

FAST TRACKS

## Phenotypic Expression of Human Hepatoma Cells in Culture

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**Abstract** Hepatomas thrive in a hypoxic environment resulting in the induction of a cluster of hypoxia related genes. The protein phenotypic expression include hypoxia inducible factor- $\alpha$ , prolyl-4-hydroxylase, vascular endothelial growth factor and erythropoietin. The present study was undertaken to determine if human hepatoma cells when cultured for 72 h in the presence of serum under normoxia would maintain their cancerous phenotypic expression of certain hypoxia inducible genes. Our positive results affords an in vitro model system to test hypoxia inhibitors on the expression and the intracellular compartmentalization or the secretion of these hypoxia-inducible proteins. *J. Cell. Biochem.* 100: 1081–1085, 2007. © 2007 Wiley-Liss, Inc.

**Key words:** human hepatoma cells (Hep 2G);  $\beta$ -subunit of prolyl-4-hydroxylase; hypoxia-inducible factor-1 $\alpha$ ; erythropoietin; vascular endothelial growth factor

A hypoxic microenvironment is of prime importance as a regulator of the expression of heat shock protein (HSP) and angiogenic factor genes by heat shock transcription factor (HSF) and hypoxia-inducible factor (HIF-1). The expression of these hypoxic genes are thought to contribute to the malignant progression of tumor cells. HIF-1 is composed of an oxygen-sensitive  $\alpha$ -subunit (HIF-1  $\alpha$  or HIF-2  $\alpha$ ) and a constitutively expressed beta-subunit. During normoxia HIF-1  $\alpha$  is destabilized by the post-translational hydroxylation of Pro-564 and Pro-402 catalyzed by oxygen-sensitive dioxygenases termed prolyl hydroxylases [Marxsen et al., 2004]. Hypoxia stimulates the synthesis

of a cluster of enzymes via HIF-1 involved in post-translational modification of collagen polypeptide chains for the formation of the triple helix [Hofbauer et al., 2003]. HIF-1 $\alpha$  is a transcription factor that senses low oxygen availability and enhances the activation of the hypoxia-inducible genes involved in energy/ion metabolism, angiogenesis, and erythropoiesis [Percy et al., 1997; Semenza, 2003; Lee et al., 2004]. Under normoxia, HIF-1 $\alpha$  protein is known to be ubiquitinated by E3 ubiquitin ligase van hippel laudue (VHL) in various cell types [Huang et al., 1998; Lisztwan et al., 1999; Maxwell et al., 1999; Ohh et al., 2000]. Under hypoxia, VHL is unable to associate with and ubiquitinate HIF-1 $\alpha$ . HIF-1 $\alpha$  then escapes from degradation, translocates to the nucleus and activates the transcription of target genes. R59949, an inhibitor of diacylglycerol kinase enhances VHL binding to endogenous HIF- $\alpha$  and stimulates prolyl hydroxylase at both 21% O<sub>2</sub> as well as at 1% O<sub>2</sub> [Temes et al., 2005]. There is now an in vivo model to test the ability of agents on oxygen tension in particular organs.

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A mouse line has been created that expresses the oxygen-dependent degradation domain of HIF-1 $\alpha$  fused to the common firefly luciferase gene, under the control of a promoter that ensures organism-wide expression. In conditions of normoxia, the protein is quickly degraded, but in hypoxic situations, the protein is not degraded and administration of substrate illuminates the areas of expression [Schmid and Young, 2006]. The present study was carried out to determine if key proteins expressed in human hepatomas *in vivo* maintain their expression in *cell culture* under normoxic conditions.

## MATERIALS AND METHODS

### Cell Culture

Hep2G human hepatoma cells (ATCC #HB-8065) were obtained from the ATCC collection (Rockville, MD). The following materials were purchased from the designated companies: penicillin, streptomycin, L-glutamine, phosphate-buffered saline (PBS) (Gibco-BRL, Grand Island, NY); fetal bovine serum (FBS), (Hyclone Laboratories, Logan, UT); modified Eagle's media (MEM) (ATCC).

The Hep2G cells were maintained under air 5% CO<sub>2</sub> in ATCC complete growth medium, MEM with 2 mM L-glutamine and Eagle's BSS adjusted to contain 1.5 g/L sodium bicarbonate, 0.1 mM non-essential amino acids and 1.0 mM sodium pyruvate, 10% FBS, (v/v) and used at passage #38.

### Protein Extracts

At 8, 12, 24, 48, and 72 h following normotoxic exposure, the cells were collected by scraping from six well plates in cold PBS and centrifuged at 500g for 3 min. Nuclear and cytoplasmic fractions were then isolated using the NE-PER Nuclear and Cytoplasmic reagent kits (Pierce Biotechnology, Rockford, IL), according to the manufacturer's protocols. The protein concentrations of the extracts were determined using the Coomassie-Plus reagent (Pierce).

