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TAGCGACATTCATAACAGAGACATGGTAACGAGATCTGTT-Reference
sgRNA

TAGCGACATTCATAACAGAGACATGGTAACGAGATCTGTT-68.56% (77995 reads)
TAGCGACATTCATAACAGAGTACATGGTAACGAGATCTGT-3.36% (3817 reads)
TAGCGACATTCATAACAGAGGACATGGTAACGAGATCTGT-3.02% (3441 reads)
TAGCGACATTCATAACAGA--CATGGTAACGAGATCTGTT-1.64% (1868 reads)
TAGCGACATTCATAACA---------TGGTAACGAGATCTGTT-1.42% (1619 reads)
TAGCGACATTCATAACAGAGAACATGGTAACGAGATCTGT-0.93% (1054 reads)
TAGCGACATTCATAACA---CATGGTAACGAGATCTGTT-0.78% (891 reads)
TAGCGACATTCATAACAGAGACATGGTCACGAGATCTGTT-0.76% (864 reads)
TAGCGACAGTCATAACAGAGACATGGTAACGAGATCTGTT-0.49% (561 reads)
TAGCGACATTCATAACAGA-ACATGGTAACGAGATCTGTT-0.47% (535 reads)
TAGCGACATTCATAACAGAG--ATGGTAACGAGATCTGTT-0.43% (493 reads)
TAGCGACATTCATAACAGAGACAGGGTAACGAGATCTGTT-0.43% (489 reads)
TAGCGACATTCATAACAGAGACCTGGTAACGAGATCTGTT-0.41% (472 reads)
TAGCGACATTCATAA-----ACATGGTAACGAGATCTGTT-0.39% (440 reads)
TAGCGACATTCATAAAAGAGACATGGTAACGAGATCTGTT-0.31% (358 reads)
TAGCGACATTCATA-----CATGGTAACGAGATCTGTT-0.30% (346 reads)
TAGCGACATTCATAACA---ACATGGTAACGAGATCTGTT-0.27% (309 reads)
TAGCGACATTCATAACAGAGAGATGGTAACGAGATCTGTT-0.27% (308 reads)
TAGCGACAT-------------GGTAACGAGATCTGTT-0.27% (307 reads)
TAGCGACATTCATAACAGAG----------ATCTGTT-0.27% (303 reads)
TCGCGACATTCATAACAGAGACATGGTAACGAGATCTGTT-0.20% (232 reads)
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bold	Substitutions
□	Insertions
-	Deletions
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