

# Mechanism of hypoxia-specific cytotoxicity of procaspase-3 fused with a VHL-mediated protein destruction motif of HIF-1 $\alpha$ containing Pro564

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**Abstract** Under normoxic conditions the alpha-subunit of hypoxia-inducible factor (HIF-1 $\alpha$ ) protein is targeted for degradation by the von Hippel-Lindau (VHL) tumor suppressor protein acting as an E3 ubiquitin ligase. Recently, we developed a hypoxia-targeting protein, TOP3, which consisted of procaspase-3 with the VHL-mediated protein destruction motif of HIF-1 $\alpha$ . This design enables procaspase-3 to be regulated similarly with HIF-1 $\alpha$ , being degraded under normoxia while stabilized under hypoxia. Furthermore, stabilized TOP3 was cleaved by the hypoxic stress-induced endogenous caspases and thus the procaspase-3 was converted to active caspase-3 specifically under hypoxic conditions. These data demonstrated that the VHL-mediated protein destruction motif of HIF-1 $\alpha$  endowed procaspase-3 with hypoxia-specific cytotoxicity.

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## 1. Introduction

Hypoxia-inducible factor (HIF) is a transcriptional complex that mediates a broad range of cellular and systemic responses to hypoxia [1]. HIF-1 is a heterodimer composed of  $\alpha$  and  $\beta$  subunits, and  $\alpha$  subunit of HIF-1 (HIF-1 $\alpha$ ) is regulated in an oxygen-dependent manner at the post-translational level [2–4]. HIF-1 $\alpha$  contains oxygen-dependent degradation (ODD) domains, which contains proline residues, proline-402 and proline-564. In normoxia, HIF-1 $\alpha$  is hydroxylated at these proline residues by prolyl-4-hydroxylases [5,6]. The modification accelerates the interaction of the HIF-1 $\alpha$  protein with the von Hippel-Lindau (VHL) tumor suppressor protein, resulting in the rapid ubiquitination and subsequent degradation of the protein by the 26S proteasome [7–10].

Recently, we screened ODD domain mutants of human HIF-1 $\alpha$  protein and determined the ODD<sub>548–603</sub>, which endowed fusion proteins with sufficient oxygen-dependent degradation regulation [11]. In order to specifically eradicate HIF-1 $\alpha$ -expressing hypoxic cells in solid tumors, we con-

structed procaspase-3 fused with ODD<sub>548–603</sub> and HIV-Tat protein-transduction domain (PTD). The final product was named TOP3 (Tat-ODD-Procaspase-3) [11–14]. The HIV-Tat PTD domain is derived from human immunodeficiency virus type-1 Tat protein and efficiently delivers TOP3 to any tissue in vivo. The procaspase-3 comes from human caspase-3 protein [15] and confers cytosolic activity to TOP3. We already reported that TOP3 had anti-tumor activity in xenografts model with various cancer cells [11–14] and induced apoptosis to hypoxic tumor cells in xenografts [14].

In the present study, we clarify the mechanism of TOP3 activation and confirm its hypoxia-specific cytotoxicity.

## 2. Materials and methods

### 2.1. Cell culture and hypoxic treatment in vitro

CFPAC-1 and MIA PaCa-2 human pancreatic cancer cell lines, HeLa human cervical epithelial adenocarcinoma cell line, A549 human lung adenocarcinoma cell line, WiDr human colorectal adenocarcinoma cell line and 786-O human renal cell carcinoma cell line were purchased from the American Type Culture Collection. HeLa/EF-Luc and HeLa/5HRE-Luc cell clones were isolated as described previously [14]. WT8, a 786-O cell clone stably transfected with a plasmid coding hemagglutinin (HA)-tagged VHL, was a kind gift from Dr. William G. Kaelin Jr. [16]. CFPAC-1 was maintained as described previously [11], and the other cells were maintained at 37 °C in 5% FBS-Dulbecco's modified Eagle's medium (Nacalai Tesque, Kyoto, Japan) supplemented with penicillin (100 units/ml) and streptomycin (100  $\mu$ g/ml).

Hypoxic condition of <0.02% of oxygen tension was attained by the use of a Bactron Anaerobic Chamber, BACLITE-1 (Sheldon Manufacturing Inc., Cornelius, OR). Cells were incubated in the chamber for at least 6 h before various treatments.

