Possible Role of PHD Inhibitors as Hypoxia-Mimicking Agents in the Maintenance of Neural Stem Cells’ Self-Renewal Properties

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Hypoxia is the most critical factor for maintaining stemness. During embryonic development, neural stem cells (NSCs) reside in hypoxic niches, and different levels of oxygen pressure and time of hypoxia exposure play important roles in the development of NSCs. Such hypoxic niches exist in adult brain tissue, where the neural precursors originate. Hypoxia-inducible factors (HIFs) are key transcription heterodimers consisting of regulatory α-subunits (HIF-1α, HIF-2α, HIF-3α) and a constitutive β-subunit (HIF-β). Regulation of downstream targets determines the fate of NSCs. In turn, the stability of HIFs-α is regulated by prolyl hydroxylases (PHDs), whose activity is principally modulated by PHD substrates like oxygen ($O_2$), α-ketoglutarate ($α$-KG), and the cofactors ascorbate (ASC) and ferrous iron ($Fe^{2+}$). It follows that the transcriptional activity of HIFs is actually determined by the contents of $O_2$, $α$-KG, ASC, and $Fe^{2+}$. In normoxia, HIFs-α are rapidly degraded via the ubiquitin-proteasome pathway, in which PHDs, activated by $O_2$, lead to hydroxylation of HIFs-α at residues 402 and 564, followed by recognition by the tumor suppressor protein von Hippel–Lindau (pVHL) as an E3 ligase and ubiquitin labeling. Conversely, in hypoxia, the activity of PHDs is inhibited by low $O_2$ levels and HIFs-α can thus be stabilized. Hence, suppression of PHD activity in normoxic conditions, mimicking the effect of hypoxia, might be beneficial for preserving the stemness of NSCs, and it is clinically relevant as a therapeutic approach for enhancing the number of NSCs in vitro and for cerebral ischemia injury in vivo. This study will review the putative role of PHD inhibitors on the self-renewal of NSCs.

Keywords: PHDs, inhibitors, NSCs, HIFs, hypoxia

INTRODUCTION

Neural stem cells (NSCs)/neural progenitor cells (NPCs) possess lifelong self-renewal abilities and can differentiate into neurons as well as glial cells such as astrocytes and oligodendrocytes in certain induced conditions (Gage, 2000). Neurogenesis in the adult brain remains controversial. The first evidence of adult hippocampal neurogenesis in the mammalian system was reported 50 years ago (Altman, 1962). However, current opinion is that neurogenesis in the hippocampus decreases rapidly during the first years of life and that neurogenesis in the dentate gyrus does not continue, or is extremely rare in adult humans (Sorrells et al., 2018). In contrast, another view is that human hippocampal neurogenesis...
IhNSCs, hNSCs were unable to proliferate in 1% O\(_2\) (IhNSCs) and raised the rate of apoptosis. In comparison with 1 or 21% O\(_2\), 5% O\(_2\) is the optimal condition for the proliferation of NSCs in normoxia. As mentioned above, in the nervous system, adult NSCs exist in quiescence in limited number (or are even completely absent). Neuronal transplants from NSCs cultured in vitro are needed for the treatment of neurodegenerative diseases such as Parkinson’s disease and Alzheimer’s disease as well as conditions like stroke (Goldman, 2016).

In this review, we present recent findings on the roles of hypoxia, HIF transcription factors and PHD inhibitors in NSC proliferation, focusing on the potential application of PHD inhibitors to mimic the effects of hypoxia.

