

G	A	G	G	C	A	A	A	G	A	G	A	C	C	G	G	G	C	C	G	C	G	G	A	G	G	C	C	A	A	G	G	A	A	A	G	G	T	A	C	Reference
sgRNA																																								
G	A	G	G	C	A	A	A	G	A	G	A	C	C	G	G	G	C	C	G	C	G	G	A	G	G	C	C	A	A	G	G	A	A	A	G	G	T	A	C	94.16% (129 reads)
C	A	C	C	C	A	A	A	T	A	T	A	C	A	C	T	C	C	C	G	C	C	T	A	C	C	C	A	A	C	G	A	A	A	T	T	C	C	C	0.73% (1 reads)	
C	A	G	T	C	A	-	-	-	-	-	-	-	-	-	-	-	-	-	-	C	T	G	T	G	G	C	A	A	A	T	C	G	T	A	G	G	C	C	G	0.73% (1 reads)
G	A	G	G	C	A	A	A	G	A	G	A	C	C	G	G	G	C	C	G	C	G	G	A	G	G	C	C	A	A	G	C	A	A	A	G	G	T	A	C	0.73% (1 reads)
G	A	G	G	C	G	A	A	G	A	G	A	C	C	G	-	-	-	-	-	C	T	G	A	G	G	C	C	A	A	G	G	A	A	A	G	G	T	A	C	0.73% (1 reads)
G	A	G	G	C	G	A	A	G	A	G	A	C	C	G	G	G	C	C	G	C	T	G	A	G	G	C	C	A	A	G	G	A	A	A	G	G	T	A	C	0.73% (1 reads)
G	C	T	T	G	A	A	A	A	A	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	A	A	G	A	A	A	A	G	G	T	A	C	0.73% (1 reads)
G	G	T	C	T	A	C	-	-	-	A	C	T	C	T	T	T	C	C	C	T	G	C	A	C	G	A	C	G	C	T	C	A	T	C	C	G	T	A	C	0.73% (1 reads)
G	T	G	G	C	A	A	T	G	A	G	A	C	C	G	C	G	A	C	C	A	G	C	A	G	G	A	T	C	T	A	C	A	C	T	C	T	T	A	C	0.73% (1 reads)

- bold** Substitutions
- Insertions
- Deletions
- Predicted cleavage position