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Hypoxia-regulated angiogenic inhibitors

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Abstract

The regulation of angiogenesis by hypoxia is an essential homeostatic mechanism that depends on a precise balance between positive and negative angiogenic regulatory molecules. Pro-angiogenic factors are well characterized; however, several *in vivo* and *in vitro* studies indicate that there are feedback mechanisms in place to inhibit angiogenesis during hypoxia. Understanding the signaling pathways leading to the negative feedback of angiogenesis will undoubtedly provide important tools to develop novel therapeutic strategies not only to enhance the angiogenic response in coronary artery disease but also to hinder deregulated angiogenesis in tumorigenesis.

Introduction

Oxygen homeostasis is critical for the survival of all multicellular organisms. The response to low oxygen levels (hypoxia) stimulates the oxygen-delivery system and provides mechanisms to activate cell death or survival pathways, depending on the context of the hypoxia signal. Systemically, highly sensitive tissues detect acute hypoxia and in response increase respiration and cardiac output. Extended hypoxia is sensed at the cellular level, which leads to the induction of angiogenesis, to increase delivery of oxygen and nutrients to tissues. This is accomplished by the *de novo* sprouting of capillaries from post-capillary venules, and in adults is stimulated mainly via the induction of hypoxia-inducible factor (HIF-1) expression.

The hypoxic response is significantly controlled in most cells by HIF-1, a heterodimeric transcription factor composed of the nearly ubiquitous HIF-1 α and its dimerization partner HIF-1 β . HIF-1 activates approximately 200 genes encoding proteins that regulate cellular metabolism, proliferation, motility, hematopoiesis, and angiogenesis (Semenza 2000). Upon initiation of the hypoxic signal, HIF1- α translocates to the nucleus, dimerizes with HIF1- β to form the HIF-1 complex and induces the expression of its transcriptional targets via binding to hypoxia-responsive elements (HREs) (Chilov et al. 1999). HREs are present in many angiogenic genes, such as VEGF, angiopoietin-2, VEGF receptors (Flt1 and KDR), and neuropilin-1 (Hickey and Simon 2006; Simons 2005). Hypoxia can up-regulate these angiogenic molecules by several mechanisms, including direct transcriptional activation by HIFs or indirect up-regulation by HIF-induced molecules. In addition, other transcription factors induced by hypoxia, such as Related Transcription Enhancer Factor-1 (RTEF-1) and early growth response 1 (EGR-1), can both target VEGF to enhance angiogenesis (Shie et al. 2004; Yan et al. 2000). Additional angiogenic growth factors such as IGF are also induced by

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hypoxia, but can signal through a HIF-1-independent pathway (Slomiany and Rosenzweig 2006).

Angiogenesis is important in physiological conditions such as embryonic development and wound healing, as well as pathological conditions including tumorigenesis, diabetic retinopathy, rheumatoid arthritis, and atherosclerosis (Fong 2008). Moreover, hypoxia is associated with virtually all forms of vascular disorders, such as coronary and peripheral arterial diseases, including stroke, myocardial and limb ischemia; lung disorders; and diabetes (Fong 2008). Severe hypoxia is also found in solid tumors, where capillary networks are insufficiently organized (Folkman 2006). Physiological stresses such as hypoxia are regulated by a complex balance of both stimulatory and inhibitory signals that promote or inhibit angiogenesis. Specifically, understanding the role and regulation of genes during angiogenesis is becoming increasingly important to elucidate the compensatory hypoxic response. In the present review, we will mainly discuss the anti-angiogenic feedback mechanisms in the HIF-1- and VEGF-related angiogenic pathways.

HIF-1-related anti-angiogenesis

As a hypoxia-induced transcription factor, HIF-1 both stimulates and represses a multitude of genes important for adaptation to the low oxygen environment. Regulator of G protein Signaling 5 (RGS5) is a HIF-1-dependent, hypoxia-induced angiogenic inhibitor (Jin et al. 2009) that functions as a negative regulator of G protein-mediated signaling (Adams et al. 2000; Bell et al. 2001).

