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Posttranslational modifications of collagens as targets of hypoxia and Hif-1 α in endochondral bone development

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Abstract

In recent years, it has been proposed that oxygen is not only an indispensable metabolic substrate, but also a regulatory signal, and that gradients of oxygenation turn on a specific genetic program. A crucial mediator of the adaptive response of cells to hypoxia is the transcription factor hypoxia-inducible factor 1 α (Hif-1 α). The fetal growth plate, which is an avascular structure of mesenchymal origin, has a unique out-in gradient of oxygenation. Hif-1 α is required for chondrogenesis *in vivo* by controlling a complex homeostatic response that allows chondrocytes to survive and differentiate in an hypoxic environment. Preliminary evidence suggests that regulation of posttranslational modifications of collagens could be an important component of this adaptive response.

Keywords

hypoxia; Hif-1 α ; VHL; chondrocytes; matrix; collagen

Introduction

Hypoxia is a relative decrease of oxygen (O₂) availability. The definition of “physiologically” normoxic conditions for either embryonic or adult cells varies significantly. Before the circulatory system is established, mammalian development proceeds in a relatively low O₂ environment of about 3%.¹ Moreover, studies that have utilized small-molecule hypoxia markers have shown the existence of specific regions of moderate to severe hypoxia in the developing embryos.² In the majority of normal adult tissues, oxygen (O₂) levels vary between 2% and 9% (as compared to ambient air, which contains 21% O₂).³ In contrast, O₂ concentrations in regions of the bone marrow, cartilage, kidney medulla, and thymus are below 1% O₂.³ Hypoxia is not only a critical factor in fetal development and differentiation, but also a pathophysiologic component of many human disorders, including cancer and ischemic diseases.⁴

Hypoxia-inducible factor-1 (Hif-1), a ubiquitously expressed transcription factor, is a major regulator of cellular adaptation to hypoxia.^{5–8} It is a heterodimeric DNA-binding complex that consists of two basic helix-loop-helix (bHLH) proteins of the PER/ARNT/SIM (PAS) subfamily, Hif-1 α and Hif-1 β . Hif-1 α and Hif-1 β mRNAs are ubiquitously expressed. Hif-1 α levels increase exponentially as O₂ levels drop below 5%. On the other hand, Hif-1 β (also known as aryl hydrocarbon nuclear translocator or ARNT) is non-oxygen-responsive.

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Conflicts of interest

The author declares no conflicts of interest.

Upon heterodimerization with Hif-1 β , the Hif-1 α /Hif-1 β complex binds to a specific sequence 5'-RCGTG-3' (where R denotes a purine residue) termed *hypoxia response elements* (HREs), and transactivates target genes containing HREs. Hif-1 α does not directly sense variations of O₂ tension; a class of 2-oxoglutarate-dependent and Fe²⁺-dependent dioxygenases are the O₂ sensors.⁹ Two types of O₂ sensors are involved in Hif-1 α action, prolyl-hydroxylase domain proteins (PHDs) and an asparaginyl hydroxylase, respectively. PHDs hydroxylate two prolyl residues (P402 and P564) in the HIF-1 α region, referred to as the O₂-dependent degradation domain (ODDD). This modification occurs in normoxic conditions and mediates the binding of the von Hippel-Lindau tumor suppressor protein (pVHL), which is an E3 ubiquitin ligase, to Hif-1 α . Hif-1 α is then marked with polyubiquitin chains and targeted for degradation by the proteasome. In well-oxygenated tissues, where O₂ tension is higher than 5%, Hif-1 α displays one of the shortest half-lives (< 5 min) among cellular proteins. Conversely, under hypoxic conditions, the activity of the PHDs is largely impaired and proline hydroxylation cannot occur. As a result, Hif-1 α protein accumulates and this initiates a multi-step pathway that includes nuclear translocation of Hif-1 α , dimerization with its partner Hif-1 β , recruitment of transcriptional co-activators, and binding to HREs within the promoters of hypoxia-responsive genes. The second type of O₂ sensor is an asparaginyl hydroxylase called factor-inhibiting Hif-1 (FIH-1). This enzyme hydroxylates an asparagine residue (N803) in the carboxy-terminal transcriptional activation domain (C-TAD) of Hif-1 α . This covalent modification blocks C-TAD interaction with transcriptional co-activators, such as p300 and CBP. Thus, the two O₂ sensors, PHD and FIH, by regulating the destruction and activity of Hif-1 α , respectively, ensure the repression of the Hif-1 pathway in well-oxygenated cells.