**NSC SELF-RENEWAL/PROLIFERATION UNDER HYPOXIC CONDITIONS**

Neural stem cells have the ability to proliferate and differentiate into astrocytes, oligodendrocytes, and neurons (Gage, 2000). Previous studies have shown that in adult mice the main NSC niches are located in the hippocampus and the subventricular zone (SVZ), where they promote stem cell proliferation at low oxygen concentrations (1–5% O\(_2\)) (Mohyeldin et al., 2010). Under anoxic conditions, mitochondrial respiration is inhibited and cellular energy is produced by anaerobic glycolysis, which provides insufficient energy to fully support cell proliferation (Papandreou et al., 2006). Indeed, mild hypoxia (2.5–5% O\(_2\)) is the optimal condition for the proliferation of NSCs in comparison with 1 or 21% O\(_2\). Santilli et al. (2010) have recently shown that in marked contrast to 2.5–5% O\(_2\), 1% O\(_2\) decreased the proliferation of immortalized human NSCs (IhNSCs) and raised the rate of apoptosis. In comparison with IhNSCs, hNSCs were unable to proliferate in 1% O\(_2\), dying after a few passages, presumably because of cell cycle arrest and inhibition of transcriptional activity (Koshiji et al., 2004; Kaidi et al., 2007; Zhang et al., 2007). Furthermore, Pistollato et al. (2007) found that neural precursor proliferation in the human postnatal brain is enhanced in hypoxic conditions (5% O\(_2\)), while raising oxygen tension to 20% depletes precursors and promotes astrocyte differentiation. Hypoxia-expanded precursors generated 17-fold more oligodendrocytes and when these precursors were expanded in hypoxia and then differentiated in normoxia, oligodendrocyte maturation was further enhanced by 2.5-fold (Pistollato et al., 2007). Hypoxia (5% O\(_2\)) also reduces apoptosis while promoting the proliferation of NSCs, and WNT/β-catenin may be involved in the regulation of NSC proliferation (Cui et al., 2011). Under hypoxia (1.7% O\(_2\)), nuclear orphan receptor TLX acts as a mediator for the proliferation and pluripotency of neural progenitors and is recruited to the Oct-3/4 (Octamer-binding transcription factor 3/4) proximal promoter, enhancing gene transcription and promoting progenitor proliferation (Chavali et al., 2011). Hypoxia/reoxygenation (H/R) is another method to stimulate NPC proliferation, via activation of the MEK (MAP kinase-ERK) and the PI3K (phosphoinositide-3-kinase)/AKT signaling pathways through a PKC (protein kinase C)-dependent mechanism. These signals were associated with proliferation of NPCs (Sung et al., 2007).

In addition to the effects on NSC proliferation or self-renewal capacity, hypoxia also increases their survival ability after transplantation into animals with intracerebral hemorrhage. Wakai et al. (2016) have demonstrated that mild hypoxia (5% O\(_2\), 24 h) enhanced NSC proliferation, upregulated p-AKT via HIF-1α and increased vascular endothelial growth factor (VEGF) production around the wound after transplantation, which favors survival of transplanted cells. Another study indicates that hypoxia regulates the expression of functional metabotropic glutamate receptor (mGluR) in proliferating NSCs and the dynamic expression of mGluRs induced by hypoxia may be one of the mechanisms of hypoxia-stimulated NSC activation (Chen et al., 2016). Interestingly, it has been reported that hypoxic conditioned media from rat cerebral cortical cells promotes NSC proliferation and differentiation into a high percentage of neurons, and these processes may be promoted through PI3K/AKT pathways (Cai et al., 2014).

Hypoxia also has a similar effect on neurogenesis in the animal brain. Our previous study showed that intermittent hypoxia (IH) increased NSC proliferation, newborn neuron survival and migration, spine morphogenesis in the dentate gyrus of the hippocampus, and neurogenesis in the olfactory bulb (Zhu et al., 2005; Zhao et al., 2008). Further evidence suggested that NOTCH1 signaling was activated following IH, which was required for hypoxia-induced neurogenesis (Zhang K. et al., 2014). Ross et al. (2012) reported another hypoxic approach to NSC expansion and neuronal differentiation. An acute (1 h) in vivo exposure to intermittent hypoxia (AIH) was conducted using neonatal C57BL/6 mice, and the isolated NPCs from SVZ tissues grew in neurospheres or adherent monolayers. AIH stimulated the proliferation of NPCs. Therefore, in vivo AIH exposure can enhance the viability of SVZ-derived NPC cultures in vitro (Ross et al., 2012). Altogether, both in vivo
and in vitro hypoxia stimulation can promote NSC proliferation and a capacity for self-renewal. Since hypoxia has a wide range of effects, revealing the mechanism by which it promotes the proliferation of NSCs will contribute to targeting NSC proliferation in vitro and in vivo.

