

# IMPROVED CAS<sub>12A</sub> PROTEINS FOR ACCURATE AND EFFICIENT GENOME EDITING

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## PATENT STATUS

Country	Type	Number	Dated	Case
United States Of America	Published Application	<a href="#">20220315914</a>	10/06/2022	2019-162

## BRIEF DESCRIPTION

Mutated versions of Cas12a that remove its non-specific ssDNA cleavage activity without affecting site-specific double-stranded DNA cutting activity. These mutant proteins, in which a short amino acid sequence is deleted or changed, provide improved genome editing tools that will avoid potential off-target editing due to random ssDNA nicking.

## SUGGESTED USES

Genome editing in animals, plants, and human cells.

## ADVANTAGES

Accurate and efficient genome editing.

Background: Cas12a (formerly called Cpf1) is a type V CRISPR-Cas enzyme derived from bacteria that is used for RNA-guided genome editing in animal, plant and human cells. However, Cas12a possesses an additional enzymatic activity in which a DNA target-bound Cas12a can rapidly and non-specifically degrade any single-stranded DNA (ssDNA) substrate in a sequence-independent manner. This enzymatic activity is endonucleolytic, which means that the ssDNA substrate does not need a free 5' or 3' end to be cut. For this reason, natural Cas12a-type enzymes have the potential to induce off-target genome editing due to nicking of exposed ssDNA in cells.

## CONTACT

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## INVENTORS

» Doudna, Jennifer A.

## OTHER INFORMATION

### KEYWORDS

genome editing, Cas12a

### CATEGORIZED AS

» **Agriculture & Animal Science**

» Animal Science

» Plant Traits

» Plant Varieties

» Transgenics

» **Biotechnology**

» Food

» Genomics

» Health

» Proteomics

» **Environment**

» Other

» **Medical**

» Gene Therapy

» Research Tools

» Screening

» Therapeutics

» **Veterinary**

» Other

» Therapeutics

## RELATED CASES

2019-162-0

## ADDITIONAL TECHNOLOGIES BY THESE INVENTORS

- ▶ COMPOSITIONS AND METHODS FOR IDENTIFYING HOST CELL TARGET PROTEINS FOR TREATING RNA VIRUS INFECTIONS
- ▶ Genome Editing via LNP-Based Delivery of Efficient and Stable CRISPR-Cas Editors
- ▶ Type III CRISPR-Cas System for Robust RNA Knockdown and Imaging in Eukaryotes
- ▶ Cas12-mediated DNA Detection Reporter Molecules
- ▶ Improved guide RNA and Protein Design for CasX-based Gene Editing Platform
- ▶ Cas13a/C2c2 - A Dual Function Programmable RNA Endoribonuclease
- ▶ RNA-directed Cleavage and Modification of DNA using CasY (CRISPR-CasY)
- ▶ CasX Nickase Designs, Tans Cleavage Designs & Structure
- ▶ In Vivo Gene Editing Of Tau Locus Via Liponanoparticle Delivery
- ▶ A Dual-RNA Guided CasZ Gene Editing Technology
- ▶ CRISPR-CAS EFFECTOR POLYPEPTIDES AND METHODS OF USE THEREOF ("Cas-VariPhi")
- ▶ Modifications To Cas9 For Passive-Delivery Into Cells
- ▶ A Protein Inhibitor Of Cas9
- ▶ RNA-directed Cleavage and Modification of DNA using CasX (CRISPR-CasX)
- ▶ Compositions and Methods for Genome Editing
- ▶ Split-Cas9 For Regulatable Genome Engineering
- ▶ Methods to Interfere with Prokaryotic and Phage Translation and Noncoding RNA
- ▶ CRISPR CASY COMPOSITIONS AND METHODS OF USE
- ▶ Single Conjugative Vector for Genome Editing by RNA-guided Transposition
- ▶ CRISPR-CAS EFFECTOR POLYPEPTIDES AND METHODS OF USE THEREOF
- ▶ Engineered/Variant Hyperactive CRISPR CasPhi Enzymes And Methods Of Use Thereof
- ▶ Engineering Cas12a Genome Editors with Minimized Trans-Activity
- ▶ Methods Of Use Of Cas12L/CasLambda In Plants
- ▶ Type V CRISPR/CAS Effector Proteins for Cleaving ssDNA and Detecting Target DNA
- ▶ THERMOSTABLE RNA-GUIDED ENDONUCLEASES AND METHODS OF USE THEREOF (GeoCas9)
- ▶ Structure-Guided Methods Of Cas9-Mediated Genome Engineering
- ▶ Endoribonucleases For Rna Detection And Analysis
- ▶ Efficient Site-Specific Integration Of New Genetic Information Into Human Cells
- ▶ CRISPR-Cas Effector Polypeptides and Methods of Use Thereof
- ▶ Class 2 CRISPR/Cas COMPOSITIONS AND METHODS OF USE
- ▶ Compositions and Methods of Use for Variant Csy4 Endoribonucleases
- ▶ Identification Of Sites For Internal Insertions Into Cas9
- ▶ Methods and Compositions for Controlling Gene Expression by RNA Processing



### University of California, Berkeley Office of Technology Licensing

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