

## Review

# The magic of the hypoxia-signaling cascade

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**Abstract.** All organisms respond to changes in their environment by activating complex signaling cascades. The “hypoxia-signaling cascade” is activated in response to low oxygen availability and this activation is central to maintaining oxygen homeostasis and hence to survival. By regulating the tran-

scriptional complex hypoxia-inducible factor, hypoxia is associated with several physiopathological processes. Several strategies, based on the targeting of the hypoxia-signaling cascade, have been developed to treat these pathologies. Our review summarizes different aspects of the hypoxic pathway.

**Keywords.** Oxygen homeostasis, hypoxia, HIF signaling, physiopathology, therapy.

## Introduction

In 1774, the scientist Joseph Priestley realized that “a mouse kept in a jar cannot survive in the absence of air renewal”. Without knowing it, he had just discovered the oxygen molecule (O<sub>2</sub>) and highlighted its absolute requirement for survival. All organisms rapidly respond to changes in oxygen availability. This response, triggered by many oxygen-sensing systems located at different levels in the organism, activates complex signaling networks, which culminate in the control of gene expression.

The signaling pathways activated by low oxygen availability, or hypoxia, allow cell adaptation and survival in the hypoxic environment. Among them, activation of the transcriptional complex hypoxia-inducible factor (HIF) contributes significantly to this adaptive response and constitutes a major regulator of oxygen homeostasis [1]. Since its discovery, the HIF

cascade has been extensively studied and its role appears clearly evident in both physiological and pathological processes such as embryogenesis, tissue repair, ischemia and cancer.

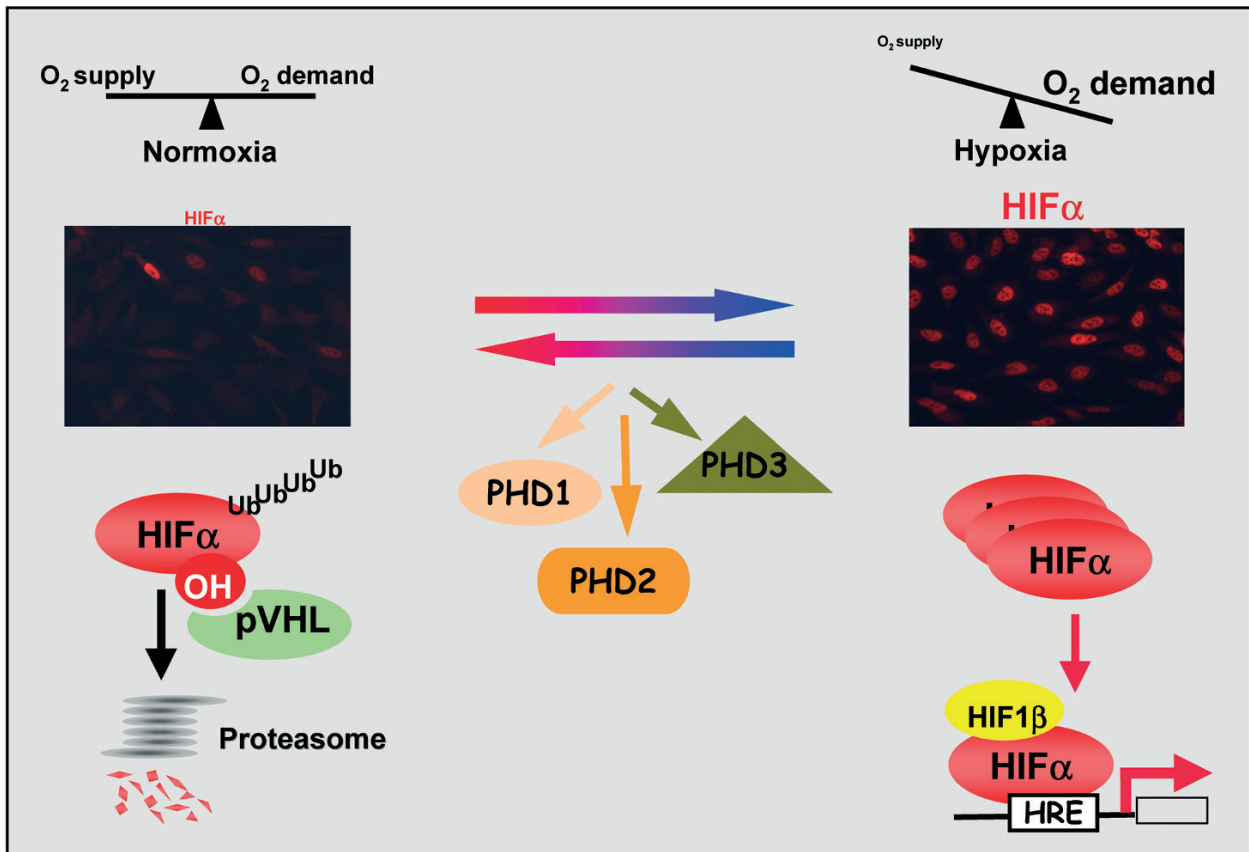
This review summarizes our current knowledge of the HIF signaling cascade, its implication in physiopathology and the different therapeutic strategies developed so far to target this pathway.

