# abcam

### Product datasheet

# Human PARP1 knockout HEK-293T cell line ab266598

#### 画像数 4

#### 製品の概要

特記事項

製品名 Human PARP1 knockout HEK-293T cell line

Parental Cell Line HEK293T
Organism Human

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

Passage number <20

**Knockout validation** Sanger Sequencing, Western Blot (WB)

アプリケーション **適用あり**: WB

Biosafety level

-

**Recommended control:** Human wild-type HEK293T cell line (<u>ab255449</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 2x10<sup>4</sup> cells/cm<sup>2</sup>. 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% CO<sub>2</sub>. 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/cm<sup>2</sup> is recommended.

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

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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

Number of cells 1 x 10<sup>6</sup> 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

Antibiotic resistance Puromycin 1.00µg/ml

Mycoplasma free Yes

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

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

#### ターゲット情報

機能 Involved in the base excision repair (BER) pathway, by catalyzing the poly(ADP-ribosyl)ation of a

limited number of acceptor proteins involved in chromatin architecture and in DNA metabolism.

This modification follows DNA damages and appears as an obligatory step in a

detection/signaling pathway leading to the reparation of DNA strand breaks. Mediates the poly(ADP-ribosyl)ation of APLF and CHFR. Positively regulates the transcription of MTUS1 and

negatively regulates the transcription of MTUS2/TIP150.

**配列類似性** Contains 1 BRCT domain.

Contains 1 PARP alpha-helical domain. Contains 1 PARP catalytic domain. Contains 2 PARP-type zinc fingers.

翻訳後修飾 Phosphorylated by PRKDC. Phosphorylated upon DNA damage, probably by ATM or ATR.

Poly-ADP-ribosylated by PARP2. Poly-ADP-ribosylation mediates the recruitment of CHD1L to

DNA damage sites.

S-nitrosylated, leading to inhibit transcription regulation activity.

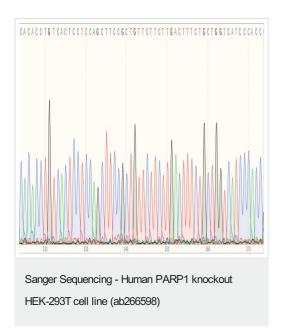
細胞内局在 Nucleus.

#### アプリケーション

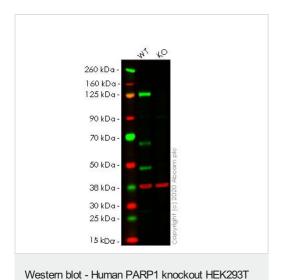
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アプリケーション	Abreviews	特記事項
WB		Use at an assay dependent concentration. Predicted molecular weight: 113 kDa.

#### 画像



Sequencing chromatogram displaying sequence edit in exon 2



cell line (ab266598)

**All lanes :** Anti-PARP1 antibody [E102] (<u>ab32138</u>) at 1/1000 dilution

Lane 1: Wild-type HEK-293T cell lysate

Lane 2: PARP1 knockout HEK-293T cell lysate

Lysates/proteins at 20 µg per lane.

Performed under reducing conditions.

**Predicted band size:** 113 kDa **Observed band size:** 113 kDa

**Lanes 1-2:** Merged signal (red and green). Green - <u>ab32138</u> observed at 113 kDa. Red - Anti-GAPDH antibody [6C5] - Loading Control (<u>ab8245</u>) observed at 37 kDa.

<u>ab32138</u> was shown to react with PARP1 in wild-type HEK-293T cells in western blot. Loss of signal was observed when knockout cell line ab266598 (knockout cell lysate <u>ab257017</u>) was used. Wild-

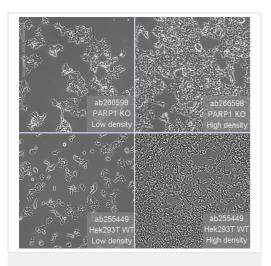
type HEK-293T and PARP1 knockout HEK-293T cell lysates were subjected to SDS-PAGE. <a href="mailto:ab32138">ab32138</a> and Anti-GAPDH antibody [6C5] - Loading Control (ab8245) overnight at 4°C at a 1 in 1000 dilution and a 1 in 20000 dilution respectively. Blots were developed with Goat anti-Rabbit IgG H&L (IRDye®800CW) preadsorbed (ab216773) and Goat anti-Mouse IgG H&L (IRDye®680RD) preadsorbed (ab216776) secondary antibodies at 1 in 20000 dilution for 1 hour at room temperature before imaging.

Mut CACACCTGTCACTCCTCCAGCTTCCGCTGTTCTTCTTGACTTTCTGCTGGTCATCCCACC

WT CACACCTGTCACTCCTCCAGCTTCCGCTGT CTTCTTGACTTTCTGCTGGTCATCCCACC

Sanger Sequencing - Human PARP1 knockout HEK293T cell line (ab266598)

Homozygous: 1 bp insertion in exon 2



Cell Culture - Human PARP1 knockout HEK293T cell line (ab266598)

Representative images of PARP1 knockout HEK293T cells, low and high confluency examples (top left and right respectively) and wild-type HEK293T cells, low and high confluency (bottom left and right respectively) showing typical adherent, epithelial-like morphology. Images were captured at 10X magnification using an EVOS M5000 microscope.

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