### 2.2. Formulation of TOP3 fusion protein

The Tat-ODD-Procaspase-3 fusion protein (TOP3) was prepared and dissolved in 10 mM Tris-HCl buffer (pH 8.0), as described previously [11]. The final concentration of TOP3 preparation was 15  $\mu$ g/ml for in vitro experiments if not indicated, and 10 mM Tris-HCl (pH 8.0) was used as the buffer in both in vivo and in vitro experiments if not indicated.

### 2.3. Experimental procedures for analyses of TOP3 activation in vitro

Cells were pre-incubated under aerobic or hypoxic condition for 6 h, added with TOP3, and incubated further for 20 h before analyses.

As for the Western blot analysis, cells were seeded at  $1 \times 10^5$  cells/well in a 6-well plate, treated as above and the lysates were prepared by suspending cells from each well in 100  $\mu$ l of 1 $\times$  loading buffer. Twenty  $\mu$ l of the lysate was electrophoresed per lane on a 15% (for

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TOP3/caspase-3), a 10% (for VHL) or a 7.5% (for HIF-1 $\alpha$ ) SDS-polyacrylamide gel. TOP3/caspase-3, VHL and HIF-1 $\alpha$  were detected by polyclonal anti-caspase-3 antibody (Cell Signaling Technology Inc., Osaka, Japan), monoclonal anti-HA antibody (Roche Diagnostics Japan, Tokyo, Japan) and monoclonal anti-HIF-1 $\alpha$  antibody (BD Bioscience Pharmingen, San Diego, CA), respectively. The polyclonal and the monoclonal primary antibodies were then reacted with anti-rabbit and anti-mouse IgG horseradish peroxidase linked antibodies (Amersham Biosciences Corp., Piscataway, NJ), respectively. Detection was carried out with a chemiluminescence-based method using the ECL-PLUS system (Amersham Biosciences Corp.).

As for the DNA fragmentation analysis, genomic DNA was isolated from  $6 \times 10^5$  cells with Quick Apoptotic Ladder Detection Kit (BioVision Research Products, Mountain View, CA). DNA was electrophoresed in a 1.5% agarose gel.

As for the FACS analysis, the cells were harvested, gently suspended in PBS and mixed with equal volume of  $2 \times$  hypotonic fluorochrome solution (100  $\mu$ g/ml propidium iodide in 0.2% sodium citrate–0.2% Triton X-100) immediately before the analysis with a flow cytometry using CELLQuest (BD Biosciences, Franklin Lakes, NJ).

As for the analysis of caspase-3 after TOP-3 treatment, the cells were seeded at  $2 \times 10^4$  cells/well in a 24-well plate, pre-incubated for 16 h and treated with TOP3 for 0, 2, 4, 6 or 8 h. As for the analysis of endogenous caspase-3 and -9 activities, similarly seeded cells were cultured with the medium, which was pre-exposed to hypoxic conditions. Then the cells were incubated under hypoxic or aerobic conditions and harvested after indicated time of incubation. Caspase-3 and -9 activi-

ties in 50  $\mu$ l lysates were measured by using Colorimetric Protease Assay Kit according to the manufacturer's instructions (MBL, Nagoya, Japan). The experiments were done in triplicate and the mean of OD<sub>405nm</sub>/50  $\mu$ g was calculated.

### 3. Results

#### 3.1. Hypoxia-dependent activation of TOP3

We previously showed cytotoxic effect of TOP3 on hypoxic cells in the xenografts of a human pancreatic cancer cell line [11,14]. TOP3 was designed to be degraded in normoxic cells through the function of ODD domain of human HIF-1 $\alpha$  protein which is sensitive to the VHL-mediated destruction in normoxia. The design allows the conversion of the procaspase-3 domain to active caspase-3 under hypoxia, executing apoptotic killing of the cells.