One of our previous reports showed that hypoxia specifically increased RGS5 expression in endothelial cells, which is confirmed in the DNA microarray data in Table 1 (Jin et al. 2009). RGS5 mRNA expression was induced by hypoxia while two other family members, RGS2 and RGS4, were not impacted. In addition to changes in oxygen levels, HIF-1 α played a key role in hypoxia-induced RGS5 expression by stimulating RGS-5 promoter activity in endothelial cells. RGS5 slowed endothelial cell growth and significantly enhanced the apoptotic protein Bax, which led to increased apoptosis due to the change in the Bcl-2/Bax ratio (Jin et al. 2009; Yang and Korsmeyer 1996). Furthermore, RGS5 inhibited VEGF-induced angiogenesis through the p38 MAPK-dependent pathway *in vitro* and *in vivo*.

Apoptosis occurring in endothelial cells causes an inhibitory effect on cell proliferation, which has a similar impact to that of anti-angiogenic factors. Hypoxia promotes apoptosis in endothelial cells, as demonstrated by changes in p53 protein levels (Hammond and Giaccia 2005; Stempien-Otero et al. 1999). Several studies suggest that HIF-1 α stabilizes p53 and contributes to hypoxia-induced p53-dependent apoptosis (Xenaki et al. 2008). A recent report proposed that P300/CBP-associated factor (PCAF), an HIF-1 α cofactor, regulates the balance between cell cycle arrest and apoptosis in hypoxia by modulating the activity and protein stability of both p53 and HIF-1 α (Xenaki et al. 2008). Previous reports demonstrated that p53 can also inhibit angiogenesis by the following mechanisms: elevating thrombospondin-1 levels, repressing both VEGF and FGF-2, and by inducing degradation of HIF-1 (Folkman 2006).

HIF-1 α mediates hypoxia-induced apoptosis through the up-regulation of genes as well as suppression of genes such as Bcl-2 (Carmeliet et al. 1998; Jin et al. 2009). One member of the Bcl-2 family, Bcl-2/E1B interacting protein (BNIP3) is up-regulated under hypoxic conditions (Chen et al. 1997; Kothari et al. 2003). Kothari and colleagues showed that blocking hypoxia-induced BNIP3 expression using siRNA or a mutant BNIP3 inhibits hypoxia-induced cell death. Additionally, hypoxia-mediated BNIP3 expression occurs by direct binding to an HRE in the human BNIP3 promoter, and mutation of this HRE site eliminates the hypoxic responsiveness of the promoter. Furthermore, hypoxia-induced BNIP3 expression was

detectable 24 h after initial up-regulation of HIF-1 α , indicating that BNIP3 expression occurs much later in the hypoxic response than that of other genes such as angiogenesis factor VEGF (Kothari et al. 2003).

While HIF-1 mainly targets pro-angiogenic genes, there are feedback molecules to counteract this up-regulation, such as endostatin, which down-regulates HIF-1 α (Abdollahi et al. 2004). Hypoxia-induced endostatin inhibits VEGF-induced angiogenesis by preventing proliferation and migration of endothelial cells (Abdollahi et al. 2004; Heljasvaara et al. 2005; Morbidelli et al. 2003) and simultaneously up-regulates anti-angiogenic genes, including thrombospondin, vasostatin, and kininogen (Abdollahi et al. 2004). Reduced oxygen supply in FVB mice exposed to a hypobaric hypoxia chamber showed enhanced MMP production leading to a subsequent increase of endostatin generation in the lung and aorta (Paddenberg et al. 2006). These results were confirmed by Suhr et al., who reported a significant increase in the endostatin and MMP-9 plasma concentration following hypoxic conditions in human cyclists (Suhr et al. 2007).

During sustained exposure to hypoxia, vascular beds may react differently. For example, a chronically hypoxic lung has structural changes in the pulmonary vascular bed which can become the major determinant of elevated vascular resistance (Fried et al. 1983). In the previously mentioned study, Paddenberg et al. questioned whether endostatin participates in the vascular alterations occurring in the lung under chronic hypoxia (Paddenberg et al. 2006). They found a hypoxia-induced increase in endostatin generation and MMP-2 activity in the lung, which corresponds with the findings of Frisdal et al. who demonstrated enhanced MMP-2 activity in rat pulmonary vessels during hypoxia-induced pulmonary hypertension (Paddenberg et al. 2006, Frisdal, 2001 #62). In the pulmonary vasculature, hypoxia-induced angiogenesis counteracts the development of pulmonary hypertension caused by vasoconstriction and vascular remodeling, while hypoxia-induced angiogenic inhibitors, i.e. endostatin would contribute to the development of pulmonary hypertension.