**HIF-1 TRANSCRIPTIONAL FACTOR UNDER HYPOXIA**

HIF-1α is a regulatory subunit of the HIF-1 transcriptional regulator. The constitutive subunit HIF-1β is present in excess and heterodimerizes with HIF-1α or other bHLH-PAS (basic helix-loop-helix–PER–ARNT–SIM) proteins. Therefore, HIF-1α protein levels determine HIF-1 transcriptional activity (Semenza, 2012). In addition to HIF-1α, HIF-2α was subsequently discovered, which also binds to the common HIF-1β subunit and forms the HIF-2 heterodimer (Tian et al., 1997). While the functions of HIF-1 and HIF-2 are similar, their distribution is different, with HIF-1α being ubiquitously expressed whereas HIF-2α is restricted to certain cell populations and mediates distinct functions (Prabhakar and Semenza, 2012). HIF-3α was originally considered as an inhibitor of HIF-1 via sequestration of HIF-1β. However, some findings indicate that HIF-3 also functions as a transcriptional activator in certain contexts such as in the hypoxic response in zebrafish embryos (Zhang P. et al., 2014). In this article, we focus on the regulation of HIF-1 because it is the most widely studied and represented. In normoxic conditions, prolyl residues 402 and 564 in the HIF-1α oxygen-dependent degradation domain (ODD) are hydroxylated by HIF prolyl hydroxylases (PHDs) (Huang et al., 1998; Semenza, 2001). Subsequently, the hydroxylated HIF-1α is recognized by von Hippel-Lindau protein (VHL), which is responsible for the recruitment of an ubiquitin ligase to target HIF-1α for proteasomal degradation (Ivan et al., 2001). In hypoxic conditions, PHDs are inactivated and the degradation of HIF-1α is blocked. Stabilized HIF-1α constitutes a heterodimer with HIF-1β and directly binds hypoxia-responsive elements (HREs) in the promoter of downstream genes, which are thereby activated (Semenza, 2007; Simon and Keith, 2008). HIF-1α activates the transcription of VEGF and erythropoietin (EPO), which encode angiogenic cytokines or growth factors, leading to angiogenesis and erythropoiesis, and so tissue perfusion improves and O2 delivery increases in normal tissues (Semenza, 2013). Another HIF target gene (EPO) also promotes proliferation of NSCs by modulating NF-kB (nuclear factor-kappa B) signaling (Shingo et al., 2001). Additionally, WNT/β-catenin signaling is implicated in the effects of HIF-1α on NSC proliferation. HIF-1α deletion leads to impaired WNT/β-catenin activity in the hippocampus, with impaired NSC proliferation (Mazumdar et al., 2010; Braunschweig et al., 2015; Hubbi and Semenza, 2015). Of note, our latest unpublished data show that autophagy induced by HIF-1 mediates NSC proliferation. Together, the above studies indicate that HIF promotes NSC self-renewal or proliferation through different signaling pathways. A number of hypoxic conditions are conducive to the self-renewal capacity of NSCs via the HIF-1 pathway. On the other hand, hypoxia does not specifically and exclusively affect NSC proliferation, due to the number of its downstream effectors, apart from the HIF-1 pathway. Furthermore, HIF-1α is unstable under normoxic conditions, and as such is easily degraded by the proteasome system (Prabhakar and Semenza, 2012).

**HIF 1 IS ASSOCIATED WITH SELF-RENEWAL CAPACITY OF NSCs**

Hypoxic preconditioning or exposure to sublethal levels of hypoxia enhanced NSC proliferation (Francis and Wei, 2010; Sart et al., 2014). Hypoxic preconditioning (5% O2, 24 h) may increase NSC proliferation by acting on the HIF-1α/AKT pathway. Moreover, VEGF as a downstream target of HIF-1α is released in response to hypoxia and promotes cell survival. It is critical for post-stroke cerebral cytoprotection that the transplanted stem cells have the capability to secrete VEGF (Wakai et al., 2016). Hypoxia (0.5% O2, 36 h) also promotes the proliferation of neural crest stem cells (NCSCs), which are a population of adult multipotent stem cells through upregulating HIF-1α (Chen et al., 2017). Similarly, intermittent hypoxia (10% O2 for 20 cycles, 40-min duration overall) increases NPC populations within the SVZ of the lateral ventricles and in the subgranular zone of the hippocampal dentate gyrus. Further experiments in vitro demonstrate that intermittent hypoxia markedly enhances the capacity for expansion of a cultured neurosphere, which is accompanied by increases in a proliferation marker (Ki67), MTT activity, and HIF-1α signaling (Ross et al., 2012).

