Mini-Review

Hypoxic Regulation of Nucleus Pulposus Cell Survival

From Niche to Notch

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This minireview examines the role of hypoxia, and hypoxia inducible factors (HIF-1 and HIF-2), in regulating the metabolism, function, and fate of cells of the nucleus pulposus in the intervertebral disk. We focus on the mechanisms by which both these hypoxia-sensitive transcription factors influence energy metabolism, radical dismutation, and expression of survival proteins. In addition, we discuss how cells of the nucleus respond to a number of hypoxia-sensitive proteins, including galectin-3, Akt, and VEGF. Where applicable, these discussions are extended to include the impact of these molecules and hypoxia on degenerating resident cells in the intervertebral niche. Finally, because the notch signaling pathway is responsive to hypoxia, we speculate that in the intervertebral niche, notch proteins participate in the regulation of disk precursor cell proliferation and differentiation. We predict that knowledge of each of these interactive proteins within the disk niche could be used to enhance renewal and promote differentiation and function of cells of the nucleus pulposus. (Am J Pathol 2010, 176:1577–1583; DOI: 10.2353/ajpath.2010.090734)

The intervertebral disk is a complex structure that displays many of the characteristics of a diarthrodial joint in that it separates opposing cartilage-covered bones (vertebrae), permits a range of motions, and accommodates high biomechanical forces. While disks from the different anatomical regions of the spine vary in shape and volume, their architecture is similar. At the disk periphery,

the outer annulus fibrosus layer forms a ligamentous structure, composed of tightly packed parallel collagen type I fibrils that are inserted into contiguous superior and inferior vertebral bodies. The inner surface of the annulus fibrosus comprises a poorly organized fibrocartilage containing collagen type II fibrils. The annulus and the cartilaginous endplates enclose the nucleus pulposus, an aggrecan-rich gel-like tissue that is sparsely populated with cells (Figure 1A).

Cells of the nucleus pulposus are often mistakenly compared with chondrocytes, although they are distinct embryologically and exist in a unique microenvironment. The nucleus pulposus is derived from the notochord, whereas annulus fibrosus and endplate cartilage is sclerotomal in origin. Throughout this review, we refer to the cells of this notochord-derived tissue as nucleus pulposus cells.

The interaction between the semifluid nucleus pulposus and the tight molecular lattice of the annulus fibrosus provides the biomechanical properties necessary for spinal stability.¹ Disturbing this relationship by compromising the stability of the nucleus pulposus, the annulus fibrosus, or the endplate cartilage results in disk degeneration, a condition that can lead to excruciating pain and loss of function, and which often results in costly surgical interventions.¹ Because the degenerative process is chronic, the nucleus pulposus cells are required to function for long time periods under suboptimal microenvironmental conditions; the goal of this review is to consider those conditions that enhance nucleus pulposus cell sur-

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Figure 1. A: A cartoon showing the structure of the intervertebral disk. Tissue is avascular except for small capillaries in the outer one third of annulus fibrosus and in cartilaginous endplates. The nucleus pulposus, which is completely devoid of any blood supply, receives its nutrition and O_2 by diffusion through the endplates and is hypoxic. **B:** Functional activity of HIF target genes. Critical functions include energy metabolism, angiogenesis, cell survival, autophagy and apoptosis, matrix synthesis, proliferation, self-renewal and differentiation, radical dismutation, and pH regulation. Many of these functions are critical for survival and functioning of the nucleus pulposus cells in the avascular niche of the intervertebral disk. Hypoxia/HIF-sensitive proteins that are identified in the nucleus pulposus cells are shown in parentheses.

vival as well as factors that disregulate the disk microenvironment and promote degenerative disk disease.

Is the Intervertebral Niche Hypoxic?