VEGF-associated anti-angiogenic regulation

While hypoxia-induced VEGF initiates angiogenesis, it also up-regulates expression of molecules within the endothelium that act as negative feedback regulators to modulate excessive vascular sprouting and endothelial cell proliferation. In addition to the aforementioned molecules, our previous report by An et al. demonstrated that Response Gene to Complement (RGC)-32 is another important VEGF-inducible gene that serves as a negative regulator of hypoxia-induced angiogenesis. RGC-32 expression was significantly increased under hypoxic conditions. Hypoxic induction of RGC-32 was also mediated by HIF-1 α at both the transcriptional and posttranscriptional levels (An et al. 2009). While HIF-1-induced RGS5 was not VEGF-mediated (Jin et al. 2009), RGC-32 expression was significantly increased by VEGF but not factors including TNF- α , FGF2 and IL-1 β (An et al. 2009). Unlike genes such as COX-2 (Wu et al. 2003), RGC-32 did not follow the canonical VEGF-induced angiogenic pathway. Overexpression of RGC-32 destabilized vascular structure formation *in vitro* by down-regulating FGF-2 and cyclin E, which caused negative feedback to VEGF activation. Additionally, when angiogenesis was examined *in vivo* in a mouse model, RGC-32 drastically inhibited VEGF-induced angiogenesis in matrigel, attenuated the recovery rate in hindlimb ischemia and reduced tumor size. (An et al. 2009).

Similarly, Delta-like ligand 4 (Dll4) is induced by VEGF (Lobov et al. 2007) yet has anti-angiogenic properties. Dll4 is part of the Notch signaling pathway (Liu et al. 2003) and highly expressed in the vascular endothelium, largely in angiogenic blood vessels (Lobov et al. 2007). Lobov, et al. demonstrated that Dll4 is an antagonistic regulator of angiogenesis by injecting a soluble version of Dll4 (Dll4-Fc) that blocks Dll4/Notch interactions into the vitreous membrane of oxygen-induced ischemic retinopathic (OIR) mice. Dll4 blockade

markedly enhanced angiogenic sprouting while suppressing ectopic pathological neovascularization in the retinal vasculature. (Lobov et al. 2007). Therefore, Dll4 plays a role as a negative regulator of sprouting angiogenesis in response to the release of hypoxia-induced factors such as VEGF (Lobov et al. 2007).

In addition to VEGF-induced anti-angiogenic genes, the hypoxia-induced VEGF, itself, is also subjected to negative regulation, such as by transcription factor E2F1 (Qin et al. 2006). Under hypoxic conditions, it is well known that VEGF is induced by HIF-1; however, another transcription factor, E2F1 down-regulates expression of VEGF by associating with p53 and specifically down-regulating VEGF expression but not other hypoxia-inducible genes. Studies of E2F1^{-/-} mice under hypoxic conditions revealed enhanced angiogenesis, dependent on deregulated VEGF, which illustrates a novel negative regulation to VEGF.

Other hypoxia-related angiogenic inhibitors

Several hypoxia-induced angiogenesis inhibitors have been discovered. The DNA array analysis in endothelial cells under hypoxic vs. normoxic conditions shown in Table 1 suggests that angiopoietin-2 (Ang2) is also important in the inhibition of angiogenesis induced by hypoxia. Previous reports determined that hypoxia enhances Ang2 expression *in vivo* as well as *in vitro* (Oh et al. 1999). The angiopoietin ligands/Tie receptors belong to a class of ligand/receptor families that plays a critical role in angiogenesis, cell migration, proliferation, and survival. Ang1 and Ang2 are both ligands with a similar binding affinity for the Tie2 receptor, which is a member of the receptor tyrosine kinases (RTKs) predominantly expressed by vascular endothelial cells (Maisonpierre et al. 1997). Ang1 induces angiogenesis via autophosphorylation of Tie2, while Ang2 competitively inhibits this effect (Maisonpierre et al. 1997). Ang2 is an antagonist for the Ang1/Tie2 pathway, as supported by data from over-expressing Ang2 mice that showed a very similar phenotype to the Ang1 and Tie2 single knockouts (Yuan et al. 2009). Thus, hypoxia-induced Ang2 plays a role in the negative regulation of angiogenesis by competing with Ang1.

