

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

sgRNA

T	G	G	C	C	C	C	G	C	A	C	A	A	C	T	C	G	T	T	G	A	G	C	A	A	T	T	C	T	T	A	A	A	A	A	G	A	T	G	G	-36.00% (36 reads)	
G	G	C	C	C	C	G	C	A	C	A	A	C	T	C	G	T	G	T	G	A	G	C	A	A	T	T	C	T	T	A	A	A	A	A	G	A	T	G	G	-2.00% (2 reads)	
T	G	G	C	C	C	C	G	C	A	A	A	A	C	T	C	G	T	T	G	A	G	C	A	A	T	T	C	T	T	A	A	A	A	A	G	A	T	G	G	-1.00% (1 reads)	
T	G	G	C	C	C	C	G	C	A	C	A	A	C	T	C	G	T	-	-	-	-	-	A	A	T	T	C	T	T	A	A	A	A	A	G	A	T	G	G	-1.00% (1 reads)	
T	G	G	C	C	C	C	G	C	A	C	A	A	C	T	C	G	T	-	G	A	G	C	A	A	T	T	C	T	T	A	A	A	A	A	G	A	T	G	G	-1.00% (1 reads)	
T	G	G	C	C	C	C	G	C	A	C	A	A	C	T	C	G	T	T	-	-	-	-	-	-	-	-	-	-	T	A	A	A	A	A	G	A	T	G	G	-1.00% (1 reads)	
T	G	G	C	C	C	C	G	C	A	C	A	A	C	T	C	G	T	T	G	-	-	-	-	A	T	T	C	T	T	A	A	A	A	A	G	A	T	G	G	-1.00% (1 reads)	
T	G	G	C	C	C	C	G	C	A	C	A	A	C	T	C	G	T	T	G	-	-	-	A	A	T	T	C	T	T	A	A	A	A	A	G	A	T	G	G	-1.00% (1 reads)	
T	G	G	C	C	C	C	G	C	A	C	A	A	C	T	C	G	T	T	G	A	-	-	-	-	-	-	C	T	T	A	A	A	A	A	G	A	T	G	G	-1.00% (1 reads)	
T	G	G	C	C	C	C	G	C	A	C	A	A	C	T	C	G	T	T	G	A	-	C	A	A	T	T	C	T	T	A	A	A	A	A	G	A	T	G	G	-1.00% (1 reads)	
T	G	G	C	C	C	C	G	C	A	C	A	A	C	T	C	G	T	T	G	A	G	-	-	-	-	-	-	-	-	-	A	A	A	A	A	G	A	T	G	G	-1.00% (1 reads)
T	G	G	C	C	C	C	G	C	A	C	A	A	C	T	C	G	T	T	G	A	G	-	-	-	-	T	C	T	T	A	A	A	A	A	G	A	T	G	G	-1.00% (1 reads)	
T	G	G	C	C	C	C	G	C	A	C	A	A	C	T	C	G	T	T	G	A	G	-	A	A	T	T	C	T	T	A	A	A	A	A	G	A	T	G	G	-1.00% (1 reads)	

bold Substitutions

Insertions

- Deletions

----- Predicted cleavage position