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# SUMO1 Knockout 293T cell lysate, Homozygous

Catalog No.: RM50191

#### **Basic Information**

#### Catalog No.

RM50191

#### Category

Cell Lysate

#### **Parental Cell line**

293T

#### Genotype

Knockout

## **Background**

This gene encodes a protein that is a member of the SUMO (small ubiquitin-like modifier) protein family. It functions in a manner similar to ubiquitin in that it is bound to target proteins as part of a post-translational modification system. However, unlike ubiquitin which targets proteins for degradation, this protein is involved in a variety of cellular processes, such as nuclear transport, transcriptional regulation, apoptosis, and protein stability. It is not active until the last four amino acids of the carboxy-terminus have been cleaved off. Several pseudogenes have been reported for this gene. Alternate transcriptional splice variants encoding different isoforms have been characterized.

#### **Gene Information**

#### **Gene Symbol**

SUM01

#### **Species**

Human

### Gene ID

7341

#### **Swiss Prot**

P63165

#### Synonyms

DAP1; GMP1; PIC1; SMT3; UBL1; OFC10; SENP2; SMT3C; SMT3H3; O1

#### **Contact**

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#### **Product Information**

#### Description

SUMO1 Knockout cell line is engineered from 293T cell line with Gene-Editing Technology. Allele-1:exon1 was deleted

Allele-2:exon1 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**

**Amount** 50μL, 2μg/μL.

## 4°C

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

#### **Protocol**

Storage

To be used as WB control. Lysate is supplied in  $1\times$  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

WT AGTGACGCGAGGCG\*GGCCAGGGCCTTCC
Mut AGTGACGCGAGGCG\*\*\*Deletion\*\*\*\*GGCCAGGGCCTTCC
Allele-1: exon1 was deleted

WT AGTGACGCGAGGCG\*\*\*\*\*\*\*\*\*\*\*GCCAGGGCCTTCCC
Mut AGTGACGCGAGGCG\*\*\*Deletion\*\*\*GCCAGGGCCTTCCC
Allele-2: exon1 was deleted

Genome sequence analysis of PCR products from parental (WT) and SUMO1 knockout (KO) 293T cells, using sanger sequencing.