

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

sgRNA

C	A	T	G	G	A	A	T	C	C	C	T	T	T	C	T	G	C	T	T	T	T	C	C	G	A	G	-	31.00% (31 reads)														
C	A	T	G	G	A	A	T	C	C	C	T	T	T	C	T	G	C	A	-	-	-	T	G	G	A	T	C	G	C	-	2.00% (2 reads)											
A	T	C	C	C	T	T	C	T	G	C	A	A	A	G	C	A	C	C	T	G	G	A	T	C	G	C	T	T	T	T	C	C	G	A	G	-	1.00% (1 reads)					
A	T	G	G	A	A	T	C	C	C	T	T	T	C	T	G	C	A	A	G	C	A	C	C	T	G	G	A	T	C	G	C	T	T	T	T	C	C	G	A	G	-	1.00% (1 reads)
C	A	T	G	G	A	A	T	C	C	C	T	T	T	C	T	G	C	A	-	-	-	-	-	-	-	-	-	-	T	C	C	G	A	G	-	1.00% (1 reads)						
C	A	T	G	G	A	A	T	C	C	C	T	T	T	C	T	G	C	A	-	-	-	-	T	G	G	A	T	C	G	C	T	T	T	T	C	C	G	A	G	-	1.00% (1 reads)	
C	A	T	G	G	A	A	T	C	C	C	T	T	T	C	T	G	C	A	-	C	A	C	C	T	G	G	A	T	C	G	C	T	T	T	T	C	C	G	A	G	-	1.00% (1 reads)
C	A	T	G	G	A	A	T	C	C	C	T	T	T	C	T	G	C	A	G	-	-	-	-	-	-	-	-	-	A	G	-	1.00% (1 reads)										
C	A	T	G	G	A	A	T	C	C	C	T	T	T	C	T	G	C	A	G	-	-	-	G	G	A	T	C	G	C	T	T	T	T	C	C	G	A	G	-	1.00% (1 reads)		
C	A	T	G	G	A	A	T	C	C	C	T	T	T	C	T	G	C	A	G	C	-	-	-	G	C	T	T	T	T	C	C	G	A	G	-	1.00% (1 reads)						
C	A	T	G	G	A	A	T	C	C	C	T	T	T	C	T	G	C	A	G	C	-	C	T	G	G	A	T	C	G	C	T	T	T	T	C	C	G	A	G	-	1.00% (1 reads)	
C	A	T	G	G	A	A	T	C	C	C	T	T	T	C	T	G	C	A	G	C	A	-	-	-	-	-	T	T	T	C	C	G	A	G	-	1.00% (1 reads)						
C	A	T	G	G	A	A	T	C	C	C	T	T	T	C	T	G	C	A	G	C	A	-	-	-	-	-	G	C	T	T	T	T	C	C	G	A	G	-	1.00% (1 reads)			
C	A	T	G	G	A	A	T	C	C	C	T	T	T	C	T	G	C	A	G	C	A	-	-	-	-	-	G	C	T	T	T	T	C	C	G	A	G	-	1.00% (1 reads)			
C	A	T	G	G	A	A	T	C	C	C	T	T	T	C	T	G	C	A	G	C	A	-	-	-	-	-	T	T	T	C	C	G	A	G	-	1.00% (1 reads)						
C	A	T	G	G	A	A	T	C	C	C	T	T	T	C	T	G	C	A	G	C	A	-	-	-	-	-	T	T	T	C	C	G	A	G	-	1.00% (1 reads)						
C	A	T	G	G	A	A	T	C	C	C	T	T	T	C	T	G	C	A	G	C	A	-	-	-	-	-	C	G	A	G	-	1.00% (1 reads)										
<b>G</b>	A	T	G	G	A	A	T	C	C	C	T	T	T	C	T	G	C	A	G	C	A	-	-	-	-	-	T	T	T	C	C	G	A	G	-	1.00% (1 reads)						

**bold** Substitutions

  Insertions

- Deletions

----- Predicted cleavage position