abcam

Product datasheet

Human TMED10 (TMP21) knockout HEK-293T cell line ab266227

4 Images

Overview

Product name Human TMED10 (TMP21) knockout HEK-293T cell line

Parental Cell Line HEK293T
Organism Human

Mutation description Knockout achieved by using CRISPR/Cas9, Homozygous: 2 bp insertion in exon 1

Passage number <20

Knockout validation Sanger Sequencing, Western Blot (WB)

Tested applications Suitable for: WB

Biosafety level 2

General notesRecommended control: Human wild-type HEK293T cell line (ab255449). Please note a wild-type cell line is not automatically included with a knockout cell line order, if required please add

recommended wild-type cell line at no additional cost using the code WILDTYPE-TMTK1.

Cryopreservation cell medium: Cell Freezing Medium-DMSO Serum free media, contains

8.7% DMSO in MEM supplemented with methyl cellulose.

Culture medium: DMEM (High Glucose) + 10% FBS

Initial handling guidelines: Upon arrival, the vial should be stored in liquid nitrogen vapor phase and not at -80°C. Storage at -80°C may result in loss of viability.

- 1. Thaw the vial in 37°C water bath for approximately 1-2 minutes.
- 2. Transfer the cell suspension (0.8 mL) to a 15 mL/50 mL conical sterile polypropylene centrifuge tube containing 8.4 mL pre-warmed culture medium, wash vial with an additional 0.8 mL culture medium (total volume 10 mL) to collect remaining cells, and centrifuge at 201 x g (rcf) for 5 minutes at room temperature. 10 mL represents minimum recommended dilution. 20 mL represents maximum recommended dilution.
- 3. Resuspend the cell pellet in 5 mL pre-warmed culture medium and count using a haemocytometer or alternative cell counting method. Based on cell count, seed cells in an appropriate cell culture flask at a density of 2x10⁴ cells/cm². Seeding density is given as a guide only and should be scaled to align with individual lab schedules.
- 4. Incubate the culture at 37°C incubator with 5% CO₂. Cultures should be monitored daily.

Subculture guidelines:

All seeding densities should be based on cell counts gained by established methods. A guide seeding density of $2x10^4$ cells/cm² is recommended.

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A partial media change 24 hours prior to subculture may be helpful to encourage growth, if required.

Cells should be passaged when they have achieved 80-90% confluence.

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We will provide viable cells that proliferate on revival.

Properties

Number of cells 1 x 10⁶ cells/vial, 1 mL

Adherent /Suspension Adherent
Tissue Kidney
Cell type epithelial

STR Analysis Amelogenin X D5S818: 8, 9 D13S317: 12, 14 D7S820: 11 D16S539: 9, 13 vWA: 16, 19 TH01:

7, 9.3 TPOX: 11 CSF1PO: 11, 12

Mycoplasma free Yes

Storage instructions Shipped on Dry Ice. Store in liquid nitrogen.

Storage buffer Constituents: 8.7% Dimethylsulfoxide, 2% Cellulose, methyl ether

Target

Function

Involved in vesicular protein trafficking. Mainly functions in the early secretory pathway. Thought to act as cargo receptor at the lumenal side for incorporation of secretory cargo molecules into transport vesicles and to be involved in vesicle coat formation at the cytoplasmic side. In COPII vesicle-mediated anterograde transport involved in the transport of GPI-anchored proteins and proposed to act togther with TMED2 as their cargo receptor; the function specifically implies SEC24C and SEC24D of the COPII vesicle coat and lipid raft-like microdomains of the ER. Recognizes GPI anchors structural remodeled in the ER by PGAP1 and MPPE1 (By similarity). In COPI vesicle-mediated retrograde transport involved in the biogenesis of COPI vesicles and vesicle coat recruitment. On Golgi membranes, acts as primary receptor for ARF1-GDP which is involved in COPI-vesicle formation. Increases coatomer-dependent GTPase-activating activity of ARFGAP2. Involved in trafficking of G protein-coupled receptors (GPCRs). Regulates F2LR1, OPRM1 and P2RY4 exocytic trafficking from the Golgi to the plasma membrane thus contributing to receptor resensitization. Involved in trafficking of amyloid beta A4 protein and soluble APP-beta release (independent of modulation of gamma-secretase activity). As part of the presenilindependent gamma-secretase complex regulates gamma-cleavages of the amyloid beta A4 protein to yield amyloid-beta 40 (Abeta40). Involved in organization of the Golgi apparatus.

Tissue specificity Ubiquitous.

Sequence similaritiesBelongs to the EMP24/GP25L family.

Contains 1 GOLD domain.

DomainThe lumenal domain mediates localization to the plasma membrane by partially overriding the ER

retention by the cytoplasmic domain.

Cellular localization Golgi apparatus > cis-Golgi network membrane. Melanosome. Endoplasmic reticulum

membrane. Endoplasmic reticulum-Golgi intermediate compartment membrane. Cytoplasmic

vesicle > secretory vesicle membrane. Cell membrane. Golgi apparatus > trans-Golgi network membrane. Identified by mass spectrometry in melanosome fractions from stage I to stage IV. Cycles between compartments of the early secretatory pathway.

