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TAGCGACATTCATAACAGAGACATGGTAACGAGATCTGTT- Reference
sgRNA
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TAGCGACATTCATAACAGAG|ACATGGTAACGAGATCTGTT- 37.57% (17740 reads)
TAGCGACATTCATAACAGA-|-CATGGTAACGAGATCTGTT- 8.43% (3979 reads)
TAGCGACATTCATAACA---|---TGGTAACGAGATCTGTT- 4.87% (2301 reads)
TAGCGACATTCATAACAGAG|-CATGGTAACGAGATCTGTT- 4.32% (2042 reads)
TAGCGACATTCATAACA---|-CATGGTAACGAGATCTGTT- 4.22% (1993 reads)
TAGCGACATTCATA------|-CATGGTAACGAGATCTGTT- 2.07% (976 reads)
TAGCGACATTCATAACAGA-|ACATGGTAACGAGATCTGTT- 1.95% (922 reads)
TAGCGACATTCATAACAGAG|----------ATCTGTT- 1.61% (758 reads)
TAGCGACATTCATAA-----|ACATGGTAACGAGATCTGTT- 1.55% (734 reads)
TAGCGACAT-----------|----GGTAACGAGATCTGTT- 1.44% (678 reads)
TAGCGACATTCATAACAGAG|--ATGGTAACGAGATCTGTT- 0.76% (359 reads)
TAGCGACATTCATAA-----|---TGGTAACGAGATCTGTT- 0.72% (339 reads)
TAGCGACATTCATAACA---|ACATGGTAACGAGATCTGTT- 0.69% (326 reads)
TAGCGACATTCATAACAGAG|-----GTAACGAGATCTGTT- 0.63% (296 reads)
TAGCGACATTCATAAC----|-CATGGTAACGAGATCTGTT- 0.59% (277 reads)
TAGCGACATTCATAACAGA-|----------ACGAGATCTGTT- 0.55% (259 reads)
TAGCGACATTCATAAC----|----------GAGATCTGTT- 0.53% (251 reads)
TAGCGACATTCATAACAGAG|------TAACGAGATCTGTT- 0.50% (236 reads)
TAGCGACATTCATAACAGA-|---TGGTAACGAGATCTGTT- 0.49% (230 reads)
TAGCGACATTCA--------|-CATGGTAACGAGATCTGTT- 0.44% (210 reads)
TAGCGACATTCAT-------|----GGTAACGAGATCTGTT- 0.35% (165 reads)
TAGCGACATTCAT------G|ACATGGTAACGAGATCTGTT- 0.34% (160 reads)
TAGCGACATTCATA-----G|ACATGGTAACGAGATCTGTT- 0.30% (142 reads)
TAGCGACATTCATA------|---TGGTAACGAGATCTGTT- 0.28% (132 reads)
TAGCGACATTCA--------|ACATGGTAACGAGATCTGTT- 0.25% (120 reads)
TAGAGACATTCATAACAGAG|ACATGGTAACGAGATCTGTT- 0.25% (118 reads)
TAGCGACATTCATAACAGAG|---TGGTAACGAGATCTGTT- 0.23% (108 reads)
TAGCGACATTCAT-------|------TAACGAGATCTGTT- 0.22% (105 reads)
TAGCGACATTCATAACAGAG|ACATTGTAACGAGATCTGTT- 0.22% (103 reads)
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bold	Substitutions
□	Insertions
-	Deletions
------	Predicted cleavage position