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TTTTGTGCCAGAGTCCTTCGATAGGGACAAGACAATTGCA- Reference
sgRNA

TTTTGTGCCAGAGTCCTTCG|ATAGGGACAAGACAATTGCA- 73.97% (68107 reads)
TTTTGTGCCAGAGTCCTTCG|GATAGGGACAAGACAATTGC- 2.59% (2388 reads)
TTTTGTGCCAGAGTCCTTC-|ATAGGGACAAGACAATTGCA- 0.82% (755 reads)
TTTTGTGCCAGAGTC-----|ATAGGGACAAGACAATTGCA- 0.64% (590 reads)
TTTTGTGCCAGAGTCCTTCG|-TAGGGACAAGACAATTGCA- 0.53% (492 reads)
TTTTGTGCCAGAGTCCTTCG|-----------AATTGCA- 0.44% (409 reads)
TTTTGTGCCAGAGTCCTT--|ATAGGGACAAGACAATTGCA- 0.40% (365 reads)
TTTTGTGCCAGAG-------|----GGACAAGACAATTGCA- 0.39% (355 reads)
TTTTGTGCCAGAG-------|ATAGGGACAAGACAATTGCA- 0.37% (340 reads)
TTTTGTGCCAGAGTCCTTCG|----GGACAAGACAATTGCA- 0.34% (317 reads)
TTTTGTGCCAGAGTCCTTCG|----------ACAATTGCA- 0.34% (316 reads)
TTTTGTGCCAG---------|----GGACAAGACAATTGCA- 0.31% (283 reads)
TTTTGTGCCAGAGTCCTTCG|--AGGGACAAGACAATTGCA- 0.29% (270 reads)
TTTTGTGCCAGAGTCCTTCG|----------CAATTGCA- 0.29% (268 reads)
TTTTGTGCCAGAGTCCTTCG|-----GACAAGACAATTGCA- 0.27% (247 reads)
TTTTGTGCCAGAGTCCTTCG|AATAGGGACAAGACAATTGC- 0.25% (233 reads)
TTTTGTGCCAGAGTCC----|----AAGACAATTGCA- 0.25% (233 reads)
TTTTGTGCCAGAGTCCTTCG|ATAGGGACAAGGCAATTGCA- 0.25% (232 reads)
TTTTGTG-------------|ATAGGGACAAGACAATTGCA- 0.25% (230 reads)
TTTTGTGCCAGAGTCCTTC-|----AAGACAATTGCA- 0.25% (229 reads)
TTTTGTGCCAGAG-------|----------ACAATTGCA- 0.23% (216 reads)
TTTTGTGCCAGAGTCCTTCG|ATAGGGGCAAGACAATTGCA- 0.23% (208 reads)
TTTTGTGCCAGAGTCCCTCG|ATAGGGACAAGACAATTGCA- 0.22% (204 reads)
TTTTGTGCCA----------|-TAGGGACAAGACAATTGCA- 0.22% (199 reads)
TTTTGTGCCAGGGTCCTTCG|ATAGGGACAAGACAATTGCA- 0.20% (188 reads)
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bold	Substitutions
□	Insertions
-	Deletions
------	Predicted cleavage position