## The hypoxia inducible factor

HIF is a transcriptional factor composed of the constitutively expressed HIF1 $\beta$  subunit and one of the three HIF $\alpha$  subunits (HIF1 $\alpha$ , HIF2 $\alpha$ , HIF3 $\alpha$ ) [2–4]. Among them, HIF1 $\alpha$  is the most extensively studied isoform. O<sub>2</sub>-dependent HIF $\alpha$  regulation is essentially post-transcriptional and involves both regulation of its stability and its activity by a mechanism that is inherently O<sub>2</sub> dependent: the hydroxylation of proline (Pro) or asparagine (Asn) residues [5–7]. Indeed, hydroxylation of the Asn<sup>803</sup> (in the

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**Figure 1.** O<sub>2</sub>-dependent regulation of hypoxia-inducible factor (HIF) $\alpha$  stability: Low oxygen availability stops the otherwise permanent HIF $\alpha$  degradation pathway, which implicates the proteasome and the product of the tumor suppressor gene von Hippel-Lindau (pVHL). The degradation pathway is triggered by hydroxylation of two proline residues within the oxygen-dependent degradation domain (ODDD) of the HIF $\alpha$  subunit by the prolyl hydroxylase domain-containing proteins (PHDs). These enzymes, acting as O<sub>2</sub> sensors, play a major role in the regulation of O<sub>2</sub> homeostasis and thus survival. Immunofluorescence inserts show the expression of HIF1 $\alpha$  in well-oxygenated cells (normoxia) or upon low O<sub>2</sub> availability (hypoxia).

human HIF1 $\alpha$ ), within the C-terminal transactivation domain (C-TAD) by factor inhibiting HIF (FIH), inhibits the binding of the p300/CBP co-activators, and hence the transcriptional activity of the protein [7, 8]. Moreover, an oxygen-dependent degradation domain (ODDD) is involved in the O<sub>2</sub>-dependent regulation of HIF $\alpha$  proteasomal degradation, which constitutes the limiting step in HIF $\alpha$  regulation, and consequently HIF activation [9, 10].

In well-oxygenated cells, HIF $\alpha$  is an exceptionally short-lived protein (half-life <5 min) and steady-state levels are very low [9]. In contrast, reduced O<sub>2</sub> availability accumulates HIF $\alpha$  by relaxing its ubiquitin-proteasome degradation. More precisely, ubiquitination and proteasomal degradation of HIF $\alpha$  require pVHL, the product of the von Hippel-Lindau tumor suppressor gene, which functions as an ubiquitin E3 ligase [11, 12]. The hydroxylation of two conserved proline residues (Pro<sup>402</sup> and Pro<sup>564</sup>, in the human HIF1 $\alpha$  sequence), contained within the ODDD, triggers pVHL binding and thus HIF1 $\alpha$

proteasome targeting [5, 6]. Stabilization of HIF $\alpha$  is due to disruption of the pVHL/HIF $\alpha$  interaction under hypoxic conditions (Fig. 1). Two independent groups identified the enzymes catalyzing the hydroxylation reaction: the HIF prolyl hydroxylases or prolyl hydroxylase domain-containing proteins (PHDs) [13, 14]. These enzymes belong to the superfamily of the Fe(II) and 2-oxoglutarate-dependent dioxygenases. They need the O<sub>2</sub> as a co-substrate, providing the molecular basis for their O<sub>2</sub>-sensing function (*in vitro* K<sub>m</sub> values for O<sub>2</sub>= 230–250  $\mu$ M) [15]. In addition, PHDs use Fe(II) and ascorbate as co-factors [16]. Fe(II) is critical for activating O<sub>2</sub> and as a template for the orderly binding of reactants. The ascorbate is necessary for maximal prolyl hydroxylase activity because it reduces the Fe(III) into Fe(II), and thus reactivates the enzymes.

Epstein et al. [14] identified three human and mouse *phd* genes (*phd1*, *phd2*, and *phd3*). The three isoforms are ubiquitously expressed, albeit at different levels [17, 18]. Fused to a green fluorescent protein (GFP),

PHD1 appears exclusively located in the nucleus, PHD2 mainly in the cytoplasm, and PHD3 in both compartments [19]. The contribution of each PHD to HIF $\alpha$  regulation depends on their relative abundance [20]. Nevertheless, we have shown that PHD2 is the rate-limiting enzyme that sets the low steady-state levels of HIF1 $\alpha$  in normoxia, whereas PHD1 and PHD3 contribute to HIF1 $\alpha$  regulation only upon a chronic hypoxia [21]. Furthermore, the expression and the activity of PHDs are tightly regulated at both transcriptional and post-transcriptional level [16, 22]. Initially identified as the regulator of the *epo* gene expression, HIF was further shown to control many hypoxia-target genes [22]. Indeed, by promoting the transcription of all these genes, HIF helps to maintain O<sub>2</sub> homeostasis. Among them, the vascular endothelial growth factor (VEGF) is essential for angiogenesis, inducible nitric oxide synthase (iNOS) and Heme Oxygenase 1 (HO1) favor vasodilatation, tyrosine hydroxylase (TH) regulates respiratory frequency, the glucose transporter GLUT-1 is involved in the anaerobic glycolysis and EPO is essential for erythropoiesis.

The activation of the HIF complex can be of major importance in certain physiological and pathological situations.

### Physiology of the HIF cascade

#### Embryonic development

Hypoxia constitutes an important feature of the embryonic development by promoting the establishment and the differentiation of the vascular and hematopoietic systems. Indeed, as early as 1 week after fertilization, the nutrients and O<sub>2</sub> supplied by passive diffusion are not sufficient for the growing embryo and hypoxic gradients are subsequently established. This process, by promoting the expression of HIF-dependent genes such as *epo* and *vegf*, is absolutely required since mice deficient in these genes die during the embryo development as a consequence of severe hematopoietic and vascular defects [23–25]. Indeed, the phenotypic features of the knockout mice for *hif1 $\alpha$* , *hif2 $\alpha$* , *hif1 $\beta$* , and *phd3* agree with the central role of the HIF signaling pathway in the establishment of key physiological systems during the embryonic development.

**HIF1 $\alpha$ .** In normal embryos, HIF1 $\alpha$  expression increases between embryonic day (E) 8.5 and E9.5. HIF1 $\alpha$  knockout mice stop their development at E8.5 and die at E10.5 because of vascular deficiencies and cardiac and neuronal abnormalities [26–28]. The vascular reduction in *hif1 $\alpha$* <sup>-/-</sup> embryos is associated

with a decrease in VEGF production. However, these vascular abnormalities occur at E9.25, whereas in *vegf*<sup>+/+</sup> embryos similar defects appear earlier, at E8.5 [23, 24]. It seems that *hif1 $\alpha$* <sup>-/-</sup> mice produce enough VEGF to initiate the vasculogenesis and to form the vascular plexus but the following formation and maturation of the vasculature established from this plexus (angiogenesis) cannot occur in the absence of HIF1 $\alpha$  expression.

