

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

PE Extension
PE spacer sgRNA

C	A	T	G	G	A	A	T	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	-	81.00% (81 reads)	
C	A	T	G	G	A	A	T	C	C	C	T	T	C	T	G	C	A	G	C	A	T	C	T	G	G	A	T	C	G	C	T	T	T	T	C	C	G	A	G	-	15.00% (15 reads)	
C	A	T	G	G	A	A	T	C	C	C	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	C	T	G	C	A	G	C	A	C	C	T	G	G	A	T	C	G	C	T	T	T	T	A	C	C	G	A	G	-	1.00% (1 reads)
C	A	T	G	G	A	A	T	C	C	C	T	T	C	T	G	C	T	G	C	C	C	G	A	C	C	T	G	G	A	T	C	G	C	T	T	T	T	C	C	-	1.00% (1 reads)	
C	A	T	T	G	A	A	T	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	-	1.00% (1 reads)	

bold

Substitutions



Insertions

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Deletions

Predicted cleavage position