

# GARS Knockdown 293T Cell Lysate, Heterozygous

Catalog No.: RM02261

## **Basic Information**

#### Catalog No.

RM02261

## Category

Cell Lysate

## **Parental Cell line**

293T

#### Genotype

Knockdown

## **Background**

This gene encodes glycyl-tRNA synthetase, one of the aminoacyl-tRNA synthetases that charge tRNAs with their cognate amino acids. The encoded enzyme is an (alpha)2 dimer which belongs to the class II family of tRNA synthetases. It has been shown to be a target of autoantibodies in the human autoimmune diseases, polymyositis or dermatomyositis. Two transcript variants encoding different isoforms have been found for this gene. [provided by RefSeq, Oct 2015]

#### **Gene Information**

## **Gene Symbol**

GARS

## **Species**

Human

## **Gene ID**

2617

#### **Swiss Prot**

P41250

## **Synonyms**

CMT2D; DSMAV; GlyRS; HMN5; SMAD1

## **Contact**

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## **Product Information**

## **Description**

GARS Knockdown 293T Cell Line is engineered from 293T cell line with Gene-Editing technology.

Allele-1:96bp deletion in exon1

Allele-2:97bp 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 Lysate and 1 vial knockout cell Lysate

## **Shipping Conditions**

Amount

4°C

50μL, 2μg/μL.

#### Storage

Lysate is stable for 12 months when stored at -20  $^{\circ}$ C. Minimizing freeze-thaw cycles.

#### **Protocol**

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 CTGCCGCCCCGGCT\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*GCGCGGGGGCTGA
Mut CTGCCGCCCCGGCT\*\*\*Deletion\*\*\*\*GGCGCGGGGGCTGA

Allele-1: 96bp deletion in exon1

WT TGCCGCCCGGCTC\*\*\*\*\*\*\*\*\*\*\*\*GGCGCGGGGGCTGA
Mut TGCCGCCCCGGCTC\*\*\*Deletion\*\*\*GGCGCGGGGGCTGA

Allele-2: 97bp deletion in exon1

Genome sequence analysis of PCR products from parental (WT) and GARS Knockdown (KD) 293T cells, using sanger sequencing.