The underlying mechanisms of HIF-1-mediated NSC proliferation or self-renewal entail the transcriptional expression of VEGF via HIF-1, which in turn stimulates neurogenesis (Harms et al., 2010). Another HIF-1 target gene (EPO) also promotes proliferation of NSCs by modulating NF-kb (nuclear factor-kappa B) signaling (Shingo et al., 2001). Additionally, WNT/β-catenin signaling is implicated in the effects of HIF-1α on NSC proliferation. HIF-1α deletion leads to impaired WNT/β-catenin activity in the hippocampus, with impaired NSC proliferation (Mazumdar et al., 2010; Braunschweig et al., 2015; Hubbi and Semenza, 2015). Of note, our latest unpublished data show that autophagy induced by HIF-1 mediates NSC proliferation. Together, the above studies indicate that HIF promotes NSC self-renewal or proliferation through different signaling pathways. A number of hypoxic conditions are conducive to the self-renewal capacity of NSCs via the HIF-1 pathway. On the other hand, hypoxia does not specifically and exclusively affect NSC proliferation, due to the number of its downstream effectors, apart from the HIF-1 pathway. Furthermore, HIF-1α is unstable under normoxic conditions, and as such is easily degraded by the proteasome system (Prabhakar and Semenza, 2012).

**HIF UBIQUITINATION DEGRADATION AND PHDs**

As previously mentioned, in normoxia HIF-1α is subjected to oxygen-dependent hydroxylation by the prolyl hydroxylase domain proteins PHD1-3 at its proline residues 402 and 564, which are followed by the binding of a von Hippel-Lindau protein VHL in an ubiquitin ligase complex (Epstein et al., 2001; Ivan et al., 2001; Jaakkola et al., 2001; Yu et al., 2001). The resulting event is the ubiquitination and proteosomal degradation of HIF-1α. The half-life of HIF-1α is only a few minutes when exposed to 20% O2 under standard tissue culture conditions (Prabhakar and Semenza, 2012).

PHDs are recognized as “oxygen sensors” at the cellular level (Kaelin and Ratcliffe, 2008). The PHD family consists of three members, PHD1, PHD2, and PHD3, which share conserved
COOH-terminal regions responsible for hydroxylase activity, while at the NH2-terminus they are greatly different (Epstein et al., 2001). Each isoform is tissue or cell line-specific as well as subcellular specific (Chen et al., 2015). PHD1 is mainly expressed in the nuclei of the testis and liver (Tambuwala et al., 2010), PHD2 is localized mainly in the cytoplasm in the heart and testis (Selvaraju et al., 2014), and PHD3 has both nuclear and cytoplasmic localization mainly in the heart and liver (Metzen et al., 2003). In addition, they have different regulatory patterns on the modulation of HIF-α: PHD2 is the primary regulator of HIF-1α, and although PHD3 is less active than PHD2 in regulating HIF-1α, it is more likely to target HIF-2α under hypoxic conditions (Appelhoff et al., 2004).

The hydroxylation reaction of HIF-1α is controlled by the activity of PHDs, which is regulated by O2, Fe2+, α-ketoglutarate, and ascorbic acid (Kaelin and Ratcliffe, 2008; Wong et al., 2013). Under hypoxic conditions, when the hydroxylation reaction is inhibited due to a lack of oxygen, HIF-1α is stabilized and enters the nucleus to form the HIF-1 heterodimer with HIF-1β, which activates the transcription of 100s of target genes and leads to a synergistic adaptation to hypoxia.

It is noteworthy that the prolyl hydroxylase enzymes contain Fe2+ in their catalytic centers; therefore, HIF-1α can also be stabilized pharmacologically through the addition of iron chelators such as desferrioxamine (DFO) or iron antagonists such as cobalt chloride (CoCl2), in addition to hypoxia. Furthermore, since α-ketoglutarate is a substrate of PHDs, its competitive antagonists [including dimethylxaloylglycine (DMOG)] can stabilize HIF-1α (Epstein et al., 2001). Additionally, multiple proteins have been identified that promote oxygen-dependent degradation of HIF-1α, for example OS9 (ortholog of fungi Yos9p) and RUNX3 (runt related transcription factor 3), which increase the efficiency of hydroxylation of HIF-1α via binding to both HIF-1α and PHD2 (Baek et al., 2005; Lee et al., 2013); and LIMD1 (LIM domains containing 1) alongside RHOBTB3 (Rho-related BTB domain-containing protein 3) can form a complex with HIF-1α, PHD2, and VHL to promote hydroxylation-mediated ubiquitination (Zhang et al., 2015). In any case, PHDs play a crucial role in the instability of HIF-1α. Therefore, inhibition of PHD activity is essential for stabilizing and activating the transcriptional activity of HIF-1.

