


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

G	G	T	G	C	G	G	A	G	C	C	C	C	T	T	C	G	A	G	C	A	G	C	C	G	C	A	G	T	A	C	T	A	C	C	T	G	G	C	G	-	37.42% (9215 reads)
G	G	T	G	C	G	G	A	G	C	C	C	C	T	T	C	G	A	G	C	-	-	-	C	G	C	A	G	T	A	C	T	A	C	C	T	G	G	C	G	-	5.13% (1264 reads)
G	G	T	G	C	G	G	A	G	C	C	C	C	T	T	C	G	A	G	C	-	-	-	-	-	-	A	G	T	A	C	T	A	C	C	T	G	G	C	G	-	1.11% (274 reads)
G	G	T	G	C	G	G	A	G	C	C	C	C	T	T	C	G	A	G	C	C	A	G	C	C	G	C	A	G	T	A	C	T	A	C	C	T	G	G	C	-	1.02% (250 reads)
G	G	T	G	C	G	G	A	G	C	C	C	C	T	T	C	G	A	G	C	A	A	G	C	C	G	C	A	G	T	A	C	T	A	C	C	T	G	G	C	-	0.54% (134 reads)
G	G	T	G	C	G	G	A	G	C	C	C	C	T	T	C	G	A	G	C	T	A	G	C	C	G	C	A	G	T	A	C	T	A	C	C	T	G	G	C	-	0.26% (64 reads)
G	G	T	G	C	G	G	A	G	C	C	C	C	T	T	C	G	A	G	C	-	-	C	C	G	C	A	G	T	A	C	T	A	C	C	T	G	G	C	G	-	0.26% (63 reads)
G	G	T	G	C	G	G	A	G	C	C	C	C	T	T	C	G	A	G	C	C	T	C	A	A	T	G	C	G	C	G	A	A	T	A	C	C	A	C	C	-	0.21% (51 reads)

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