

Result	Query tag	Search result	Query type	Mismatch	Ends with RG	Chr Position	Strand	Cut site	Score
ATGCAAGGCAGCGACATCCGAGG -- hit	Query:	hit:	No indel	0	Yes	Chr9:73158762-73158784	-	73158768	0
ATGCAAGGCAGCGACATCCGNGG -- query	ATGCAAGGCAGCGACATCCGNGG	ATGCAAGGCAGCGACATCCGAGG							
TGCAAGGCAGCGACATCCGAGG -- hit	Query:	hit:	Del 20	0	Yes	Chr9:73158762-73158783	-	73158768	0.63
TGCAAGGCAGCGACATCCGNGG -- query	TGCAAGGCAGCGACATCCGNGG	TGCAAGGCAGCGACATCCGAGG							
TGCAAGGCAGCGACATCCGAGG -- hit	Query:	hit:	Del 19	1	Yes	Chr9:73158762-73158783	-	73158768	0.76
AGCAAGGCAGCGACATCCGNGG -- query	AGCAAGGCAGCGACATCCGNGG	TGCAAGGCAGCGACATCCGAGG							
ATGCAAGGCAGCGACATCCGAG -- hit	Query:	hit:	Del 1	1	Yes	Chr9:73158763-73158784	-	73158769	26.51
ATGCAAGGCAGCGACATCCNGG -- query	ATGCAAGGCAGCGACATCCNGG	ATGCAAGGCAGCGACATCCGAG							
ATGCAAGGCAGCGACATCCGAG -- hit	Query:	hit:	Del PAM 3	1	Yes	Chr9:73158763-73158784	-	73158769	40.51
ATGCAAGGCAGCGACATCCGGG -- query	ATGCAAGGCAGCGACATCCGGG	ATGCAAGGCAGCGACATCCGAG	Del PAM 1, or Del						
ATGCAAGGCAGCGACATCCGAG -- hit	Query:	hit:	PAM 2	0	Yes	Chr9:73158763-73158784	-	73158769	20.51
ATGCAAGGCAGCGACATCCGNG -- query	ATGCAAGGCAGCGACATCCGNG	ATGCAAGGCAGCGACATCCGAG							
TATGCAAGGCAGCGACATCCGAGG -- hit	Query:	hit:	Ins 19	1	Yes	Chr9:73158762-73158785	-	73158768	0.95
ANTGCAAGGCAGCGACATCCGNGG -- query	ANTGCAAGGCAGCGACATCCGNGG	TATGCAAGGCAGCGACATCCGAGG	Ins PAM 2, or Ins						
ATGCAAGGCAGCGACATCCGAGG -- hit	Query:	hit:	PAM 3	1	No	Chr9:73158761-73158784	-	73158767	40.7
ATGCAAGGCAGCGACATCCGNGG -- query	ATGCAAGGCAGCGACATCCGNGG	ATGCAAGGCAGCGACATCCGAGGA							
ATGCAAGGCAGCGACATCCGAGG -- hit	Query:	hit:	Ins PAM 1	1	No	Chr9:73158761-73158784	-	73158767	40.7
ATGCAAGGCAGCGACATCCGNGNG -- query	ATGCAAGGCAGCGACATCCGNGNG	ATGCAAGGCAGCGACATCCGAGGA							