


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

G	A	G	T	A	G	G	T	G	G	A	C	T	A	T	G	C	A	G	T	C	A	A	T	G	G	A	C	G	A	A	C	T	G	C	C	G	C	T	A	-49.26% (144806 reads)
A	G	T	<b>T</b>	A	G	G	T	G	G	A	C	T	<b>C</b>	T	G	C	A	G	T	C	A	A	T	G	G	A	C	G	A	A	C	T	G	C	C	G	C	T	A	-34.73% (102098 reads)
A	G	<b>A</b>	<b>T</b>	A	G	G	T	G	G	A	C	T	A	T	G	C	A	G	T	C	A	A	T	G	G	A	C	G	A	A	C	T	G	C	C	G	C	T	A	-6.34% (18628 reads)
A	G	<b>A</b>	<b>T</b>	A	G	G	T	G	G	A	C	T	A	T	G	C	A	G	T	C	A	A	T	G	G	A	C	G	A	A	C	T	<b>T</b>	C	C	G	C	T	A	-0.83% (2446 reads)
A	G	T	<b>T</b>	A	G	G	T	G	G	A	C	T	A	T	G	C	A	G	T	C	A	A	T	G	G	A	C	G	A	A	C	T	G	C	C	G	C	T	A	-0.47% (1386 reads)
G	A	G	T	A	G	G	T	G	G	A	C	T	<b>C</b>	T	G	C	A	G	T	C	A	A	T	G	G	A	C	G	A	A	C	T	G	C	C	G	C	T	A	-0.34% (1001 reads)
A	G	<b>A</b>	<b>G</b>	A	G	G	T	G	G	A	C	T	<b>C</b>	T	G	C	A	G	T	C	A	A	T	G	G	A	C	G	A	A	C	T	G	C	C	G	C	T	A	-0.32% (950 reads)

**bold** Substitutions

 Insertions

- Deletions

----- Predicted cleavage position