.43% (166 reads) CTAGTTTTCTCGCACTTAAGCTCTGTTTTCAGACAGAAACG-0.37% (145 reads) ctagttttctctctcacttaagcttctggttcagacagaaacg-0.33% (128 reads) CTAGTGTTCTCGCACTTAAGCTTCTGGTTCAGACAGAAACG-0.29% (115 reads CTATTTTCTCGCACTTAAGCTCTGGTTCAGACAGAAACG-0.29% CTAGTTTTCTCGCACTTAAGCTTCTTGTTCAGACAGAAACG-0.27% CTAGTTTTCTCGCACTTAATCTCTGGTTCAGACAGAAACG-0.26% CTAGTCTTCGCACTTAAGCTCTGGTTCAGACAGAAACG-0.25% (97 reads) TCTCGCACTTAAGCTCTGGTTCAGACAGAAACG-0.23% CTAGTTTTCTCGCACTTAAGCTTCTGGTTCAGACAGAAGCG-0.23% (89 reads) CTAGTTTTCTCGCACTTAAGCTCTGGTTCAGACAGAAACT-0.23% (88 reads) CTAGTTTTCTCGCACTTAAGCTTCTGGCTCAGACAGAAACG-0.22% (87 reads)