**HIF2 $\alpha$ .** *hif2 $\alpha$* <sup>-/-</sup> mice also present a high embryonic lethality between E9.5 and E16.5 [29–32]. HIF2 $\alpha$  seems to be required for the remodeling of the vascular network in the embryo and the heart function due to the catecholamines produced by the Zuckerman organ [30, 32]. Two independent laboratories successfully obtained live *hif2 $\alpha$* <sup>-/-</sup> embryos [29, 31]. However, the respiratory, cardiac and hematopoietic functions of these mice were strongly affected. In addition, the replacement of HIF1 $\alpha$  by HIF2 $\alpha$  expression in knockout mice cannot rescue embryo survival [33]. Thus, in spite of the high sequence similarities, HIF1 $\alpha$  and HIF2 $\alpha$  have distinct and non-redundant functions that are essential, at least, for the embryo development.

**HIF1 $\beta$ .** The invalidation of HIF1 $\beta$  in mice also promotes a delayed embryo development together with vascular and placenta formation abnormalities leading to death at E10.5 [34, 35].

**PHDs.** The invalidation of the PHDs isoforms in mice highlighted specific functions for each PHD in the embryonic development [36, 37]. Indeed, *phd2* knockout promotes severe defects in the placenta and heart formation, leading to embryonic lethality between E12.5 and E14.5, whereas *phd1*<sup>-/-</sup> and *phd3*<sup>-/-</sup> mice are viable and apparently normal. Hence, PHD1 and PHD3 expression cannot rescue the *phd2*<sup>-/-</sup> phenotype. Accordingly to our previous report showing the key and unique role of PHD2 in the HIF1 $\alpha$  stability regulation *in cellulo*, these results highlight the critical role of PHD2 during embryonic development.

#### Postnatal life

Hypoxia, by activating the HIF transcription factor, controls the expression of several genes involved in the regulation of essential cellular processes during postnatal life.

**Growth.** The development of *Drosophila* flies in hypoxic conditions is normal but the size of the adults is smaller, due to a reduced cell growth [37]. In *Drosophila*, the Cdk4/CycD signaling pathway regulates cell growth. In Cdk4/CycD mutants whose

growth is impaired, the ectopic expression of the dmPHD (named Fatiga) is sufficient to rescue the normal phenotype [38, 39]. Cdk4/CycD induces the ribosomal protein mRpl12 that is required for cell growth stimulation and, in the case of mRpl12 mutants, in spite of normal Fatiga protein level, Fatiga activity is blocked promoting HIF activity [40]. The role of Fatiga involves the dmHIF1 $\alpha$  (named Sima) since its mutants are able to rescue the phenotype observed in those of Fatiga [41].

**Proliferation.** The activation of the angiogenic process and the anaerobic glycolysis *via* HIF stimulate cellular proliferation. However, Carmeliet et al. [26] reported that proliferation of tumors derived from ES HIF1 $\alpha$ <sup>-/-</sup> cells is more important than those from ES HIF1 $\alpha$ <sup>+/+</sup> cells. This discrepancy could be due in part to an increased p21 expression that promotes cell cycle arrest in hypoxia [42].

**Differentiation.** Hypoxia, in general, helps to maintain cells in an undifferentiated state. Indeed, hypoxia inhibits the differentiation of preadipocyte fibroblasts into adipocytes and myoblasts into myocytes [43, 44]. Nevertheless, some studies have shown that it can promote the differentiation of certain cell types [45, 46].

Recently, a link has been established between the HIF signaling and the Notch signaling, implicated in the differentiation of many stem cells; this link could explain, at least partially, the mechanism of dedifferentiation under hypoxia [47]. Indeed, under hypoxia, HIF1 $\alpha$  interacts with the intracellular domain (ICD) of Notch and so promotes the recruitment of HIF on the Notch-induced genes promoters and their expression. Furthermore, the Notch ligand Dll4 is up-regulated under hypoxia [48]. In endothelial progenitor cells, the hypoxic inductions of the arterial factors Dll4 and hey2 repress the expression of the venous regulator COUP-TFII and block the differentiation in veins of these cells. In addition, as a negative feedback loop, the factors hey1 and hey2 are able to repress HIF1 $\alpha$  transcriptional activity.

HIF-target genes can also regulate the functions of stem cells under hypoxia in a direct way. Indeed, during the embryonic development, the hypoxic induction of HIF2 $\alpha$  controls the expression of the Oct-4 transcription factor, which is essential to maintain the hematopoietic stem cells in an undifferentiated state and to regulate the embryonic stem cell differentiation [33]. Oct-4 is also expressed in several cancer cell lines and is induced under hypoxia in a renal carcinoma cell line strongly expressing HIF2 $\alpha$  [49].

In addition, two recent studies have also revealed some cross talk between HIF and the transforming

growth factor  $\beta$  (TGF- $\beta$ ) and Wnt-dependent signaling pathways [50, 51].

**Migration.** The ability of the leukocytes to migrate is enhanced by the HIF-dependent induction of CD11b/CD18, a member of the integrins superfamily [52, 53]. Hypoxia also favors the migration of dermal fibroblasts and keratinocytes [54, 55].

**Death.** The mechanisms regulating cell death under limited oxygenation conditions are complex. Indeed, not only the O<sub>2</sub> level, but also the degree of activation of certain oncogenes and nutrient deprivation determine the future behavior of the cells [56].

In general, drastic oxygen conditions (0.1% O<sub>2</sub> or anoxia) triggers cell death through the activation of pro-apoptotic proteins, like Bax, cytochrome c release, and caspases activation [57]. More precisely, this cell death is mediated by HIF. Indeed, HIF1 promotes the expression of the pro-apoptotic protein BNIP3, a member of the Bcl-2 family (Bcl2/adenovirus E1B 19Kda interacting protein 3), as well as the expression of NIX, homologous to BNIP3 [58]. The absence of HIF1 prevents cells from up-regulating BNIP3 and reduces the apoptotic process induced by anoxia [59]. Anoxia also activates the JNK pathway (c-Jun NH<sub>2</sub>-terminal kinase), which can promote the apoptotic process [60].