To date, more than 100 putative Hif-1 target genes have been identified.^{10–12} They are involved in a wide variety of biological processes including energy metabolism, angiogenesis, erythropoiesis, cell survival, apoptosis, redox, and pH regulation.¹¹ Mouse embryos lacking Hif-1 α exhibit multiple morphologic defects as early as embryonic day E8.5, and die *in utero* by E10.5.^{13–15} Many malignant cancers contain regions of severe hypoxia, resulting in high levels of Hif-1 α that drive tumor progression,^{5,8} and inhibition of Hif-1 α has been proposed as a potentially powerful approach.

Hif-1 α is necessary for chondrocyte survival and differentiation and regulates joint development

The fetal growth plate is a striking example of the critical and nonredundant role of the transcription factor Hif-1 α in survival and differentiation of hypoxic cells *in vivo*.

Skeletal development depends on two mechanisms, intramembranous and endochondral.¹⁶ In the first, mesenchymal cells develop directly into osteoblasts and form the flat bones of the skull. The second, accounting for the development of most other bones, involves a two-stage mechanism, whereby chondrocytes form a matrix template, the growth plate, which is then replaced by bone. During endochondral bone development, growth plate chondrocytes undergo well-ordered and controlled phases of cell proliferation, maturation, and death. Proliferative chondrocytes synthesize collagen type II and form a columnar layer. They then stop proliferating and differentiate into postmitotic hypertrophic cells. Hypertrophic chondrocytes express predominantly collagen type X and mineralize their surrounding matrix. Differentiation is followed by death of hypertrophic chondrocytes, followed by blood vessel invasion, and finally by replacement of the cartilaginous matrix with bone.

The cartilaginous matrix is formed by two components, the proteoglycans and the collagens. Proteoglycans are macromolecules containing a core protein with multiple attached polysaccharide chains.¹⁷ Because of their high content of charged polysaccharides,

proteoglycans are highly hydrated. In regard to collagens, there are three different types of collagens in the growth plate matrix, fibrillar, fibril-associated, and sheet-forming collagens.¹⁸ Fibrillar collagens comprise collagen type II and collagen type XI. Collagen type II is also found in the vitreous. In cartilage, collagen type II is produced by proliferating chondrocytes and by the upper hypertrophic chondrocytes. Fibril-associated collagens comprise collagen type IX, which, as collagen type II, is also found in the vitreous and binds proteoglycans. Sheet-forming collagens comprise collagen type X. Collagen type X is exclusively expressed by hypertrophic chondrocytes.

Cartilage is an avascular and hypoxic mesenchymal tissue,¹⁹ although it requires an angiogenic switch in order to be replaced by bone. The fetal growth plate contains a hypoxic central region.¹⁹ By using the marker of hypoxia EF5, the fetal chondrocytic growth plate has been shown to bind EF5, with no binding detected in surrounding soft tissues. The most hypoxic chondrocytes are in the round proliferative layer near the joint space, in the center of the columnar proliferative layer and in the upper portion of the hypertrophic zone.¹⁹ The EF5 findings document a gradient of oxygenation, from the proliferative to the hypertrophic zone, as well as from the outer to the inner region of the fetal growth plate. The high rate of O₂ diffusion to the mineralized hypertrophic layer may be the reason for the hypoxic condition of the early hypertrophic chondrocytes, even though they are located near the blood vessels of the primary spongiosa.¹⁹

Analysis of *in vivo* genetic models using the Cre-LoxP system has shown that Hif-1 α is necessary for chondrogenesis *in vivo* by turning on a complex homeostatic response that allows chondrocytes to survive and differentiate in a hypoxic environment.^{19,20}

Moreover, Hif-1 α protein is also particularly abundant in the highly hypoxic developing joints, possibly because the avascular perichondrium surrounding them is thickened.²⁰ Even after the joint space has formed, articular chondrocytes are significantly more hypoxic than the rest of the cartilage.²⁰ Lack of Hif-1 α in limb bud mesenchyme delays joint development, without altering thickening of the perichondrium.^{20,21} Thickening of the perichondrium therefore precedes joint formation and is likely to be critical for joint development. The delay of early chondrogenesis, secondary to the lack of Hif-1 α , may impair joint formation. However, because of the pronounced expression of Hif-1 α in the prospective joint, the delayed joint formation associated with loss of Hif-1 α may not be the only consequence of a delay in early chondrogenesis.

