

# MAP1LC3A Knockout 293T Cell Lysate, Homozygous

Catalog No.: RM50108

## Basic Information

### Catalog No.

RM50108

### Category

Cell Lysate

### Parental Cell line

293T

### Genotype

Knockout

## Gene Information

### Gene Symbol

MAP1LC3A<sup>293T</sup>

### Species

Human

### Gene ID

84557

### Swiss Prot

Q9H492

### Synonyms

LC3; LC3A; ATG8E; MAP1ALC3;  
MAP1BLC3; MAP1LC3A

## Contact

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## Background

MAP1A and MAP1B are microtubule-associated proteins which mediate the physical interactions between microtubules and components of the cytoskeleton. MAP1A and MAP1B each consist of a heavy chain subunit and multiple light chain subunits. The protein encoded by this gene is one of the light chain subunits and can associate with either MAP1A or MAP1B. Two transcript variants encoding different isoforms have been found for this gene. The expression of variant 1 is suppressed in many tumor cell lines, suggesting that may be involved in carcinogenesis.

## Product Information

### Description

MAP1LC3A Knockout cell line is engineered from 293T cell line with Gene-Editing Technology.

Allele-1:exon2 was deleted

Allele-2:exon2 was deleted

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 Lysate and 1 vial knockout cell Lysate

### Shipping Conditions

4°C

### Amount

50μL, 2μg/μL.

### Storage

Lysate is stable for 12 months when stored at -20°C. Minimizing freeze-thaw cycles.

### Protocol

To be used as WB control. Lysate is supplied in 1× SDS sample buffer (2% SDS, 60 mM Tris-HCl pH 6.8, 10% Glycerol, 0.02% Bromophenol blue, 60 mM beta-mercaptoethanol). Lysate should be boiled for 3 - 5 minutes before loading onto gel.

## Sequencing data

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WT GCTGTCCGAG\*\*\*\*\*GCGTCCCGACA  
Mut GCTGTCCGAG\*\*\*\*\*Deletion\*\*\*\*\*GCGTCCCGACA  
Allele-1: exon2 was deleted

WT GCTGTCCGAG\*\*\*\*\*GTTCCCGACAG  
Mut GCTGTCCGAG\*\*\*\*\*Deletion\*\*\*\*\*GTTCCCGACAG  
Allele-2: exon2 was deleted

Genome sequence analysis of PCR products from parental (WT) and MAP1LC3A knockout (KO) 293T cells, using sanger sequencing.