The negative feedback of angiogenesis during hypoxia, shown in the proposed schematic in Figure 1, may explain the inadequate natural collateral circulation in hypoxic heart diseases, in which up-regulated VEGF inhibits other pro-angiogenic signaling or triggers angiogenic inhibitor signaling. Therapeutic angiogenesis with either HIF-1 or VEGF resulted in a marked increase in blood flow and improved cardiac function in animal studies without apparent toxicity (Simons 2005). However, the results of clinical trials have been inconsistent and largely disappointing (Simons 2005), presumably because exogenous pro-angiogenic factors such as VEGF induce endogenous angiogenic inhibitor signaling to balance the effect; therefore, these feedback inhibitors will likely be new targets in future studies.

Concluding remarks

Intensive studies in the last decade have established the hypoxia signaling pathway as an important therapeutic target for various vascular diseases. Yet many questions still remain. Which pro- or anti-angiogenic factors will dominate in certain conditions? Which transcription factors regulate hypoxia-induced pro- or anti-angiogenic molecules? Understanding the signaling pathways leading to the negative feedback regulation of angiogenesis will undoubtedly provide an important means to develop therapeutic strategies that enhance neovascularization of ischemic tissue and interfere with focal, dysregulated vascular remodeling, the key mechanism for atherosclerotic disease progression as well as pulmonary hypertension. In contrast, selective stimulation of these negative feedback regulators is a promising prospect for tumor therapy.

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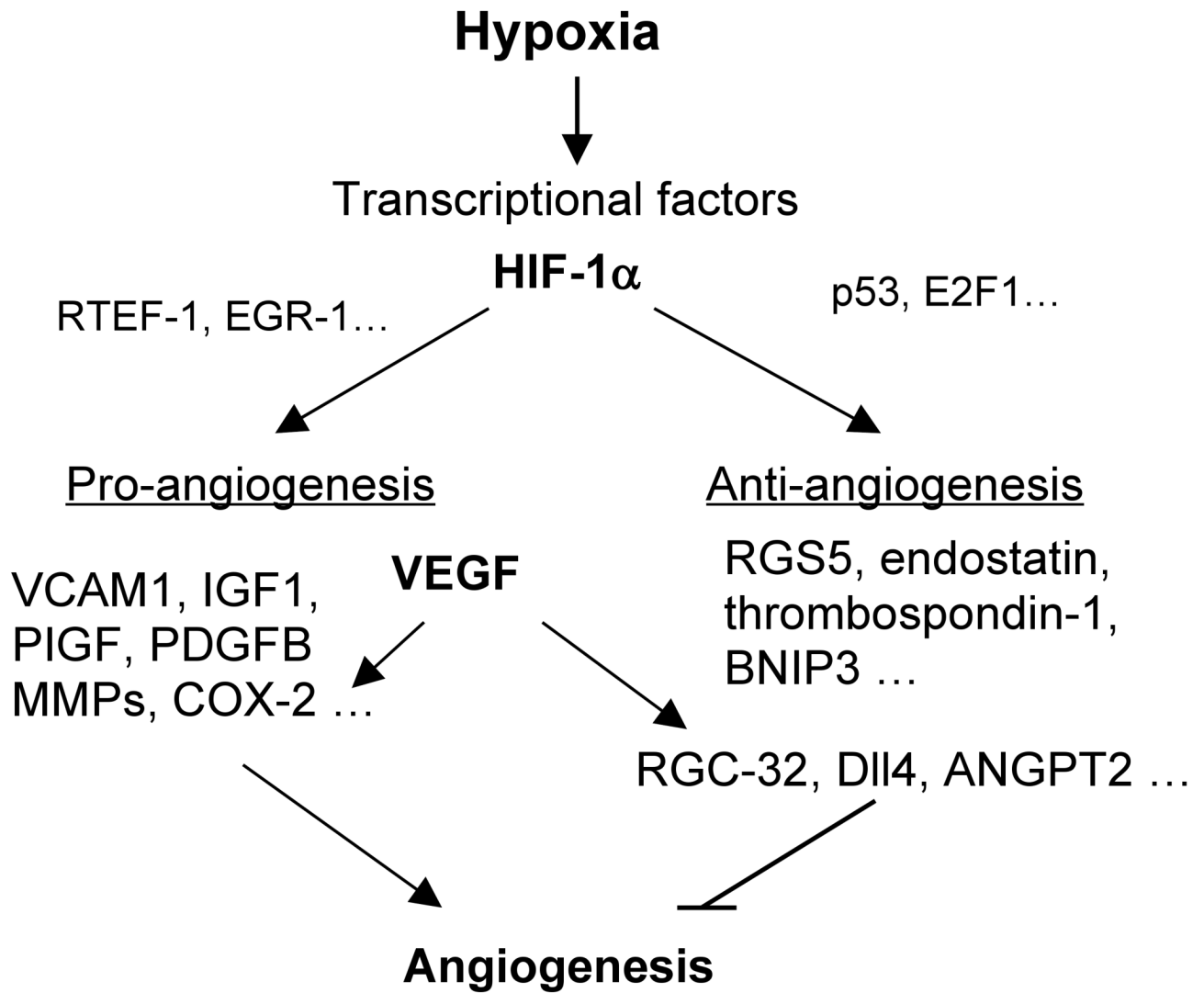


