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# **ADAMTS4 Knockout 293T Cell Line, Homozygous**

Catalog No.: RM02096

#### **Basic Information**

#### Catalog No.

RM02096

#### Category

Cell Line

#### **Parental Cell line**

293T

#### Genotype

Knockout

#### **Gene Information**

#### **Gene Symbol**

ADAMTS4

#### **Species**

Human

#### **Gene ID**

9507

#### **Swiss Prot**

075173

#### Synonyms

ADAMTS-2; ADAMTS-4; ADMP-1

#### **Contact**

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### **Background**

This gene encodes a member of the ADAMTS (a disintegrin and metalloproteinase with thrombospondin motifs) protein family. Members of this family share several distinct protein modules, including a propeptide region, a metalloproteinase domain, a disintegrin-like domain, and a thrombospondin type 1 (TS) motif. Individual members of this family differ in the number of C-terminal TS motifs, and some have unique C-terminal domains. The enzyme encoded by this gene lacks a C-terminal TS motif. The encoded preproprotein is proteolytically processed to generate the mature protease. This protease is responsible for the degradation of aggrecan, a major proteoglycan of cartilage, and brevican, a brain-specific extracellular matrix protein. The expression of this gene is upregulated in arthritic disease and this may contribute to disease progression through the degradation of aggrecan. Alternative splicing results in multiple transcript variants, at least one of which encodes an isoform that is proteolytically processed. [provided by RefSeq, Feb 2016]

#### **Product Information**

#### Description

ADAMTS4 Knockout 293T Cell Line is engineered from 293T cell line with Gene-Editing Technology.

Allele-1:86bp deletion in exon1

Allele-2:86bp deletion in exon1

Mammalian cells such as human, rat and mouse cells are normally diploid with two alleles. Homozygote: both alleles were knocked out, mRNA has no signal, no expression of proteins. Heterozygote: only one allele was knocked out, the mRNA transcript levels was decreased compared to wild type, and the protein expression levels was also lower than that of the wild type.

#### **Packaging**

1 vial parental cell line and 1 vial knockout cell line

## **Shipping Conditions**

**Amount** 

Dry ice

1~5x10<sup>6</sup> cells/vial

#### Storage

Stored in liquid nitrogen for a long time less than -130°C. Minimizing freeze-thaw cycles.

#### Protoco

Upon arrival, it should be maintained in DMEM medium with 10%(v/v) fetal bovine serum and 100U penicillin-streptomycin, at  $37^{\circ}C$  with 5% CO<sub>2</sub> condition.

- 1. Thaw the vial in 37°C water bath ,and shake it to melt as soon as possible.
- 2. Transfer the cell suspension to a 15mL conical tube with pre-warmed 5mL complete medium and centrifuge 1000rpm for approximately 5 minutes at room temperature.
- 3. Remove and discard the supernatant.
- 4. Resuspend the cell pellet with 1mL pre-warmed complete medium and seed in 10cm dish.
- 5. Add 8-10mL of complete medium.
- 6. Incubate the culture at 37°C incubator with 5% CO<sub>2</sub>.
- 7. A subcultivation ratio of 1:2-1:4 is recommended.

## Sequencing data

WT TCCTGCTCCCCATT\*\*\*\*\*\*\*\*\*\*\*GGAGGAGGAGATCG
Mut TCCTGCTCCCCATT\*\*\*Deletion\*\*\*GGAGGAGGAGATCG

Allele-1: 86bp deletion in exon1

WT TCCTGCTCCCCATT\*\*\*\*\*\*\*\*GGAGGAGGAGATCG
Mut TCCTGCTCCCCATT\*\*\*Deletion\*\*\*GGAGGAGGAGATCG

Allele-2: 86bp deletion in exon1

Genome sequence analysis of PCR products from parental (WT) and ADAMTS4 knockout (KO) 293T cells, using sanger sequencing.