

C	C	A	G	T	T	T	T	C	T	C	G	C	A	C	T	T	A	A	G	C	T	C	T	G	G	T	T	T	G	G	A	T	C	G	A	A	A	G	A	-	Reference
sgRNA																																									
C	C	A	G	T	T	T	T	C	T	C	G	C	A	C	T	T	A	A	G	C	T	C	T	G	G	T	T	T	G	G	A	T	C	G	A	A	A	G	A	-	76.62% (10465 reads)
C	C	A	G	T	T	T	T	C	T	C	G	C	A	C	T	T	A	A	G	A	T	C	T	G	G	T	T	T	G	G	A	T	C	G	A	A	A	G	A	-	1.72% (235 reads)
C	C	A	G	T	T	T	T	C	T	C	G	C	A	C	T	T	A	A	G	C	T	C	T	G	G	T	T	C	G	G	A	C	A	G	A	A	A	T	A	-	1.00% (137 reads)
C	C	A	G	T	T	T	T	C	T	C	G	C	T	C	T	T	A	A	G	C	T	C	T	G	G	T	T	T	G	G	A	T	C	G	A	A	A	G	A	-	0.92% (126 reads)
C	A	A	G	T	T	T	T	C	T	C	G	C	A	C	T	T	A	A	G	C	T	C	T	G	G	T	T	T	G	G	A	T	C	G	A	A	A	G	A	-	0.81% (111 reads)
C	C	A	G	T	G	T	T	C	T	C	G	C	A	C	T	T	A	A	G	C	T	C	T	G	G	T	T	T	G	G	A	T	C	G	A	A	A	G	A	-	0.76% (104 reads)
C	C	A	G	T	T	T	T	C	T	C	G	C	A	C	T	T	A	A	G	C	T	C	T	G	G	T	T	T	G	G	A	T	C	G	A	C	A	G	A	-	0.66% (90 reads)
C	C	A	G	A	T	T	T	C	T	C	G	C	A	C	T	T	A	A	G	C	T	C	T	G	G	T	T	T	G	G	A	T	C	G	A	A	A	G	A	-	0.64% (87 reads)
C	C	A	G	T	T	T	T	C	T	C	T	C	A	C	T	T	A	A	G	C	T	C	T	G	G	T	T	T	G	G	A	T	C	G	A	A	A	G	A	-	0.55% (75 reads)
C	C	A	G	T	T	T	T	C	T	C	G	C	A	C	T	T	A	A	G	C	T	C	T	G	G	T	G	T	G	G	A	T	C	G	A	A	A	G	A	-	0.40% (55 reads)
C	C	A	G	T	T	T	T	C	T	C	G	C	A	C	T	T	A	A	G	C	T	C	T	T	G	T	T	T	G	G	A	T	C	G	A	A	A	G	A	-	0.37% (51 reads)
C	C	A	G	T	A	T	T	C	T	C	G	C	A	C	T	T	A	A	G	C	T	C	T	G	G	T	T	T	G	G	A	T	C	G	A	A	A	G	A	-	0.34% (46 reads)
C	C	A	G	T	T	T	T	C	T	C	G	C	A	C	T	T	A	A	G	C	T	C	T	G	G	T	T	T	T	G	A	T	C	G	A	A	A	G	A	-	0.26% (36 reads)
C	C	A	G	T	T	T	T	C	T	C	G	C	A	C	T	A	A	G	C	T	C	T	G	G	T	T	T	G	G	A	T	C	G	A	A	A	G	A	-	0.25% (34 reads)	
C	C	A	G	T	T	T	T	C	T	C	G	A	A	C	T	T	A	A	G	C	T	C	T	G	G	T	T	T	G	G	A	T	C	G	A	A	A	G	A	-	0.23% (31 reads)
C	C	A	G	T	T	A	T	C	T	C	G	C	A	C	T	T	A	A	G	C	T	C	T	G	G	T	T	T	G	G	A	T	C	G	A	A	A	G	A	-	0.21% (29 reads)

bold Substitutions

Insertions

- Deletions

----- Predicted cleavage position