Contrary to anoxia, moderate hypoxia ( $\geq 1\%$  O<sub>2</sub>) does not affect cell proliferation and cell survival [61]. This cell survival is ensured by the PI3K/Akt activation, which inhibits the pro-apoptotic protein, Bad and favors NF- $\kappa$ B activity [62, 63]. In a same way, Degenhardt et al. [56] have recently shown that cells are able to survive by inducing the autophagic process. Hypoxia can also favor apoptotic resistance. In renal epithelial cells, Dong Z et al. [64] reported a better resistance of cells under hypoxia to a pro-apoptotic treatment such as staurosporine. In these cells, the hypoxic induction of the apoptotic inhibitory protein IAP-2 prevents Bax mitochondrial translocation. All these cellular functions regulated by the hypoxic signaling pathway are integrated in several biological processes such as inflammation or wound healing.

**Inflammation.** Low O<sub>2</sub> and glucose levels often characterize the environment of inflammatory sites such as the cutaneous infection, arthritis, or the necrotic regions of solid tumors [65–68]. At these sites, the myeloid cells have to evolve in spite of reduced oxygenation.

Hypoxia profoundly affects many properties of the myeloid cells: phagocytosis capacity, migration, adhesion, cytokines secretion, *etc.* More precisely, functional inactivation of the HIF1-dependent signaling

pathway has been shown to inhibit cell aggregation, motility, invasion and the capacity of cells to finally destroy the pathogen bacteria after their phagocytosis [69].

In addition, hypoxia promotes the secretion of inflammatory chemokines that attract neutrophils and monocytes. Furthermore, the macrophages-secreted cytokines interleukin 1 (IL-1) or tumor necrosis factor- $\alpha$  (TNF- $\alpha$ ) favor the expression and DNA binding of HIF1 $\alpha$  *via* the MAPK and PI3K pathways [70]. They also promote the production of reactive oxygen species, and thus contribute to HIF1 $\alpha$  protein stabilization [71].

Finally, hypoxia, by inhibiting PHDs, increases also the IKK $\beta$  activity that is involved in the inflammatory process by activating the NF- $\kappa$ B pathway [72].

**Wound healing.** Hypoxic regions are also present around the wound healing sites. The O<sub>2</sub> reduction in the wounded tissue is due in part to the modification of the vascular network but also, in a large part, to the O<sub>2</sub> consumption of the cells that strongly proliferate and are metabolically over-activated. These low-oxygenated areas contribute to the stimulation of the angiogenic process and tissue repair by the induction of several genes, such as *tgf $\beta$ 1*, *pro-collagen  $\alpha$ 1*, *vegf*, and *pdgf* (platelet-derived growth factor) [73]. Hypoxia around the wound promotes also the motility of the skin cells such as keratinocytes and favors wound closure by activating re-epithelialization [55]. Recently, Li et al. [74] have shown that hypoxia, by inducing HSP90 $\alpha$  secretion in the extracellular environment, promotes dermal fibroblasts migration. More precisely, HSP90 $\alpha$ , whose secretion is induced by HIF1 appears sufficient to increase fibroblasts motility.

### Pathology of the HIF cascade

HIF activity is involved in several pathologies. In the case of ischemia or amyotrophic lateral sclerosis (ALS), the insufficient activation of HIF, in spite of reduced oxygenation, promotes the progression of the disease, whereas in Alzheimer's disease (AD), cancer or pre-eclampsia, the sustained and strong activation of the pathway forms part of the basis of these pathologies.

### Neurological diseases

Mounting evidence shows a crucial role for hypoxia in neurological disorders such as ALS, Parkinson disease, AD or schizophrenia [75]. However, the implication of the HIF transcription factor has only been shown for the ALS and AD.

**ALS.** This is a late-onset progressive neurodegenerative disease affecting the motor neurons of the brain and the spinal cord [76]. Denervation of the respiratory muscles and diaphragm is generally the fatal event. The lifetime risk is at about 1 in 1000. Even now, 10 years after the approval of Riluzole, there is no effective therapy, although a few drugs are presently in Phase I, II and III clinical trials.

In mice, ALS can be mimicked by the deletion of the *vegf* HRE sequence [77]. Indeed, this disorder seems to be caused by a reduced blood perfusion of the neurons. Moreover, in ALS-affected rats, the direct administration of VEGF in the brain extends the motor neurons survival, protects the neuro-muscular junctions and finally favors the lifespan of these animals. These data highlighted the role of the HIF-dependent VEGF induction on motor neurons survival. Furthermore, Subramanian et al. [78] have recently identified a mutation in the gene coding for angiogenin (another hypoxia-induced protein involved in angiogenesis) in patients showing sporadic and familial ALS.

**AD.** An important feature of the AD is the formation of extracellular senile plaques in the brain, the major components of which are small peptides called  $\beta$ -amyloid (A $\beta$ ) derived from  $\beta$ -amyloid precursor protein (APP) [79, 80]. APP is sequentially cleaved first by the  $\beta$ -secretase ( $\beta$ -site amyloid precursor protein cleaving enzyme, BACE) and then by the  $\gamma$ -secretase complex to generate A $\beta$ . The molecular mechanism of this sporadic disease is still unknown. However, a reduced cerebral perfusion is a classical feature and one of the major contributors for the AD pathogenesis development. Ischemia or stroke gives rise to hypoxic conditions, which greatly increase the incidence of AD. Indeed, activation of the HIF1 signaling pathway could favor AD development at different levels: (i) APP, the expression of which is up-regulated in the post-ischemic brain [81]; (ii) the isoform 1 of the BACE  $\beta$ -secretase (BACE 1), which contains a functional HRE sequence [82]; (iii) the *aph-1a* promoter gene, which is a key component of the  $\gamma$ -secretase complex, and also contains an HRE site [83]. Thus, the hypoxic inductions of APP, BACE 1 and *aph-1* would promote the production and the deposition of A $\beta$  and consequently affect the learning and memory capacities, as observed in AD patients.

