

# SLC38A8 Knockout 293T Cell Lysate, Homozygous

**Catalog No.:** RM50104

## Basic Information

**Catalog No.**

RM50104

**Category**

Cell Lysate

**Parental Cell line**

293T

**Genotype**

Knockout

## Gene Information

**Gene Symbol**

SLC38A8

**Species**

Human

**Gene ID**

146167

**Swiss Prot**

A6NNN8

**Synonyms**

FVH2; FHASD; SNAT8

## Contact

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## Background

This gene encodes a putative sodium-dependent amino-acid/proton antiporter. The protein has eleven transmembrane domains, an extracellular N-terminus and an intracellular C-terminal tail. The protein is a member of the SLC38 sodium-coupled neutral amino acid transporter family of proteins. Mutations in this gene result in foveal hypoplasia with or without optic nerve misrouting and/or anterior segment dysgenesis.

## Product Information

**Description**

SLC38A8 Knockout cell line is engineered from 293T cell line with Gene-Editing Technology.

Allele-1:158bp deletion in exon1

Allele-2:158bp 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**

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 GTCTTCCTGAT\*\*\*\*\*TCGGGGACCAGC  
Mut GTCTTCCTGAT\*\*Deletion(158bp)\*\*TCGGGGACCAGC  
Allele-1: 158 bp deletion in exon1

WT GTCTTCCTGAT\*\*\*\*\*TCGGGGACCAGC  
Mut GTCTTCCTGAT\*\*Deletion(158bp)\*\*TCGGGGACCAGC  
Allele-2: 158 bp deletion in exon1

Genome sequence analysis of PCR products from parental (WT) and SLC38A8 knockout (KO) 293T cells, using sanger sequencing.