


C G C A C C T 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 - Reference
 sgRNA 

C	G	C	A	C	C	T	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	- 91.68% (21445 reads)
C	G	C	A	C	C	T	C	A	T	G	G	A	A	T	T	T	T	T	T	C	T	G	C	A	G	C	A	C	C	T	G	G	A	T	C	G	C	T	T	- 2.81% (658 reads)
C	G	C	A	C	C	T	C	A	T	G	G	A	A	T	T	C	C	T	T	C	T	G	C	A	G	C	A	C	C	T	G	G	A	T	C	G	C	T	T	- 0.55% (128 reads)
C	G	C	A	C	C	T	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	T	A	T	C	G	C	T	T	- 0.29% (68 reads)
C	G	C	A	C	C	T	C	A	T	G	G	A	A	T	T	T	T	T	T	T	T	G	C	A	G	C	A	C	C	T	G	G	A	T	C	G	C	T	T	- 0.27% (63 reads)

bold Substitutions
 Insertions
 - Deletions