Although the clinical outcomes of degenerative disk disease are well documented, biological events that regulate nucleus pulposus cell survival are not understood. One overriding aspect of disk cell biology is that cells of the nucleus pulposus and cells residing in the inner annulus are removed from the blood supply. For example, blood vessels originating in the vertebral body traverse the superficial region of the endplates; none of these vessels infiltrate the nucleus pulposus. With respect to the annulus, Gruber et al pointed out that this tissue is avascular except for small discrete capillary beds in the dorsal and ventral surfaces—in no case, does the annulus vasculature enter the nucleus pulposus.^{2–4} Modeling studies by Urban and colleagues indicate that the pO_2 within the disk is low⁵; related to this observation, cells in the transitional zone between the inner annulus and the nucleus as well as in the nucleus itself evidence a robust hypoxic signal.⁶ For these and other reasons, it

would not be unreasonable to predict that the nucleus pulposus cells reside in a hypoxic tissue niche.⁷

Herein, we use the term niche to describe the confines of the nucleus, bounded laterally and medially by the annulus and superiorly and inferiorly by the end plate cartilage. Although the concept of a niche was originally directed at anatomical structures, more recently the term has been used to describe interactions between communities of cells that are in close proximity to each other. For example, within the bone marrow niche, stem cell commitment to a particular lineage is dependent on local micro-environmental conditions that regulate the interactions between resident hematopoietic as well as stromal cells.

If the concept of a regulatory niche, composed of a number of cell types responsive to local microenvironmental conditions, is valid, then this begs the question: is cell survival in the hypoxic niche hypoxia inducible factor (HIF)-dependent? Thanks to the brilliant studies of Semenza and colleagues, it is now recognized that the key molecule regulating energy metabolism and survival activity is HIF-1.⁸ This molecule is a member of the basic helix-loop-helix (bHLH)–PER-ARNT-SIM (PAS) family of proteins and composed of a constitutively expressed β subunit and an α subunit. The latter subunit is stable under hypoxic conditions but is rapidly degraded in normoxia.⁹ Transactivation of HIF-1 target genes involves dimerization of the two subunits and binding to an enhancer, the hypoxia-response element in target genes. HIF-1 serves as a key transcription factor that regulates the expression of enzymes concerned with glycolysis, the activity of the TCA cycle and oxidative phosphorylation.^{8,10,11} Additional target genes include those required for survival, apoptosis, autophagy, and matrix synthesis.^{12–14} Details of these relationships are shown schematically in Figure 1B. It should be added that other isoforms of HIF exist, the most important being HIF-2 α . Recent evidence suggests that HIF-1 α and HIF-2 α are not redundant, and that the relative importance of each of the homologues, in response to hypoxia, varies among different cell types.¹⁵ For example, unlike HIF-1, HIF-2 regulates expression of a number of unique genes including superoxide dismutase 2 (SOD2), catalase, frataxin, and cited2.^{16–18} In addition to these genes, the Sox family of transcription factors that are essential for the development and function of the nucleus pulposus are hypoxia- and HIF-sensitive.^{19–22} Lafont et al showed that HIF-2 but not HIF-1 regulated the expression of Sox9 and the phenotype of primary human chondrocytes.²⁰ Similarly, expression of Sox9 and Sox5 and Sox6 is hypoxia- (5% O_2) and HIF-2-sensitive during in chondrogenic differentiation of stem cells derived from infrapatellar fat pads of osteoarthritic patients.²¹ In contrast, using marrow mesenchymal stem cells, Kanichai et al showed involvement of HIF-1 α in regulating Sox9 expression during chondrogenesis under hypoxia (2% O_2).²² However, the relationship between Sox proteins and HIF in the hypoxic niche of the disk is not as yet known.

To return to the question raised above concerning the importance of the HIF system, a considerable number of reports now clearly show that there is a robust HIF re-

Figure 2. O_2 -dependent metabolism of HIF-1 α . **A:** Normoxic regulation of HIF-1 α . Under normoxic conditions, the α subunit of HIF-1 is hydroxylated at specific proline residues by prolyl hydroxylases (PHD1-3) that require Fe^{2+} , 2-oxoglutarate (2-OG), and ascorbate. Of these, PHD1 plays a major role in HIF-1 α regulation. Once hydroxylated, HIF-1 α is recognized and bound by von Hippel Lindau, an E3 ubiquitin ligase; it is subsequently ubiquitinated and rapidly degraded through the 26S proteasomal pathway. **B:** Hypoxic regulation of HIF-1 α . Under hypoxic conditions, HIF-1 α is stabilized, translocates to the nucleus, and dimerizes with a constitutively expressed HIF-1 β subunit. The dimer associates with several coactivators, including p300/CBP and the DNA polymerase II (Pol II) complex. This protein complex then binds to an enhancer, the Hypoxia-Response Element (HRE), in HIF-1 target genes to initiate gene transcription. Of note, in nucleus pulposus cells, HIF-1 α is stabilized even in normoxia.

