

CHP1 Knockout 293T Cell Lysate, Homozygous

Catalog No.: RM50101

Basic Information

Catalog No.

RM50101

Category

Cell Lysate

Parental Cell line

293T

Genotype

Knockout

Gene Information

Gene Symbol

CHP1

Species

Human

Gene ID

11261

Swiss Prot

Q99653

Synonyms

CHP; p22; p24; SPAX9; Sid470p;
SLC9A1BP; CHP1

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Background

This gene encodes a phosphoprotein that binds to the Na⁺/H⁺ exchanger NHE1. This protein serves as an essential cofactor which supports the physiological activity of NHE family members and may play a role in the mitogenic regulation of NHE1. The protein shares similarity with calcineurin B and calmodulin and it is also known to be an endogenous inhibitor of calcineurin activity.

Product Information

Description

CHP1 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

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

WT GGAGCTCCCGCAC*****TAAGGTCTGGAGC
Mut GGAGCTCCCGCAC***Deletion***TAAGGTCTGGAGC
Allele-1: exon1 was deleted

WT AGGAGTCCCGCA*****TAAGGTCTGGAGC
Mut AGGAGTCCCGCA***Deletion***TAAGGTCTGGAGC
Allele-2: exon1 was deleted

Genome sequence analysis of PCR products from parental (WT) and CHP1 knockout (KO) 293T cells, using sanger sequencing.