The role of Hif-1 α in cell differentiation is tissue-specific, because Hif-1 α maintains stem cells in an undifferentiated state^{3,22–25} and inhibits differentiation of mesenchymal cells into osteoblasts, adipocytes, and myocytes,^{3,26–28} yet stimulates differentiation of trophoblastic cells and dopaminergic neurons and chondrocytes.^{3,29–31}

Role of posttranslational modifications of collagens in mediating the survival and differentiation functions of Hif-1 α in growth plate chondrocytes

A variety of mechanisms can be invoked downstream of Hif-1 α as a survival and differentiation factor, including regulation of VEGF expression and modulation of metabolic pathways and autophagy.^{20,32} However, recent experimental evidence indicates that regulation of posttranslational modification of collagens, with hydroxylation of collagen prolines in particular, could be one of the modalities by which Hif-1 α regulates chondrocyte survival and differentiation. Prolyl-4-hydroxylases I and II (P4HaI and P4HaII) are the enzymes responsible for generating 4-hydroxyprolines in the collagens; these are essential for the formation of triple-helical collagens. P4Has have much lower K_m for O₂ than the

PHDs, which trigger Hif-1 α degradation (20 versus 250, respectively).³³ This indicates that P4Hs require a minimal amount of O₂ for proper function, that is, they still function enzymatically at low O₂ levels. The alpha subunits of P4HaI and P4HaII are targets of hypoxia in chondrocytes and other cell types in a Hif-1 α -dependent fashion.^{20,34} Proper accumulation of extracellular matrix is not only essential for organ development, but also promotes cell differentiation and survival through specific cell-matrix interactions.³⁵ Hif-1 α may thus operate as a survival and differentiation factor in chondrocytes, improving the efficiency of posttranslational modifications of collagen type II and, in so doing, promoting the formation of a proper extracellular matrix. A defect in posttranslational hydroxylation of collagens leads to a decrease in extracellular matrix and an increase in under-hydroxylated collagens. This in turn may trigger an unfolded protein response (UPR),³⁶ and may be a cause of the delayed chondrogenesis in mice that lack Hif-1 α in limb bud mesenchyme.

The positive effect of Hif-1 α on matrix accumulation in chondrocytes is consistent with the role of hypoxia in promoting fibrosis in pathologic conditions.³⁷ Moreover, it has been reported that lysyl oxidase, which is responsible for the formation of cross-links between collagen molecules, is induced by hypoxia and is essential for metastasis of highly malignant and hypoxic tumors.³⁸

Hypoxia and Hif-1 α may also modulate matrix formation by chondrocytes by upregulating expression of Sox9,^{21,39} a master regulator of chondro-genesis.⁴⁰ In mouse bone marrow stromal (ST2) cells, in particular, hypoxia brings about an increase in nuclear accumulation of Hif-1 α and Sox9 transcription.³⁹ Similar findings have been reported in limb bud micromass cultures,²¹ but not in primary chondrocytes or *ex vivo* metatarsal explants.²⁰ Notably, no evidence has been reported of a role of hypoxia in directly regulating expression of mRNAs encoding collagens in growth plate chondrocytes.²⁰

Summary

In this brief review, we have highlighted the role of hypoxia and Hif-1 α in regulating posttranslational modifications of collagens, and its importance in mediating the survival and differentiation functions of HIF-1 α in the developing cartilage and joints. It will be now important to investigate whether regulation of collagen P4Hs is the main mechanism adopted by this transcription factor to control matrix accumulation in the developing growth plate. To study regulation of posttranslational modifications of collagens by hypoxia and Hif-1 α in cartilage and joints may significantly expand our understanding of cellular adaptation to hypoxia as well as cartilage homeostasis.

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