sponse by cells of the nucleus pulposus. The response is evident across species, it is seen *in vivo* and *in vitro*, and more importantly, HIF-1 α activity is unresponsive to the oxemic state of the tissue.^{7,23,24} Accordingly, when compared with most other tissues, there are substantive underlying differences in the HIF status and reactivity of disk cells: HIF-1 α expression and activity is always 'on.' This unusual response suggests that stabilization of HIF-1 α in cells of the nucleus pulposus ensures that transcriptional activity is a major determinant of cell function. The second HIF homologue, HIF-2 α , is robustly expressed by nucleus pulposus cells. Like HIF-1 α , steady-state protein levels are similar in both hypoxia and normoxia, suggesting that it too is constitutively expressed.²⁵

Before leaving this topic, it is important to comment on the possible mechanism of stabilization of HIF-1 α in nucleus pulposus cells. A glance at Figure 2 indicates that stabilization can be achieved at a number of different levels. For example, suppression of the von Hippel Lindau or low activity of one or more of the O_2 -sensing prolyl hydroxylase (PHD) enzymes would be expected to decrease HIF-1 α turnover.²⁶ In the intervertebral disk, nucleus pulposus cells express PHD1 and PHD3 whereas PHD2 is expressed at a low level (unpublished). Inhibition of PHD function by pyruvate and TCA cycle intermediates isocitrate, succinate, fumarate, and oxaloacetate can modulate HIF expression and activity.²⁷ Besides prolyl hydroxylases and von Hippel Lindau, HIF-1 α stabilization is achieved by controlling expression of important constituents of the multiprotein complex such as osteosarcoma-9 that regulates HIF-1 α levels in an O_2 -dependent

manner.²⁸ Whether normoxic stabilization of HIF-1 α and HIF-2 α in the disk is an adaptive response to an imposed metabolic need, related to the unique embryonic origins of the disk, is currently unknown. However, it is important to note that because disks are hypoxic *in vivo*, stabilization of HIF-1/2 expression would serve to maintain cell metabolism and survival activities when disk integrity is breached during disk herniation²⁹ or at an early stage of degeneration.¹

Does HIF-1 Maintain Energy Conserving Activities in the Disc?

Earlier classical biochemical studies have shown that when the pO_2 is low, there is almost complete reliance on glycolysis to generate ATP and reducing equivalents. As indicated in the previous section, one of the consequences of low oxygen tension in the nucleus pulposus is the reliance on glycolysis for energy generation.^{24,30} Glycolysis may be viewed as a relatively inefficient process: it generates 2 moles of ATP/mole glucose; in contrast, mitochondrial metabolism is slow, but it creates about 30 moles of ATP/mole glucose. In the trade-off between rate and yield, the glycolytic pathway generates a small number of ATP molecules at a very fast rate and maintains the reducing status of the cell. In this way, glycolysis provides sufficient energy for both housekeeping functions and for protein synthesis.

One of the logical outcomes of stabilization of HIF-1 α is the robust expression of glucose transporters and enzymes required for anaerobic glycolysis. When the ex-

pression of three target genes (glucose transporter [GLUT]–1 and –3 and GAPDH) at 2% and 21% O₂ were evaluated, it was found that the activities are comparable and remain constant over time.²⁴ Although these genes were not responsive to the oxemic state of the culture, we have observed a small induction in enolase-1 and phosphofructokinase 2 (PFKFB) promoter activities. This result is surprising as the later protein is regarded as the glycolytic ‘pacemaker’; however, because this intermediary step is sensitive to a number of hormones, and metabolic intermediates, it is more than likely that induction is in response to other regulatory factors. Nevertheless, the muted response does not detract from the conclusion that the glycolytic flux in disk cells, even in normoxia, is high.