### Antibody Sources

Antibodies recognizing the following proteins were purchased: hypoxia inducible factor-1 $\alpha$ , (HIF- $\alpha$ ) 1:8,000 (cat#07-628, Upstate Biotech., Charlottesville, VA), vascular endothelial growth factor (VEGF), 1:8,000 (cat #sc-507, Santa Cruz Biotechnology, Santa Cruz, CA), erythropoietin (EPO), 1:8,000 (cat #sc1310,

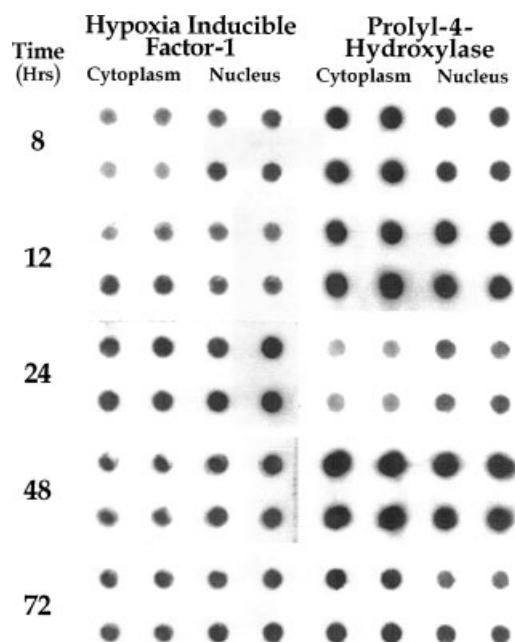
Santa Cruz Biotechnology) and human  $\beta$ -subunit of prolyl-4-hydroxylase (P-4-H) 1:8000 (cat # 63-164, ICN Biomedicals, Aurora, OH). These antibodies have all been shown to recognize their corresponding human antigens by their respective vendor. Five micrograms protein of nuclear or cytoplasmic extract was assayed per well. DAR (1:100,000) was used as secondary antibody for HIF-1 $\alpha$  while DAM was used as secondary antibody for P-4-H. For media 10  $\mu$ l/well was assayed. DAG at a 1:100,000 dilution was used as secondary antibody for VEGF and DAR at this same dilution was used as secondary antibody for EPO.

### Dot Blot Hybridization

Hep2G cells were plated in 96 well dishes and grown to confluency in DMEM containing 10% (v/v) FBS. Equal protein aliquots of the media, nuclear or cytoplasm were dot blotted onto a PVDF membrane (Millipore) using a dot blot template instrument to analyze a large number of samples. Proteins were transferred by electrophoresis onto a PVDF membrane which have any free protein binding sites blocked in a 5%(w/v) dried milk solution. The membrane is incubated with a primary antibody and after multiple washes, the membrane is incubated with peroxidase-conjugated secondary antibody, followed by another series of washes and the proteins are detected on X-ray film by BCA chemiluminescence (Pierce). To document equivalent protein loading an equal amount of protein of each fraction is dot blotted at each time of cell incubation. X-ray film is scanned and the protein dot density is used to quantify relative intracellular and extracellular protein levels and protein subcellular compartmentalization.

## RESULTS

Hypoxia-inducible gene expression occurs in spontaneous, chemical-induced and transplantable hepatocarcinomas under an acidic and hypoxic environment maintained by anaerobic glycolysis *in vivo*. As can be seen in Figure 1 HIF-1 and P-4-H are expressed in both the nuclear and cytoplasmic fractions of Hep2G human hepatoma cells up to 72 h in culture under normoxia. In addition at 8 h, the earliest time tested, under normoxia there is a decrease of P-4-H and an increase of HIF-1 in the nuclear fraction which represents a

