

G C A G T C A T C T T A G T C A T T A C C T G A G G T G T T C G T T G T A A C T

PE Extension

PE spacer sgRNA

G	C	A	G	T	C	A	T	C	T	T	A	G	T	C	A	T	T	A	C	C	T	G	A	G	G	T	G	T	T	C	G	T	T	G	T	A	A	C	T	-54.34% (13537 reads)
G	C	A	G	T	C	A	T	C	T	T	A	G	T	C	A	T	T	A	C	C	T	G	A	G	G	T	G	T	T	C	G	T	T	G	T	A	A	C	T	-0.29% (72 reads)
G	C	A	G	T	C	A	T	C	T	T	A	G	T	C	A	T	T	A	C	C	T	G	A	G	G	C	G	T	T	C	G	T	T	G	T	A	A	C	T	-0.22% (56 reads)

bold Substitutions

Insertions

- Deletions

----- Predicted cleavage position