In normoxia, basal concentrations of ATP in nucleus pulposus cells are between 20 and 25 nmol/L/mg protein.²⁴ These values are comparable with levels reported for articular chondrocytes, another cell type that uses glycolysis to generate energy.³¹ In the presence of 2-deoxyglucose, a potent inhibitor of glycolysis, ATP generation is suppressed by almost 80%.²⁴ The sensitivity of the cells to this inhibitor emphasizes the reliance on glycolysis for energy generation. Based on this observation, it is likely that the oxemic stability of HIF-1 α in nucleus pulposus cells is optimal for survival in an environment where there are frequent shifts in vascular supply and O₂ delivery; in the intervertebral disk, these shifts may reflect minute to minute or day/night variations in biomechanical forces applied to the spinal units.

Although glycolysis is clearly the major ATP-generating pathway, the possibility exists that some high energy intermediates may be produced through mitochondrial oxidative phosphorylation. However, current studies indicate that inhibitors of mitochondrial function do not influence ATP production, nor nucleus pulposus cell viability.²⁴ As for a role, if any, for mitochondria, little is known. Gan et al have reported that although nucleus pulposus cells contain mitochondria with normal architecture, the total number of organelles per cell is low.³² Nevertheless, nucleus pulposus cells can perform mitochondrial oxidative metabolism: thus, they oxidize fatty acid and generate ATP.²⁴ Based on all of these studies, there is strong support for the notion that although glucose and anaerobic glycolysis represent the major fuel and pathway for energy generation, respectively, mitochondria in the nucleus pulposus are functional; they retain the capacity to metabolize fatty acids through mitochondrial oxidative metabolism.

The conclusion that disk cell energy metabolism is dependent on glycolysis fits well with current observations concerning the regulatory functions of HIF-1. It is known that HIF-1 plays a major role in directing the interplay between glycolysis and oxidative phosphorylation.¹⁰ HIF-1 inhibits mitochondrial function by *trans*-activating the gene encoding pyruvate dehydrogenase kinase 1. Because this protein suppresses pyruvate dehydrogenase, pyruvate cannot be converted into acetyl-CoA, and as a result the TCA cycle is blocked.¹⁰ More recently, Fukuda et al showed that HIF-1 reciprocally regulates mitochondrial cytochrome c oxidase (COX)–4 subunit

expression by activating transcription of the genes encoding COX4–2 and a protease that is required for COX4–1 degradation.¹¹ Thus, HIF regulates not just the entry of reducing equivalents into the mitochondria, but also oxidative phosphorylation. Based on these observations, it is concluded that although mitochondrial function is retained by cells of the intervertebral disk, it is reasonable to assume that normoxic expression of HIF-1 α by nucleus pulposus cells serves to suppress oxidative phosphorylation and promote glycolytic ATP generation. More than likely, nucleus pulposus mitochondria are required for nonenergy related metabolic functions, while oxidative phosphorylation is used to a very minor degree.

Does Hypoxia Promote Cell Survival in the Intervertebral Disc?

If the premise is correct that the HIF signaling network serves to promote nucleus pulposus function, then the cells should be adapted to survive and grow in a hypoxic environment. To test this assumption, experiments have been performed in which nucleus pulposus cells were treated with low levels of common apoptogens and survival measured.^{33,34} Notably, when the pO₂ was below 5%, there was maximum disk cell survival. Studies with other hypoxia-adapted cells (chondrocytes) showed that when HIF-1 α is silenced, viability is maintained in the face of an O₂ challenge.³⁵ Hence, some effects of hypoxia on nucleus pulposus survival are probably mediated by other signaling molecules in an HIF-1-independent fashion. The latter observation raises the question, which signaling pathways are up-regulated in hypoxia? Work from a number of labs indicates that a variety of hypoxia sensitive proteins exist, including vascular endothelial growth factor (VEGF), galectin-3, and Akt/PI3K. We have found that expression levels of phospho Akt in nucleus pulposus cells is high in hypoxia and when serum starved confers resistance to apoptosis.³³ Activation of this protein is of considerable interest as it has been shown to modulate apoptosis by inactivating (phosphorylating) Bad and caspase-9 and modulating the transcription of proapoptotic transcription factors.³⁶ Relevant to nucleus pulposus cells, activation of PI3K/Akt signaling has been shown to regulate HIF-1 α protein levels in other cell types.²² Like Akt, extracellular signal-regulated kinase (ERK)1/2 is induced in hypoxic nucleus pulposus cells. Because activation of ERK has been linked to survival, possibly by regulating nitric oxide synthase and caspase activities, the possibility exists that activation of ERK in concert with Akt serves to maintain the viability of the disk cells at a low pO₂.^{33,34}

