

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

sgRNA

A	C	A	T	C	C	A	T	C	G	G	C	G	C	T	T	T	G	G	T	C	G	G	-	93.19% (219 reads)
A	C	A	T	C	C	A	T	C	G	G	C	G	C	T	T	T	G	G	T	C	G	G	-	0.85% (2 reads)
A	C	<b>C</b>	T	C	C	A	T	C	G	G	C	G	C	T	T	T	G	G	T	C	G	G	-	0.85% (2 reads)
-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	0.43% (1 reads)
A	<b>A</b>	C	<b>T</b>	<b>C</b>	T	T	G	<b>C</b>	<b>T</b>	T	C	<b>C</b>	C	T	<b>C</b>	<b>C</b>	T	G	<b>A</b>	C	G	<b>C</b>	-	0.43% (1 reads)
A	C	A	T	C	C	A	<b>C</b>	<b>T</b>	G	<b>C</b>	C	G	C	T	T	T	G	<b>T</b>	T	C	G	G	-	0.43% (1 reads)
A	C	A	T	C	C	A	T	C	<b>A</b>	G	C	G	C	T	T	T	G	G	T	C	T	G	-	0.43% (1 reads)
A	C	A	T	C	C	A	T	C	G	<b>C</b>	<b>A</b>	G	C	T	T	T	G	<b>T</b>	<b>C</b>	C	G	G	-	0.43% (1 reads)
A	C	A	T	C	C	A	T	C	G	<b>C</b>	<b>T</b>	G	C	T	T	T	G	G	T	C	G	G	-	0.43% (1 reads)
A	C	A	T	C	C	A	T	C	G	G	C	G	C	T	T	T	G	G	<b>C</b>	C	C	G	-	0.43% (1 reads)
A	C	A	T	C	C	A	T	C	G	G	C	G	C	T	T	T	G	G	<b>C</b>	C	G	G	-	0.43% (1 reads)
A	C	A	T	C	C	A	T	C	G	G	C	G	C	T	T	T	G	G	<b>G</b>	C	G	G	-	0.43% (1 reads)
A	C	<b>C</b>	T	C	C	A	T	C	G	G	C	G	C	T	<b>C</b>	<b>C</b>	G	G	<b>C</b>	C	G	C	-	0.43% (1 reads)
<b>C</b>	<b>C</b>	A	T	<b>T</b>	C	<b>C</b>	T	<b>T</b>	<b>C</b>	G	C	<b>C</b>	C	T	T	T	<b>T</b>	G	<b>G</b>	C	G	G	-	0.43% (1 reads)
C	T	A	C	A	G	A	G	C	T	G	A	C	C	C	T	G	A	G	T	-	-	-	-	0.43% (1 reads)

**bold** Substitutions

Insertions

- Deletions

----- Predicted cleavage position