[Cat. No.]

ATC-0100, ATC-0101

Introduction

CjCas9 is derived from Campylobacter jejuni (C. jejuni) and is the smallest Cas9 orthologue (2.9 kb), which improves the disadvantages of large-sized SpCas9 (4 kb), derived from Streptococcus pyogenes (S. pyogenes).

The CiCas9 gene can be packaged in an adeno-associated virus (AAV) vector together with a reporter gene and is highly specific in vitro and in vivo, enabling precise genome editing applicable for treatment

DNA-directed RNA-Guided Endonucleases (dRGENs) are efficient, affordable, and convenient tools for genome editing experiments. AccuTool™ dRGEN-CjCas9 is a custom-designed sgRNA that can be used with CjCas9, which is provided as a CjCas9-sgRNA expression plasmid. The CjCas9 expression plasmid uses the CMV/T7 promoter for strong transient expression of CjCas9. Plasmids can be delivered to your cell of interest by any standard methods like lipofection, nanoparticle, or electroporation to achieve highly efficient delivery.

Applications

- Genome editina
- Drug discovery: CRISPR library screening, target validation
- Bioprocessing: Cell line engineering
- Agriculture: Plant breeding

Components

Components	Amount	
Lyophilized dRGEN-CjCas9 sgRNA	2 µg or 50 µg	
Plasmid		

^{*} Note: For research use only. Not for use in diagnostic or therapeutic

Specifications

Expression vector for CiCas9 guide RNAs (CiCas9 dRGEN-U6sgRNA) are ampicillin-resistance and stable in general E. coli strains such as DH5α or XL1.

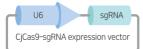


Figure 1. CjCas9 dRGEN-U6-sgRNA

Storage

- AccuTool™ dRGEN-CjCas9 is lyophilized and delivered at ambient temperature.
- Store at -20°C after adding distilled water (D.W.) or TE buffer. Do not store in a frost-free freezer.

sgRNA Information

Name	Target Sequence

Online Resources





Visit our product page for additional information and protocols.

Ordering Information

Description	Cat. No.
AccuTool™ dRGEN-CjCas9 (sgRNA plasmid) - 2 μg	ATC-0100
AccuTool™ dRGEN-CjCas9 (sgRNA plasmid) - 50 μg	ATC-0101

Notice

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Explanation of Symbols







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BQ-042-101-04

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Experimental Procedures

Steps		Procedure Details				
Ge	Gene knock-out cell establishment					
1	Cell transfection	Transfer CRISPR plasmids (sgRNA and Cas9) to target cells. Any DNA delivery method (lipofection, electroporation, nucleofection, or microinjection) optimal for your cell line can be used to deliver CRISPR plasmids. The recommended ratio of sgRNA expression plasmids and Cas9 gene expression plasmids is 1:1-5:1.				
2	Isolation and expanding monoclonal cell colonies	2. 2-3 days after CRISPR plasmids treatment, plate appropriate cell density to isolate monoclonal cell colonies. • The colony formation efficiency could vary among cell lines. Thus, an optimal density of the cell population needs to be determined empirically. • Dish method: plate 50, 250, 1,000, and 5,000 cells/100 mm dish (2 plates/cell population) • Limiting dilution method: plate 0.4 cell/well of 96-well plate (2-3 plates) (It is recommended to proceed with the test for each cell, which means screening after colony seeding.) • Save some of the cell population treated with CRISPR plasmids, and confirm the efficient mutation induction by CRISPR plasmids in your target cell by a T7E1 assay. 3. Isolate and expand monoclonal cell colonies after 10-20 days of plating (50-100 colonies are recommended).				
	Identification of the knock-out cell clones					
3	Genotyping	4. Prepare genomic DNA from each clone between a 48-well plate and a 12-well plate (the plate wells may change depending on the purpose of the experiment). 5. Identify the knock-out cell clones by genotyping. (In/del analysis by NGS or/and T7E1 assay screening.) • T7E1 assay screening: Screening of isolated colonies by T7E1 assay will identify cell clones with a mutation at the target site but will not be able to discern the heterozygous and homozygous knockout cell. Refer to the Mutation Detection Kit Manual for additional information regarding the T7E1 assay screening.	Most mutation induced by CRISPR plasmids at the target site is small deletions and insertions (-20 to +10 bp). When these mutations are causing the frameshift, it will function as a knockout mutation. Cell lines usually			
	Immunoblot analysis	4. Prepare protein from each clone between a 48-well plate and a 12-well plate (the plate wells may change depending on the purpose of the experiment). 5. Identify the knock-out cell clones by immunoblot analysis.	contain more than 2 alleles (polyploidy). The complete knockout cell lines will have frameshift mutation on all alleles of the target gene.			