These expectations were tested as follows. Firstly, the stability of TOP3 was compared with that of HIF-1 $\alpha$  protein under aerobic and hypoxic conditions (Fig. 1A). CFPAC-1 cells were cultured under aerobic (lanes 1 and 2) and hypoxic (lanes 3 and 4) conditions and the protein levels of TOP3 and HIF-1 $\alpha$  were tested with (lanes 2 and 4) or without (lanes 1 and 3) addition

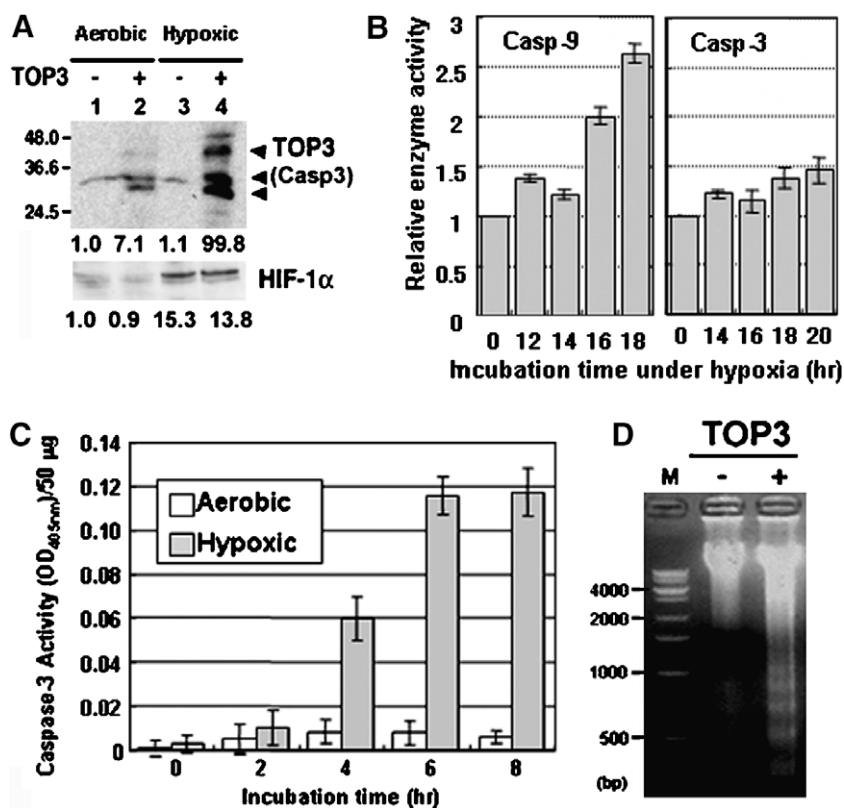


Fig. 1. Hypoxia-dependent stabilization and activation of TOP3 in vitro. (A) CFPAC-1 cells were treated with the buffer only (–; lanes 1 and 3) or TOP3 (+; lanes 2 and 4) under aerobic (lanes 1 and 2) or hypoxic (lanes 3 and 4) conditions for 20 h. TOP3 (upper panel) and HIF-1 $\alpha$  (lower panel) proteins in the cells were analyzed by Western blotting with a polyclonal anti-caspase-3 and a monoclonal HIF-1 $\alpha$  antibody, respectively. TOP3, its major derivative proteins and endogenous caspase-3 are indicated by arrowheads. The relative density of the bands of TOP3 (upper panel) and HIF-1 $\alpha$  (lower panel) proteins of the lanes 2–4 to the lane 1 are indicated below each panel, respectively. (B) CFPAC-1 cells were cultured under hypoxic conditions for indicated period of time and then cell lysates were prepared. Caspase-9 (left) and -3 (right) activities in the cell lysates were measured and relative caspase activities to the 0 h are indicated. (C) CFPAC-1 cells were cultured under aerobic (open bars) and hypoxic (gray bars) conditions for 16 h and then treated with TOP3 or the buffer for the indicated period of time. Total cell lysates were prepared and then OD<sub>405 nm</sub> of the lysates was measured. TOP3-derived caspase-3 activity was calculated using the following formula: (OD<sub>405 nm</sub>/50  $\mu$ g of the TOP3-treated cells) – (OD<sub>405 nm</sub>/50  $\mu$ g of the buffer-treated cells). Results are the means of three independent experiments  $\pm$  S.D. (D) The cells were treated with the buffer (–) and TOP3 (+) for 20 h under hypoxic conditions. Then the genomic DNA was isolated and analyzed by electrophoresis with a 1.5% agarose gel.

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