### Pre-eclampsia

Pre-eclampsia is a human pregnancy-specific disorder affecting 2–10% of pregnancies. It is a leading cause of prenatal morbidity and mortality for the fetus and the mother, and the only intervention that effectively reverses the syndrome is giving birth. During the early

stages of pregnancy, the cytotrophoblasts play a major role in the placentation process: the differentiation of the proliferative cytotrophoblasts into invasive cytotrophoblasts is absolutely required for this process [84]. In the case of pre-eclampsia, it does not occur and the cytotrophoblasts are blocked in their proliferative state. Consequently, the utero-placental circulation is impaired and promotes a placental ischemia. This defect in adequate vascular network formation can be rescued by the use of HIF1 $\alpha$  antisense [85]. Thus, sustained activation of HIF, by inhibiting cytotrophoblasts differentiation, promotes the development of the disease.

### Ischemia

Several diseases, such as cerebral, cardiac or vascular ischemia, occur when cells do not adapt correctly to a reduced oxygenation [1]. Atherosclerosis leads to a decrease in oxygen tissue perfusion and thus to the formation of a hypoxic area. Although HIF is activated in these hypoxic regions, its activation is not sufficient to restore oxygen homeostasis. In coronary arteries, atherosclerosis lesions cause a cardiac ischemia due to the decreased delivery of oxygen and nutrients to the heart and can lead to a myocardium infarction.

### Cancer

Tumor progression is characterized by an anarchic proliferation of the cancer cells. This proliferation is so major that the tumor is rapidly deprived of oxygen and nutrients, leading to the appearance of hypoxic regions. This process occurs in most solid tumors. At the cellular level, intra-tumoral hypoxia is associated to several phenomena that ensure survival and growth of the cancer cells. Tumor hypoxia promotes a radio- and chemo-resistant environment [86, 87]. Hypoxia stimulates the angiogenic process to increase tumor perfusion, switches tumor cells metabolism, and keeps their proliferative rate constant by increasing the transport of glucose and the glycolytic pathway to maintain a suitable ATP production [22]. Furthermore, hypoxia deregulates the pro- and anti-apoptotic balance to favor cell survival and also increases the invasive properties of cancer cells by up-regulating different genes. In all these processes induced by hypoxia, the HIF transcription factor plays a decisive role.

### HIF $\alpha$ expression in tumors

HIF1 $\alpha$  is overexpressed in many different types of human cancers [88]. Its expression is associated to an aggressive phenotype and is a marker for a poor prognosis for many types of tumors, such as prostate tumor, oropharyngeal, esophageal or head and neck

tumors, lung cancer, ovarian cancer and breast cancer [89, 90].

HIF2 $\alpha$  is more frequently overexpressed in hepatic primary or metastatic cancer and in the macrophages associated to the tumor (tumor-associated macrophages, TAMs) whose invading rate is correlated to the tumor progression and to a poor prognosis [91, 92]. The topography of HIF $\alpha$  subunit expression in tumors is usually focal and mainly restricted around hypoxic/necrotic areas, even if there is not always a strict colocalization of HIF $\alpha$  and the hypoxic regions (Figs. 2, 3). Nevertheless, in certain tumors, such as in clear cell renal carcinoma (RCCs), mutation in gene coding for the pVHL protein leads to a constitutive stabilization of HIF1 $\alpha$  and/or HIF2 $\alpha$  [12]. In paragangliomas with a mutation in the gene for the succinate dehydrogenase kinase, HIF1 $\alpha$  levels are also up-regulated due to the PHD inhibition [93]. More recently, in endometrial tumor, mutation in the *phd2* gene has also been reported and shown to over-activate HIF1 [94].

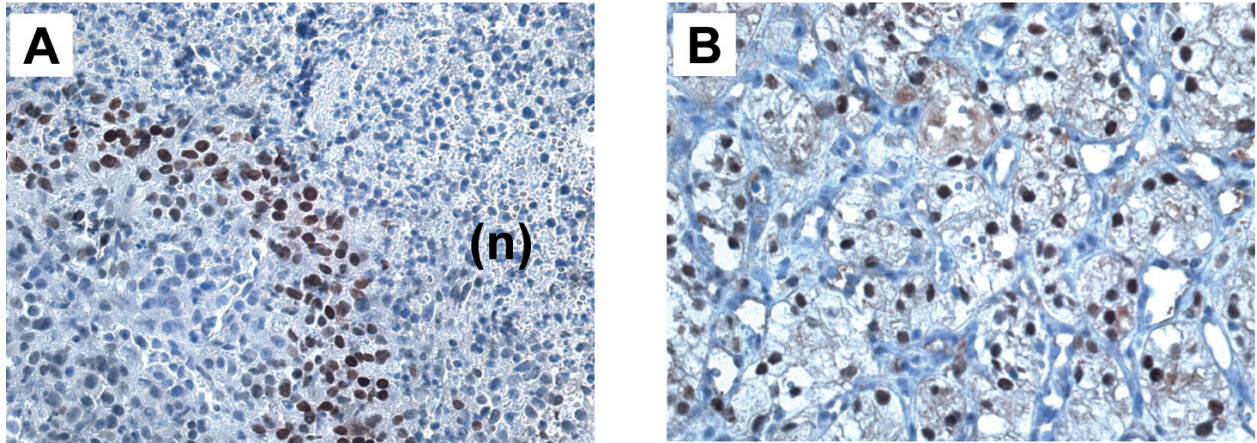
### HIF $\alpha$ contribution to tumor progression

As previously mentioned, hypoxia-induced HIF activation controls the expression of several genes involved in many aspects of cancer progression.

