abcam

Product datasheet

Human PARD6A (PAR6) knockout HeLa cell line ab265171

画像数 2

製品の概要

製品名 Human PARD6A (PAR6) knockout HeLa cell line

Parental Cell Line HeLa
Organism Human

Mutation description Knockout achieved by using CRISPR/Cas9, 1 bp deletion in exon 3 and 1 bp insertion in exon 3

Passage number <20

Knockout validation Sanger Sequencing

Biosafety level

特記事項

Recommended control: Human wild-type HeLa cell line (<u>ab255928</u>). 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 quidelines:

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

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.

製品の特性

Cell type

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

Adherent / Suspension Adherent
Tissue Cervix

Disease Adenocarcinoma

Gender Female

STR Analysis Amelogenin X D5S818: 11, 12 D13S317: 12, 13.3 D7S820: 8, 12 D16S539: 9, 10 vWA: 16, 18

TH01: 7 TPOX: 8,12 CSF1PO: 9, 10

Mycoplasma free Yes

保存方法 Shipped on Dry Ice. Store in liquid nitrogen.

epithelial

パップァー Constituents: 8.7% Dimethylsulfoxide, 2% Cellulose, methyl ether

ターゲット情報

機能 Adapter protein involved in asymmetrical cell division and cell polarization processes. Probably

involved in the formation of epithelial tight junctions. Association with PARD3 may prevent the interaction of PARD3 with F11R/JAM1, thereby preventing tight junction assembly. The PARD6-PARD3 complex links GTP-bound Rho small GTPases to atypical protein kinase C proteins.

組織特異性 Expressed in pancreas, skeletal muscle, brain and heart. Weakly expressed in kidney and

placenta.

配列類似性 Belongs to the PAR6 family.

Contains 1 OPR domain.
Contains 1 PDZ (DHR) domain.
Contains 1 pseudo-CRIB domain.

ドメイン The pseudo-CRIB domain together with the PDZ domain is required for the interaction with Rho

small GTPases.

The OPR domain mediates interactions with MAP2K5. The PDZ domain mediates the interaction with CRB3.

細胞内局在 Cytoplasm. Cell membrane. Cell projection > ruffle. Cell junction > tight junction. Colocalizes with

GTP-bound CDC42 or RAC1 at membrane ruffles and with PARD3 and PRKCI at epithelial tight

junctions.

画像

Mut	TAGACGT GGACCTACT GCCT GAGACCCACC - ACGGGT GCGGCT GCACAAGCAT GGT T CAG	
WT	${\sf TAGACGTGGACCTACTGCCTGAGACCCACCGACGGGTGCGGCTGCACAAGCATGGTTCAG}$	
Sanger Sequencing - Human PARD6A knockout		
Sanger Sequencing - Human FANDON Knockout		
HeLa cell line (ab265171)		

Allele-1: 1 bp deletion in exon 3.

Mut	TAGACGT GGACCTACTGCCTGAGACCCACC <mark>A</mark> GACGGGT GCGGCTGCACAAGCATGGTTCA
WT	TAGACGTGGACCTACTGCCTGAGACCCACC GACGGGTGCGGCTGCACAAGCATGGTTCA

Allele-2: 1 bp insertion in exon 3.

Sanger Sequencing - Human PARD6A knockout HeLa cell line (ab265171)

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