Applications

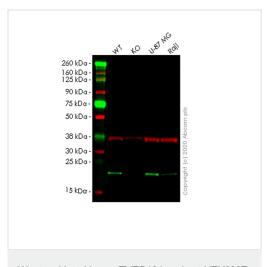
The Abpromise guarantee

Our <u>Abpromise guarantee</u> covers the use of ab266227 in the following tested applications.

The application notes include recommended starting dilutions; optimal dilutions/concentrations should be determined by the end user.

Application	Abreviews	Notes
WB		Use at an assay dependent concentration. Predicted molecular weight: 25 kDa.

Images



Western blot - Human TMED10 knockout HEK293T cell line (ab266227)

All lanes : Anti-TMP21 antibody [EPR9036(B)] (ab134948) at 1/1000 dilution

Lane 1: Wild-type HEK293T cell lysate

Lane 2: TMED10 knockout HEK293T cell lysate

Lane 3: U-87 MG cell lysate

Lane 4: Raji cell lysate

Lysates/proteins at 20 µg per lane.

Secondary

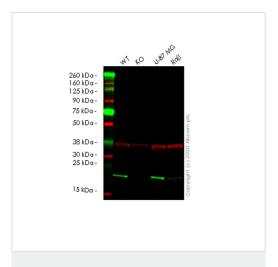
All lanes : Goat anti-Rabbit lgG H&L (IRDye® 800CW) preadsorbed (ab216773) at 1/10000 dilution

Predicted band size: 25 kDa **Observed band size:** 19 kDa

Lanes 1-4: Merged signal (red and green). Green - <u>ab134948</u> observed at 19 kDa. Red - loading control <u>ab8245</u> observed at 36 kDa.

<u>ab134948</u> Anti-TMP21 antibody [EPR9036(B)] was shown to specifically react with TMP21 in wild-type HEK293T cells. Loss of signal was observed when knockout cell line ab266227 (knockout cell lysate <u>ab258233</u>) was used. Wild-type and TMP21 knockout samples were subjected to SDS-PAGE. <u>ab134948</u> and Anti-GAPDH antibody [6C5] - Loading Control (<u>ab8245</u>) were incubated

at room temperature for 2.5 hours at 1 in 1000 dilution and 1 in 20000 dilution respectively. Blots were developed with Goat anti-Rabbit lgG H&L (IRDye[®] 800CW) preadsorbed (ab216773) and Goat anti-Mouse lgG H&L (IRDye[®] 680RD) preadsorbed (ab216776) secondary antibodies at 1 in 20000 dilution for 1 hour at room temperature before imaging.



Western blot - Human TMED10 knockout HEK293T cell line (ab266227)

All lanes : Anti-TMP21 antibody [EPR9037(B)] (ab133771) at 1/1000 dilution

Lane 1: Wild-type HEK293T cell lysate

Lane 2: TMED10 knockout HEK293T cell lysate

Lane 3: U-87 MG cell lysate

Lane 4: Raji cell lysate

Lysates/proteins at 20 µg per lane.

Secondary

All lanes : Goat anti-Rabbit lgG H&L (IRDye® 800CW) preadsorbed (**ab216773**) at 1/10000 dilution

Predicted band size: 25 kDa **Observed band size:** 19 kDa

Lanes 1-4: Merged signal (red and green). Green - <u>ab133771</u> observed at 19 kDa. Red - loading control <u>ab8245</u> observed at 36 kDa.

<u>ab133771</u> Anti-TMP21 antibody [EPR9037(B)] was shown to specifically react with TMP21 in wild-type HEK293T cells. Loss of signal was observed when knockout cell line ab266227 (knockout cell lysate <u>ab258233</u>) was used. Wild-type and TMP21 knockout samples were subjected to SDS-PAGE. <u>ab133771</u> and Anti-GAPDH antibody [6C5] - Loading Control (<u>ab8245</u>) were incubated at room temperature for 2.5 hours at 1 in 1000 dilution and 1 in 20000 dilution respectively. Blots were developed with Goat anti-Rabbit lgG H&L (IRDye[®] 800CW) preadsorbed (<u>ab216773</u>) and Goat anti-Mouse lgG H&L (IRDye[®] 680RD) preadsorbed (<u>ab216776</u>) secondary antibodies at 1 in 20000 dilution for 1 hour at room temperature before imaging.

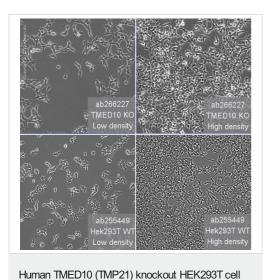
Mut CATGTCTGGTTTGTCTGGCCCACCAGCCCGCGGCGCCCCTTTTCCGTTAGCGTTGCTG

WT CATGTCTGGTTTGTCTGGCCCACCAGCCCG GCGCGGCCCTTTTCCGTTAGCGTTGCTG

Sanger Sequencing - Human TMED10 knockout

HEK293T cell line (ab266227)

Homozygous: 2 bp insertion in exon1



line (ab266227)

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