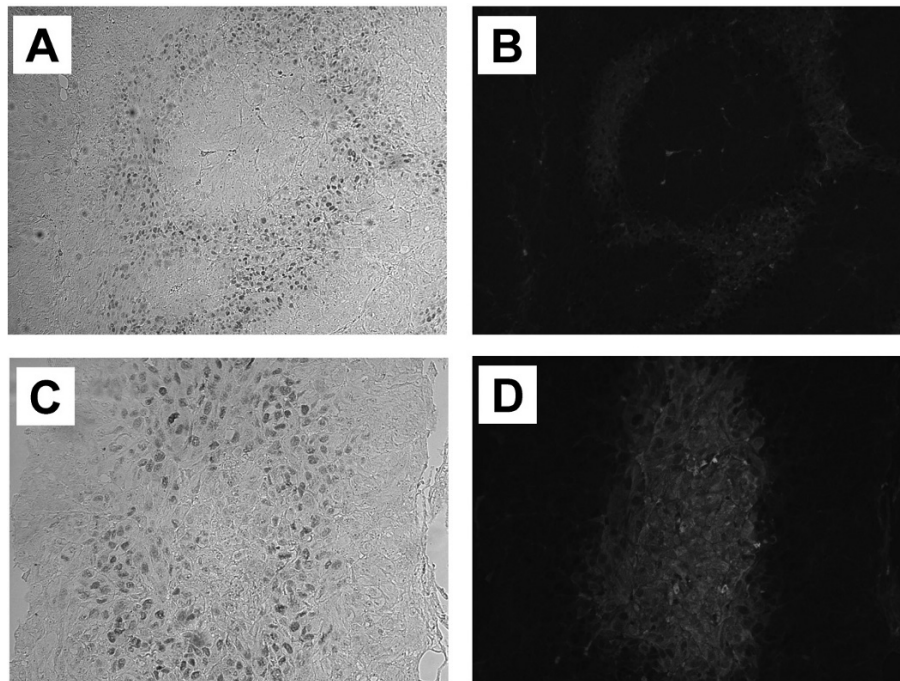
**Tumor angiogenesis.** The angiogenic switch that represents one of the major events in tumor progression is triggered by hypoxia and HIF by inducing the expression of VEGF and its receptors (Flt1, Tie2). These results agree with findings of *in vivo* experiments by Carmeliet et al. [26]: HIF1 $\alpha$ <sup>-/-</sup> ES-derived tumors are smaller and poorly vascularized compared to those from HIF1 $\alpha$ <sup>+/+</sup> ES cells. In contrast, Ryan et al. [28] showed that HIF1 $\alpha$  inactivation affects tumor growth but has no impact on vascular density.

Furthermore, it seems that the role of HIF is strongly dependent on the tumor environment [95]. HIF1 $\alpha$ <sup>-/-</sup> astrocytes injected subcutaneously induce some poorly perfused and very necrotic tumors, whereas the same astrocytes injected into the brain give rise to highly vascularized tumors.

**Tumor invasion.** Several genes that induce extracellular matrix remodeling and basal membrane digestion are regulated by hypoxia and HIF, such as *vimentin*, *fibronectin*, *keratin 14, 18, 19*, *matrix metalloproteinase 2*, *cathepsin D*, and *urokinase plasminogen activator receptor*. In parallel, HIF1 $\alpha$  stabilization triggers the loss of E-cadherin expression and promotes epithelial-mesenchymal transition [96]. Indeed, HIF1 induces the expression of the lysyl oxidase, which regulates the SNAIL transcription factor whose activation represses the E-cadherin expression [97–99]. Moreover, several genes favoring cancer



**Figure 2.** HIF1 $\alpha$  expression in tumors. (A) HIF1 $\alpha$  shows nuclear staining surrounding necrotic areas (n) in a sarcoma xenograft model; (B) HIF1 $\alpha$  is constitutively expressed in a human renal clear cell carcinoma (RCC) because of the inactivating *vhl* mutation.



**Figure 3.** Staining of HIF1 $\alpha$  protein and hypoxic regions in human tumors. Xenograft tumor of human melanoma: HIF1 $\alpha$  staining (A) co-localizes with the hypoxic marker, pimonidazole (B). Xenograft tumor of human breast carcinoma: HIF1 $\alpha$  staining (C) is not perfectly correlated to the hypoxic marker staining, pimonidazole (D).

cells migration, and metastatic properties are also regulated by the activation of the HIF complex: the proto-oncogene *c-MET*, the receptor *CXCR4* and its unique chemokine ligand stromal cell-derived factor1 (*SDF1* also called *CXCL12*).

**Metabolic adaptation.** HIF stimulates the expression of the glucose transporters and the glycolytic enzymes to assure the ATP delivery inside the tumor. Because the anaerobic glycolysis leads to an acidification of the tumoral microenvironment, tumor cells also induce, *via* HIF, the expression of exchangers such as the  $\text{Na}^+/\text{H}^+$  exchanger (*NHE1*) or transporters such as the monocarboxylate transport-

er (*MCT4*) to maintain the pH homeostasis [100, 101]. In addition, tumor cells up-regulate the expression of carbonic anhydrases *CAIX* and *CAXII*, which are linked to the extracellular membrane and catalyze the conversion of  $\text{CO}_2$  in  $\text{HCO}_3^-$  [102, 103]. The bicarbonates are then driven into the cells by the  $\text{Cl}^-/\text{HCO}_3^-$  exchangers and restore the intracellular compartment pH assuring the adaptation of these cells to the acidic environment.

#### Hypoxia and HIF cascade as therapeutic targets

The role of hypoxia and HIF in several pathologies is clearly established, particularly in cancer and cardiovascular diseases. For this reason, their modulation

might be a good strategy for treating hypoxic tumors or ischemia-related pathologies.

## Cancer

In this section, we present an overview of the most relevant approaches established to improve survival based on the targeting of hypoxia and/or the HIF cascade.

### Hypoxia as a therapeutic target in cancer

Tumor hypoxia has been exploited as a way to activate some drug compounds.

