

C A T T T T C G G T T C T C C A C T A A A T T T G G A C A G G C C T T G A T T G - Reference

sgRNA

C	A	T	T	T	T	C	G	G	T	T	C	T	C	C	A	C	T	A	A	A	T	T	T	G	G	A	C	A	G	G	C	C	T	T	G	A	T	T	G	-	75.68% (11679 reads)
C	A	T	T	T	T	C	G	G	T	T	C	T	C	C	A	C	T	A	A	-	T	T	T	G	G	A	C	A	G	G	C	C	T	T	G	A	T	T	G	-	2.22% (342 reads)
C	A	T	T	T	T	C	G	G	T	T	C	T	C	C	A	-	-	-	-	-	T	T	T	G	G	A	C	A	G	G	C	C	T	T	G	A	T	T	G	-	1.78% (274 reads)
C	A	T	T	T	T	C	G	G	T	T	C	T	C	C	A	C	T	A	A	T	A	T	T	T	G	G	A	C	A	G	G	C	C	T	T	G	A	T	T	-	1.39% (215 reads)
C	A	T	T	T	T	C	G	G	T	T	C	T	C	C	A	C	-	-	-	A	T	T	T	G	G	A	C	A	G	G	C	C	T	T	G	A	T	T	G	-	1.10% (170 reads)
C	A	T	T	T	T	C	G	G	T	T	C	T	C	C	A	C	T	A	-	-	T	T	T	G	G	A	C	A	G	G	C	C	T	T	G	A	T	T	G	-	0.81% (125 reads)
C	A	T	T	T	T	C	G	G	T	T	C	T	C	C	A	-	-	-	-	A	T	T	T	G	G	A	C	A	G	G	C	C	T	T	G	A	T	T	G	-	0.80% (124 reads)
C	A	T	T	T	T	C	G	G	T	T	C	T	C	C	A	C	T	A	A	C	T	T	T	G	G	A	C	A	G	G	C	C	T	T	G	A	T	T	G	-	0.51% (79 reads)
C	A	T	T	T	T	C	G	G	T	T	C	-	-	-	-	-	-	-	-	A	T	T	T	G	G	A	C	A	G	G	C	C	T	T	G	A	T	T	G	-	0.49% (76 reads)
C	A	T	T	T	T	C	G	G	T	T	C	T	C	C	A	C	T	A	C	A	T	T	T	G	G	A	C	A	G	G	C	C	T	T	G	A	T	T	G	-	0.41% (63 reads)
C	A	T	T	T	T	C	T	G	T	T	C	T	C	C	A	C	T	A	A	A	T	T	T	G	G	A	C	A	G	G	C	C	T	T	G	A	T	T	G	-	0.38% (59 reads)
C	A	T	T	T	T	C	G	G	G	T	C	T	C	C	A	C	T	A	A	A	T	T	T	G	G	A	C	A	G	G	C	C	T	T	G	A	T	T	G	-	0.36% (56 reads)
C	A	T	T	T	T	C	G	G	T	T	C	T	C	C	A	C	T	-	-	-	-	T	T	G	G	A	C	A	G	G	C	C	T	T	G	A	T	T	G	-	0.35% (54 reads)
C	A	T	T	T	T	C	G	G	T	T	C	T	C	C	A	C	T	A	A	A	A	T	T	T	G	G	A	C	A	G	G	C	C	T	T	G	A	T	T	-	0.30% (47 reads)
C	A	T	T	T	T	C	G	G	T	T	C	T	A	C	A	C	T	A	A	A	T	T	T	G	G	A	C	A	G	G	C	C	T	T	G	A	T	T	G	-	0.27% (42 reads)
C	A	T	T	T	T	C	G	G	T	T	C	T	C	C	A	C	T	A	A	A	T	T	T	G	G	A	C	A	G	G	C	C	G	T	G	A	T	T	G	-	0.24% (37 reads)
C	A	T	T	T	T	C	G	G	T	T	C	T	C	C	A	C	T	A	A	A	T	T	T	G	G	A	C	A	G	G	C	A	T	T	G	A	T	T	G	-	0.23% (36 reads)
C	A	T	T	T	T	C	G	G	C	T	C	T	C	C	A	C	T	A	A	A	T	T	T	G	G	A	C	A	G	G	C	C	T	T	G	A	T	T	G	-	0.21% (32 reads)

bold

Substitutions

Insertions

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Deletions

Predicted cleavage position