

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

sgRNA

C	C	A	G	T	T	T	T	C	T	C	G	C	A	C	T	T	A	A	G	C	T	C	T	G	G	T	T	C	G	G	A	C	A	G	A	A	A	T	A	-	83.23% (2233 reads)	
C	C	A	G	T	T	T	T	C	T	C	G	C	A	C	T	T	A	A	G	A	C	T	C	T	G	G	T	T	C	G	G	A	C	A	G	A	A	A	T	-	1.53% (41 reads)	
C	C	A	G	T	T	T	T	C	T	C	G	C	A	C	T	T	A	A	G	C	T	C	T	G	G	T	T	T	G	G	A	T	C	G	A	A	A	G	A	-	0.89% (24 reads)	
C	C	A	G	T	T	T	T	C	T	C	G	C	A	C	T	T	A	A	G	T	C	T	C	T	G	G	T	T	C	G	G	A	C	A	G	A	A	A	T	-	0.75% (20 reads)	
C	C	A	G	T	T	T	T	C	T	C	G	C	A	C	T	T	A	A	-	C	T	C	T	G	G	T	T	C	G	G	A	C	A	G	A	A	A	T	A	-	0.56% (15 reads)	
C	C	A	G	T	T	G	T	C	T	C	G	C	A	C	T	T	A	A	G	C	T	C	T	G	G	T	T	C	G	G	A	C	A	G	A	A	A	T	A	-	0.45% (12 reads)	
C	C	A	G	T	T	T	T	C	T	C	T	C	A	C	T	T	A	A	G	C	T	C	T	G	G	T	T	C	G	G	A	C	A	G	A	A	A	T	A	-	0.37% (10 reads)	
C	C	A	G	T	T	T	T	C	T	C	G	C	A	C	T	T	A	A	G	C	T	C	T	G	G	T	T	C	G	G	A	C	A	G	A	G	A	T	A	-	0.37% (10 reads)	
C	C	A	G	T	T	T	T	C	T	C	G	C	A	C	T	T	A	A	G	C	T	C	T	G	G	T	T	C	G	G	A	C	A	G	G	A	A	T	A	-	0.37% (10 reads)	
C	C	A	G	T	T	T	T	C	T	C	G	C	A	C	T	T	A	A	G	C	C	T	C	T	G	G	T	T	C	G	G	A	C	A	G	A	A	A	T	-	0.34% (9 reads)	
C	C	A	G	T	T	T	T	C	T	C	G	C	A	C	T	C	A	A	G	C	T	C	T	G	G	T	T	C	G	G	A	C	A	G	A	A	A	T	A	-	0.30% (8 reads)	
C	C	A	G	T	T	T	T	C	T	C	G	C	A	C	T	T	C	A	G	C	T	C	T	G	G	T	T	C	G	G	A	C	A	G	A	A	A	T	A	-	0.26% (7 reads)	
C	C	A	G	T	T	T	T	C	T	C	G	C	G	C	T	T	A	A	G	C	T	C	T	G	G	T	T	C	G	G	A	C	A	G	A	A	A	T	A	-	0.26% (7 reads)	
C	C	A	G	T	C	T	T	C	T	C	G	C	A	C	T	T	A	A	G	C	T	C	T	G	G	T	T	C	G	G	A	C	A	G	A	A	A	T	A	-	0.26% (7 reads)	
C	C	A	G	T	T	T	T	C	T	C	A	C	A	C	T	T	A	A	G	C	T	C	T	G	G	T	T	C	G	G	A	C	A	G	A	A	A	T	A	-	0.22% (6 reads)	
C	C	A	G	T	T	T	T	C	T	C	G	C	A	C	T	-	-	-	-	-	-	-	C	T	G	G	T	T	C	G	G	A	C	A	G	A	A	A	T	A	-	0.22% (6 reads)
C	C	A	G	T	G	T	T	C	T	C	G	C	A	C	T	T	A	A	G	C	T	C	T	G	G	T	T	C	G	G	A	C	A	G	A	A	A	T	A	-	0.22% (6 reads)	
C	C	A	G	T	T	C	T	C	G	C	A	C	A	C	T	T	A	A	G	C	T	C	T	G	G	T	T	C	G	G	A	C	A	G	A	A	A	T	A	-	0.22% (6 reads)	
C	C	A	G	T	T	T	T	C	T	C	G	C	A	C	T	T	A	A	G	C	T	C	T	G	G	T	T	C	G	G	G	C	A	G	A	A	A	T	A	-	0.22% (6 reads)	
C	C	A	G	T	T	T	T	C	T	C	G	C	A	C	T	T	A	A	T	C	T	C	T	G	G	T	T	C	G	G	A	C	A	G	A	A	A	T	A	-	0.22% (6 reads)	

bold Substitutions

Insertions

- Deletions

----- Predicted cleavage position