**Hypoxia-activated drugs.** Some anti-cancer agents (N-oxides, quinones, nitro-aromatics) have been explored as drugs that exploit the hypoxia itself. The most widely studied of these compounds is tirapazamine [104]. This 'hypoxic cytotoxin' is bioreductively activated in hypoxic cells and potentiates the cytotoxicity of radiation and chemotherapy, in particular that of platinum and taxanes family [105–107]. Tirapazamine has been studied in a Phase III clinical trial in combination with *cis*-platin and paclitaxel for non-small-cell lung cancer [108]. However, the association of this hypoxia-activated drug to the chemotherapy did not improved response rates and survival. These data are in contradiction with another study of head and neck cancer where the adjunction of tirapazamine increases the 3-year failure-free survival rate [109]. Consequently, other evaluations and studies are justified.

In addition, the capacity of anaerobic bacteria to proliferate under hypoxic conditions has been exploited as a tool for hypoxia-targeted anticancer therapy. In this context, the bacteria *Clostridium acetobutylicum* has been modified to overexpress cytosine deaminase, which is able to metabolize the 5-fluorocytosine into the toxic 5-fluorouracile (5-FU). Following the administration of these bacteria to rhabdomyosarcoma-bearing rats, cytosine deaminase is detected in the tumor, suggesting that this original approach might be useful [110].

Therapeutic failure is associated with hypoxic tumors and thus several approaches have been explored to reduce tissue hypoxia. The most obvious strategy to enhance radio- and chemo-sensitivity is the administration of high-pressure oxygen. In addition, the development of sensitizers is another method to potentiate the effect of classical therapies.

**Erythropoietin.** The administration of erythropoietin (Epo, the glycoprotein hormone endogenously secreted by renal fibroblasts under hypoxic conditions

via an HIF-dependent mechanism) is used to increase tumor oxygenation [111]. Recombinant human Epo (rHuEpo) is indeed a crucial therapeutic tool in cancer therapy for patients suffering from anemia. Many studies have shown that correction of anemia, and consequently of hypoxia, is a prognostic factor of disease control and survival. Thus, in preclinical and clinical studies, rHuEpo may improve radio- and chemo-sensitivity of solid tumor possibly by increasing tumoral oxygenation [112, 113].

**Hyperbaric oxygen therapy.** Hyperbaric oxygen (HBO) therapy involves the administration of pure oxygen at a pressure superior than 1 atmosphere. Experimental studies suggest that hypoxia reduces the radio-sensitivity of cells as they require three times more radiation [114]. By increasing the oxygen tension of hypoxic cells, HBO may sensitize radiation therapy. However, it is known that reoxygenation of hypoxic tumor cells may stimulate tumor growth. Nevertheless, Feldmeier's group elegantly concluded, from clinical and preclinical studies, that intermittent HBO therapy has no stimulatory effect on tumor growth and dissemination [115]. Furthermore, intermittent HBO therapy increased the radio-curability of several cancers. In particular, it improved local tumor control and reduced mortality in head and neck cancer [116].

As seen for radiotherapy, HBO therapy is able to sensitize chemotherapy by increasing tumor perfusion and cellular sensitivity. This has been demonstrated *in vitro* with 5-FU, doxorubicin, and taxol [117]. Thus, HBO therapy could be considered as adjuvant therapy in hypoxic tumors.

**ARCON.** Accelerated radiotherapy with carbogen and nicotinamide is another approach that makes it possible to circumvent the mechanisms of radio resistance. In this method, radiotherapy is associated with inhalation of hyperoxic gas to decrease diffusion-limited hypoxia and nicotinamide to decrease perfusion-limited hypoxia. In a breast xenograft model the radiation dose is decreased by 50% with ARCON [118]. However, the results obtained with ARCON in clinical studies have been disappointing. In non-small-cell lung cancer, there were no significant responses with ARCON [119]. Nevertheless, in bladder cancer, ARCON showed significant gains on local control, disease-free survival and overall survival compared with a group control [120].

**Sensitizer agents.** By simulating the action of oxygen, sensitizer agents compensate the low oxygen concentration and increase radiation-induced damage. The compounds most frequently used in clinical studies



**Table 1.** Summary of the most relevant anti-cancer therapeutic approaches targeting the HIF cascade developed so far.

Molecular screenings			
Target	Compound	Mechanism of action	Efficacy
HIF-targeting high-throughout	PX-478	?	Inhibition of tumor growth in animal models
	Topotecan	Topoisomerase-1 inhibitors	Inhibition of tumor growth in animal models
	ARC-111	HIF DNA-binding activity	Inhibition of HIF1 $\alpha$ expression <i>in cellulo</i>
	Echinomycin	HIF / p300 binding	Disappointing results in clinical trials
PHDs	Chetomin	HIF / p300 binding	Inhibition of tumor growth in animal models
	Cyclosporine A R59949	PHDs activators	Inhibition of HIF1 $\alpha$ expression <i>in cellulo</i> (Doubts about future application because of the immunosuppressive effect) Inhibition of HIF1 $\alpha$ expression <i>in cellulo</i>
Others	Rapamycin	PI3K/AKT/mTor pathway inhibitors	Disappointing results in clinical trials
	CCI-779		Used in clinic against Breast cancer
	Trastuzumab	MAPK pathway inhibitors	Inhibition of tumor growth in animal models
	LY294002,		Inhibition of HIF1 $\alpha$ expression <i>in cellulo</i>
	Wortmannin		
	PD98059	?	Inhibition of tumor growth in animal models
	GL331		
	YC-1	Hsp90 inhibitor	Inhibition of tumor growth in animal models
Geldanamycin			
Radicalcol	Microtubules	Inhibition of tumor growth in animal models	
2ME2			
Gene therapy			
Strategies			Efficacy
Antisense therapy			Inhibition of tumor growth in animal models
Gene suicide			
RNA interference			

over the last few years were nitroimidazoles, but these again have had disappointing results. In carcinoma of cervix, Grigsby et al. [121] did not show any difference in a study comparing irradiation with or without misonidazole. On the other hand, in head and neck carcinomas, a randomized study showed encouraging results with nimorazole [122]. Indeed, in association with radiotherapy, nimorazole induced a better loco-regional control and cancer-related survival than the control group. In the same way, a meta-analysis suggested that radio-sensitization with misonidazole for astrocytomas increases 1-year survival by 8% [123].