Like HIF-1, one of the critical functions of Akt is regulation of glucose metabolism. It is thought that Akt may promote cell survival by maintaining GLUT-1 transcription under conditions of growth factor withdrawal (serum-starvation).³⁷ Indeed, the high level of expression of GLUT-1 protein by nucleus pulposus cells *in vivo*^{7,38} indicates that this tissue may adapt to its hypoxic environment by increasing glucose uptake. This activity would serve to promote and enhance glycolysis, thereby preventing

ischemia-induced injury. Taken together, these studies suggest that the PI3K-Akt and ERK signaling pathways in conjunction with HIF-1 provide a mechanism by which nucleus pulposus cells remain viable and maintain their specialized physiological function, despite environmental limitations in O_2 and even possibly changes in nutrient availability.

Two proteins that have been linked to nucleus pulposus survival in hypoxia are VEGF-A and galectin-3. In disk cells, it has been reported that HIF-1 regulates galectin-3 expression.³⁹ From a functional perspective, by forming complexes with integrins, externalized galectin-3 influences cell adhesion and spreading.⁴⁰ Accordingly, galectin-3 is most likely involved with matrix stability and in concert with HIF-1 provides the discal cells with both a mechanico-transduction as well as a survival function. Other studies have shown that galectin-3 regulates survival by suppressing signaling through the TNF family of proteins.⁴¹ This finding is particularly pertinent to disk disease, as $TNF\alpha$ together with other cytokines are known to play a major role in the etiology, as well as progression, of the degenerative state. Based on these findings, it is possible that in the hypoxic intervertebral disk, the robust expression of HIF-1 α serves to maintain galectin-3 levels, which then serve to promote cell survival and disk function.³⁹

With respect to VEGF, not surprisingly, levels of this protein are high in herniated disks or in degenerative disks where there is evidence of neovascularization.⁴² In the normative state, because the disk environment is avascular, it would be reasonable to assume that VEGF expression is low. However, this is not the case as there is robust expression of this protein and its receptor in the nucleus pulposus.^{25,43} This level of expression is likely related to HIF-2, which preferentially upregulates VEGF-A expression and promoter activity.²⁵ This observation leads to the question of what is the function of VEGF in the disk? Clearly, it cannot serve to promote angiogenesis as this activity would promote vascularization and compromise disk function. There is some information to indicate that VEGF supports cell survival.⁴⁴ Indeed, in a recent study, Fujita et al⁴³ confirmed that VEGF and its receptors are expressed by nucleus cells in hypoxia and showed that this protein promoted nucleus pulposus survival. Thus, from a functional viewpoint, VEGF could serve as to maintain nucleus pulposus viability in the face of shifts in environmental pO_2 .

As indicated earlier, there is growing interest in the second HIF homologue, HIF-2 α . This protein is robustly expressed by nucleus pulposus cells. With respect to functional activities, unlike most other tissues, hypoxia failed to increase the transcriptional activities of SOD2 and frataxin, two common HIF-2 target genes concerned with radical dismutation.^{16,17} This finding could explain why this tissue is susceptible to radical attack associated with annular lesions or nucleus herniation. Notably, there is evidence to indicate that kyphosis, scoliosis, and radiculopathies are linked to defective radical dismutation, whereas Friedreich's Ataxia is now known to be attributable to low frataxin levels and loss of antioxidant defenses.^{45,46} It would be important to know whether

these conditions are also linked to the inability of vertebral tissues to mount a robust HIF-2-dependent scavenging response.