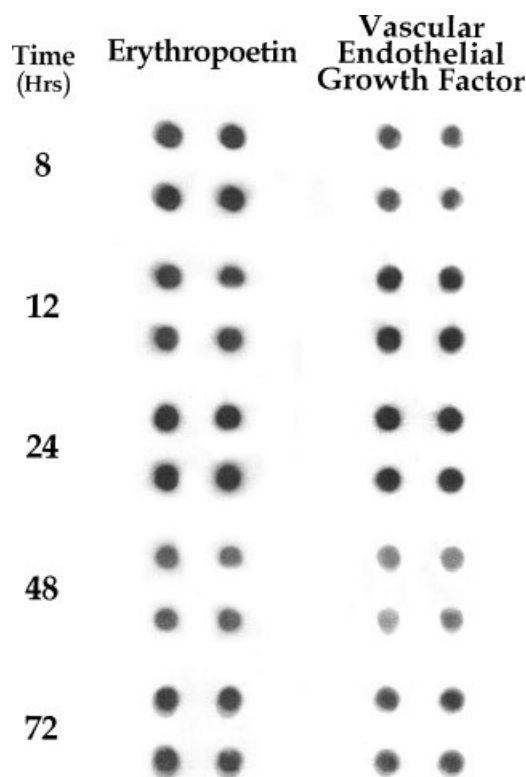


**Fig. 1.** Time course of hypoxia-inducible factor-1 and prolyl-4-hydroxylase in cultured Hep2G cells under normoxia in cytoplasmic (5  $\mu$ g protein) and nuclear (5  $\mu$ g protein) cellular fractions. All experiments were performed on duplicate cultures with duplicate dot blots of each subcellular fraction or media sample.

decreased translocation of this enzyme from the cytoplasm and an increased translocation of this latter factor from the cytoplasm. This enzyme is involved in the degradation of the HIF-1 through VHL binding and 26S proteasome degradation. This post-translational modification enzyme also is involved in collagen synthesis in the cytoplasm. The only other time point under normoxia that the Hep2G cells demonstrated a translocation of P-4-H from the cytoplasm to the nuclear fraction was at 24 h in culture with the HIF-1 remaining unchanged in both the cytoplasmic and nuclear fractions. As seen in Figure 2 no changes were observed for the hypoxia-inducible expressed gene phenotypes of VEGF and EPO secreted into the media of Hep2G carcinoma cells cultured up to 72 h under normoxia. However, both of these factors in the media of 48 h Hep2G cultures were significantly decreased, possibly due to turnover.

#### DISCUSSION

Tumors exist *in vivo* in an acidic, hypoxic microenvironment undergoing anaerobic glyco-



**Fig. 2.** Time course of media (10  $\mu$ l) vascular endothelial growth factor and erythropoietin of Hep2G cells cultured under normoxia.

lysis, angiogenesis with the HIF-1 induction of VEGF [Jung et al., 2005; Kim et al., 2005; Zhang et al., 2005] and EPO [Percy et al., 1997]. Collagen prolyl hydroxylases are required for proper collagen helix synthesis. These enzymes belong to the group of four oxoglutarate and iron-dependent oxygenases. P-4-H has been characterized as an  $\alpha_2\beta_2$  tetramer in which protein disulfide isomerase (PDI) is the  $\beta$  subunit isoforms (EC 5.3.4.1) with two different  $\alpha$  subunit (EC 1.14.11.2) isoforms. The enzymatically active  $\alpha$  subunit (64 kDa) of both the human and chick enzymes are glycosylated [Guzman et al., 1992]. PDI, the  $\beta$ -60 kDa subunit is a constituent of P-4-H which catalyses the co- and posttranslational hydroxylation of certain sequences specific collagen and HIF-1 $\alpha$  prolyl residues. The vertebrate P-4-H $\beta$  subunit is identical to the multifunctional enzyme PDI. Moreover, unassembled  $\alpha$ -subunits are maintained in an assembly competent inactive form by interaction with the molecular chaperone BiP [John and Bulleid, 1996].

PDI has two interrelated activities, that is, the ability to catalyze the formation, reduction,

and isomerization of disulfide bonds and the ability to bind to polypeptide chains [Freedman et al., 1994; Pirneskoski et al., 2001]. This latter activity enables PDI to function as a molecular chaperone and to assist in the folding of polypeptides. It has been suggested that PDI forms more permanent associations with specific proteins, for example, to become a subunit of P-4-H and microsomal triglyceride transfer protein. Inhibition of P-4-H activity can be obtained with a monoclonal antibody to the beta subunit of this enzyme [Kaska et al., 1987].

We recently published a comprehensive review on the relationship of tissue fibrosis to the final outcome of carcinogenesis in various tissues [Cutroneo et al., 2006]. Hypoxia is the microenvironment in which carcinomas thrive. Hypoxia also increases ProCOL1A1 mRNA [Falanga et al., 1993], COL1A1 transcription and collagen synthesis in normal fibroblasts [Falanga et al., 2002; Papakonstantinou et al., 2003].

In the present study a human hepatoma cell culture model of key proteins elevated in carcinomas maintained in an acidic hypoxic environment in vivo were shown to be phenotypically expressed in cells cultured under normoxia for at least 72 h. This affords an in vitro model to access the effects of inhibitors of HIF-1 synthesis and the subsequent effects on its cluster of hypoxia inducible proteins.

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