**PHD INHIBITORS AND PROLIFERATION OF NSCs**

As discussed above, the employment of hypoxia-mimetics to provide HIF-1α stabilization might facilitate the expansion of NSCs in vitro. However, the commonly used agents for stabilizing HIF-1α, including CoCl2, DFO or DMOG, not being specific inhibitors of PHD, have side effects on vital cellular processes. Some agents demonstrate cytotoxicity, spanning from long- (e.g., DFO) to short- (e.g., CoCl2) term, to proliferating NPCs, leading to decreased expression of cyclin D1 and nestin in

![FIGURE 1](https://example.com/figure1.png)
MSCs (Lovejoy and Richardson, 2002; Parmar et al., 2007; Milosevic et al., 2009), while others showed a combination of antiproliferative and cytotoxic effects (e.g., DMOG). However, DMOG-treated human-induced pluripotent stem cell-derived MSCs (hiPSC-MSCs) show improved expressions of HIF-1α and its downstream genes, indicating that DMOG significantly enhances the expression profiles of angiogenic-related factors in hiPSC-MSCs via maintaining HIF-1α stability. In a critical-sized calvarial defect model in rats, DMOG-treated hiPSC-MSCs showed increased angiogenic capacity in the tissue-engineered bone, which results in bone regeneration (Zhang et al., 2016). However, in this study, the side effects of DMOG were not evaluated. A new selective PHD inhibitor FG-4497 has been developed, which stabilizes HIF-1α and induces the expression of HIF-1 target genes. It has been reported that FG-4497 ameliorates mucosal damage in experimental murine colitis (Robinson et al., 2008). In addition, FG-4497 is also reported in HSCs to stabilize HIF-1α protein in vivo in the bone marrow (BM), increase the proportion of HSCs and primitive hematopoietic progenitor cells (HPCs) into quiescence, and accelerate blood recovery after severe irradiation (Forristal et al., 2013). Importantly, FG-4497 promotes cell proliferation without being toxic to NPCs and increases the number of cells in normoxia, comparable to hypoxia-induced effects, and is also beneficial for dopaminergic differentiation (Milosevic et al., 2009). Thus, FG-4497 appears to be a non-toxic small molecule, capable of inducing neurogenesis in vitro.

Endogenous inhibitors such as reactive oxygen species (ROS) can also block the activity of PHDs, thereby stabilizing HIF-1α. Mitochondria are the main intracellular source of ROS (Solaini et al., 2010; Scherz-Shouval and Elazar, 2011). Other intracellular sources include NADPH oxidases (NOX) (Finkel, 2003). The amount of ROS produced at a cellular level may play a key role in determining NSC fate. In one study, high levels of ROS produced by NOX activity in hypoxic conditions (4% O2) were associated with NSC regeneration, while low endogenous ROS levels resulted in a decrease in NSCs’ capacity for self-renewal (Le Belle et al., 2011). The proposed mechanism entailed enhanced ROS-dependent NSC self-renewal and neurogenesis via the PI3K/AKT signaling pathway. Conversely, decreased levels of cellular ROS as a result of pharmacological or genetic manipulations were shown to interfere with NSC function in vitro and in vivo (Kim and Wong, 2009; Chuikov et al., 2010).

PHD activity is closely related to ROS levels in many systems. Relatively low levels of ROS increase PHD activity, whereas high ROS levels stabilize HIF-1α as a result of inactivating PHDs (Callapina et al., 2005; Fedotcheva et al., 2006; Hamanaka and Chandel, 2009). However, to date, the mechanism by which ROS directly affect PHD function has not been clarified, since stabilization of HIF-1α can occur independently of mitochondrial ROS production during hypoxia, indicating that PHD activity may not be directly regulated by ROS (Wong et al., 2013). One study proposed that H2O2 may affect PHD activity more directly through a local Fenton reaction (Gerald et al., 2004). Although ROS have been implicated in the regulation of PHD activity, this modulation in NSCs has not been reported. The graphical abstract reported in Figure 1 summarizes these studies and puts forward the concept that the proliferation and differentiation of NSCs are regulated by upstream PHD activity and the core transcription factor HIF-1.

CONCLUSION AND FUTURE PERSPECTIVES

As previously described, hypoxia-dependent regulation of NSC proliferation can be simulated with pharmacological interventions acting via HIF-1α stabilization. Several PHD inhibitors have the potential to promote NSC proliferation; however, their side effects hinder their use. Therefore, the development of new specific and effective PHD inhibitors may offer a valid tool for the application of NSC transplantation in neurodegenerative diseases. Additionally, targeted hypoxia-mimicking agents need to be developed specifically for neural cells for future clinical application.

AUTHOR CONTRIBUTIONS

L-YW and L-LZ discussed and wrote the manuscript. L-YW and Y-LH approved the final version to be published.

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