Figure 1. Schematic of hypoxia-mediated angiogenesis and anti-angiogenesis

Table 1
Hypoxia-induced anti-angiogenic factors

DNA Microarray data from hypoxia-treated human umbilical vein endothelial cells (HUVECs) analyzed by Ingenuity Pathway Analysis software. This table shows molecules associated with inhibition of angiogenesis that were enhanced upon hypoxia in comparison with normoxia.

Symbol	Entrez Gene Name	p value
TIE1	tyrosine kinase with immunoglobulin-like and EGF-like domains 1	2.00E-05
RGS5	Regulator of G protein signaling 5	2.00E-05
COL181A	collagen, type XVIII, alpha 1/endostatin	2.00E-05
HHEX	hematopoietically expressed homeobox	2.00E-05
LAMA5	laminin, alpha 5	2.00E-05
SEMA3F	sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3F	2.10E-05
STAB1	stabilin 1	4.00E-05
THDS1	thrombospondin, type I domain containing I	4.60E-05
ANGPT2	angiopoietin 2	1.89E-04
PLAT	plasminogen activator, tissue	2.19E-04
PGK1	phosphoglycerate kinase 1	2.19E-04
ADAMTS1	ADAM metalloproteinase with thrombospondin type 1 motif, 1	3.46E-04
SPARC	secreted protein, acidic, cysteine-rich (osteonectin)	3.88E-04
PDE4B	Homo sapiens phosphodiesterase 4B (cyclic nucleotide phosphodiesterases)	3.89E-04
NOTCH1	Notch homolog 1 (Drosophila)	3.89E-04
WARS	tryptophanyl-tRNA synthetase	3.89E-04
BMPRI1A	bone morphogenetic protein receptor, type 1A	4.68E-04
PTX3	pentraxin-related gene, rapidly induced by IL-1 beta	5.52E-04
PDCD5	programmed cell death 5	6.18E-04
SERPINE1	serpin peptidase inhibitor, clade E (nexin, plasminogen activator inhibitor type 1), member 1	6.92E-04
TGFB2	transforming growth factor, beta 2	1.32E-03
PAWR	PRKC, apoptosis, WT1, regulator	2.25E-03
BNIP3	BCL2/adenovirus E1B 19kDa interacting protein 3	4.48E-03
PTRPM	protein tyrosine phosphatase, receptor type, M	5.64E-03