


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

T	A	G	T	T	G	G	C	C	A	A	G	T	C	A	T	T	C	A	A	T	T	T	T	A	A	A	C	G	T	A	C	A	A	A	G	T	T	C	C	- 61.69% (49073 reads)
T	A	G	T	T	G	G	C	C	A	A	G	T	C	A	T	T	C	A	A	T	T	T	T	G	G	A	C	G	T	A	C	A	A	A	G	T	T	C	C	- 30.09% (23932 reads)
T	A	G	T	T	T	G	C	C	A	A	G	T	C	A	T	T	C	A	A	T	T	T	T	A	A	A	C	G	T	A	C	A	A	A	G	T	T	C	C	- 1.97% (1564 reads)
T	A	G	T	T	G	G	C	C	A	A	G	T	C	A	T	T	C	A	A	T	T	T	T	G	A	A	C	G	T	A	C	A	A	A	G	T	T	C	C	- 1.76% (1403 reads)
T	A	G	T	T	T	G	C	C	A	A	G	T	C	A	T	T	C	A	A	T	T	T	T	G	G	A	C	G	T	A	C	A	A	A	G	T	T	C	C	- 0.94% (746 reads)
T	A	G	T	T	G	G	C	C	A	A	G	T	C	A	T	T	C	A	C	T	T	T	T	A	A	A	C	G	T	A	C	A	A	A	G	T	T	C	C	- 0.48% (383 reads)

bold Substitutions

 Insertions

- Deletions