It is important to comment that both HIF-1 and HIF-2 are involved in survival of endplate chondrocytes by activation of the autophagic pathway.^{47,48} The importance of this pathway for removal of misfolded proteins and damaged organelles has been emphasized by a number of workers, and its role in directing the maturation of connective tissue cells has been discussed by Srinivas and his colleagues.⁴⁸ Noteworthy, although autophagy is viewed as a survival pathway, there is little doubt that continued macromolecular breakdown, while serving as a source of nutrients and energy for the stressed cell, inevitably leads to increased susceptibility to apoptosis (Type II apoptosis). Hence, HIF activity and ultimately HIF targets serve as key proteins that straddle both the apoptotic and survival pathways.

A very recent study by Bohensky et al pointed out that HIF-2 was also involved in regulating survival by modulating autophagy.⁴⁷ HIF-2 was expressed abundantly by cells in human and murine articular cartilage, hypertrophic cartilage, and in the endplate cartilage. When HIF-2 α was suppressed, ROS generation was elevated, and there was a decrease in the activity of the ROS dismutating enzymes catalase and superoxide dismutase. Suppression of HIF-2 α was associated with decreased Akt-1, reduced Bcl-x(L) expression, and a robust autophagic response, even under nutrient-replete conditions.⁴⁷ Relevant to disk disease, it is generally agreed that the degenerative state is exacerbated by decrease in permeability of the endplate cartilage and the concomitant reduction in nutrient availability. It is possible that under these nutritionally challenging conditions, increased HIF-1/-2 expression may serve to maintain nucleus pulposus survival by promoting the induction of autophagy.

Some Final Comments: Does HIF Promote Nucleus Pulposus Renewal through the Notch Signaling Pathway?

In this review, we have drawn attention to the critical role of the tissue pO_2 on the function and survival of cells of the intervertebral disk. We have focused on the mechanisms by which the hypoxia sensitive transcription factors HIF-1 and -2 influence energy metabolism and expression of survival proteins. In addition, we have discussed how cells of the nucleus respond to hypoxia-sensitive proteins, galectin-3, Akt, and VEGF. Where applicable, we have extended these discussions to include the impact of these molecules and hypoxia on degenerating resident cells in the intervertebral niche; not discussed is the effect of hypoxia on cell renewal in the disk. Although this topic is beyond the scope of the review, it should be stated that in concert with most connective tissues, cell turnover within the niche is slow. Moreover, like most of these tissues, progenitor cells are present in the disk that can differentiate along the mesengenic pathway to re-

place resident cells.⁴⁹ Thus, tissue renewal in the intervertebral disk is dependent on the ability of progenitor cells to commit to the nucleus pulposus lineage and undergo terminal differentiation.

The notch signaling pathway is central to these progenitor activities and pertinent to the ideas discussed earlier, the notch signaling pathway is responsive to hypoxia. Interaction between HIF-1 α and the intracellular domain of the notch protein inhibits differentiation of myogenic and neural precursor cells.⁵⁰ In skeletal tissues, disruption of notch signaling markedly increases trabecular bone mass: with aging, the mice become osteopenic due to a sharp reduction in mesenchymal progenitor populations.^{51,52} Hypoxia also increases the expression of known notch target genes such as Hes1 and Hey1.⁵⁰ Accordingly, in the nucleus, HIF-1 α may directly interact with the notch intracellular domain and direct cell fate. Based on what is known of cell replacement in other tissues, this HIF-1-regulated pathway is a critical component of cell renewal and replacement.

From a disease viewpoint, an oxemic shift, possibly mediated by alterations in the vascular supply to the endplate cartilage or even the annulus fibrosus would be expected to lead to a failure in progenitor cell activation and a decrease in the number of differentiated cells. In turn, this would lead to decrements in function and enhancement of the effect of agents that are known to promote disk degeneration. From a therapeutic viewpoint, it should be possible to modulate the niche environment to enhance renewal and promote differentiation of precursors into functional cells of the nucleus or the annulus. Accordingly, rather than relying on surgical and other interventional strategies, which may themselves damage the disk or cause infection, it should be possible to promote tissue repair by manipulating oxemic conditions within the niche, or use proteins of the notch signaling pathway to reactivate the endogenous progenitor cells in the annulus fibrosus or nucleus pulposus. Restoration of disk cell function and prevention of degeneration remains the ultimate goal of current intervertebral disk research.

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