#### HIF cascade as a therapeutic target in cancer

The inhibition of HIF1 $\alpha$  expression impairs tumor growth when using HIF1 $\alpha$ <sup>-/-</sup>-derived ES cells [26]. Similarly, the expression of a dominant negative mutant of HIF1 (deprived of the two TAD domains) in PCI43 cells leads to the inhibition of tumor growth [124]. Inhibiting the hypoxia signaling cascade as a therapeutic approach to block tumor progression might target different processes: HIF1 $\alpha$  protein stability, nuclear translocation, DNA-binding capacity or even the association with transcriptional repressors and/or co-activators. Several approaches have been

developed using not only conventional strategies (as molecular screenings) but also more innovative strategies, such as RNA interference (Table 1).

**HIF-targeting drug screening.** High-throughout screenings have led to the discovery a certain number of efficient molecules targeting the HIF cascade:

- PX-478 is a drug developed by ProlX Pharmaceuticals. PX-478 suppresses HIF1 $\alpha$  protein expression under normoxic and hypoxic condition, although its precise mechanism of action remains unclear. Moreover, PX-478 showed anti-tumor activity in a variety of xenograft animal models [125]. This effect correlates with a reduction of HIF1 $\alpha$  protein level measured by immunohistochemistry. Furthermore, toxicological studies showed a moderated toxicity. To date, no clinical trial is being carried out with this promising drug.
- Topoisomerase-I inhibitors: Out of the 2000 agents tested by Rapisarda et al. [126], 4 were identified as HIF1 inhibitors. Three of these compounds are related to camptothecin, which inhibits topoisomerase-I activity. The best characterized in clinical study is NSC-609699 (topotecan). In a glioma xenograft model, topotecan inhibits tumor growth,

with a concomitant and dramatic reduction of the protein levels of HIF1 $\alpha$  and HIF1-target gene products [127]. Recently, another topoisomerase I-inhibitor (ARC-111) has been tested *in cellulo*, showing inhibition of HIF1 $\alpha$  expression upon hypoxia [128]. However, no clinical or preclinical studies are available at this moment using these drugs.

- Echinomycin is an antibiotic derived from quinoxaline family, which inhibits HIF1 $\alpha$  DNA-binding activity [129]. This is not the only effect of this small molecule. This drug has been evaluated in numerous preclinical and clinical trials; however, with disappointing results [130, 131].
- Chetomin is small molecule that is able to disrupt HIF1 binding to the p300 co-activator. *In vivo*, this transcriptional inhibition reduces tumor growth by increasing the number of necrotic regions in treated xenograft tumors [132]. These effects seem to be specific for HIF1, since the expression of a p300-independent form of HIF1 $\alpha$  in tumor cells rescues, at least partially, the phenotype induced by chetomin. This molecule is on standby for clinical evaluation.

**PHDs activators.** PHDs are clearly implicated in the regulation of HIF1 $\alpha$  protein stability. The activation of these PHDs could be an interesting anti-HIF therapeutic strategy:

- Cyclosporin A is an immunosuppressive agent used in organ transplantation. Cyclosporin A also activates PHD activity [133], through this mechanism this drug inhibits hypoxia-induced HIF1 $\alpha$  stabilization and HIF1-mediated cellular responses in glioma cells. However, these results observed *in cellulo* cannot be extrapolated to the *in vivo* situation. Indeed, because of its immunosuppressive functions, cyclosporin should not be used in cancer therapy where immunity is crucial.
- R59949 is a PHD activator. This agent is an inhibitor of diacylglycerol kinase [134]. *In cellulo*, R59949 inhibits HIF1 $\alpha$  protein accumulation by stimulating PHD activity. This molecule offers a new possibility against cancer but has not yet been tested *in vivo*.

#### Indirect inhibitors of the HIF complex

**Inhibitors of the PI3K/Akt/mTor cascade.** Drugs interfering with the PI3K/Akt/mTor pathway can block the expression of HIF1 $\alpha$  protein and HIF1-dependent gene products. Rapamycin inhibits HIF1 $\alpha$  expression *in cellulo* [135]. An ester analogue of

rapamycin, CCI-779, inhibits tumor growth in a rhabdomyosarcoma xenograft model [136]. This inhibition is related to the reduction of HIF1 $\alpha$  and VEGF expression. CCI-779 has, however, shown contradictory results in numerous clinical studies [137, 138]. Other example is trastuzumab (herceptin). This humanized monoclonal antibody targets the human EGF receptor-2 (HER2) [139]. HER2 induces the expression of HIF1 $\alpha$  and VEGF in breast cancer cells *via* the PI3K/AKT pathway [140]. Treatment of breast cancer cells with the HER2 inhibitor induces a dose-dependent inhibition of EGF levels but the impact on HIF1 $\alpha$  still remains controversial [141]. LY294002 and wortmannin represent other tested agents that reduce HIF1 $\alpha$  expression *in cellulo* [142]. *In vivo* and in clinical studies, these drugs reduce tumor growth but the impact on the HIF pathway has not yet been elucidated [143].

**Inhibitors of the MAPK cascade.** Some inhibitors have been developed, such as PD98059 that inhibits the transactivation ability of HIF1 $\alpha$  but does not change the stabilization or DNA binding ability of HIF1 $\alpha$  [144]. GL331 is a plant-derived MAPK inhibitor (the podophylotoxin). GL331 is known to block the ERK pathway and inhibit HIF1 $\alpha$  expression in human lung cancer cells [145]. Nevertheless, there are no reported data on its use as an anticancer drug.

**YC-1.** YC-1 [3-(5'-hydroxymethyl-2'-furyl)-1-benzylindazole] is another molecule that can inhibit HIF1 $\alpha$ . Initially developed for circulatory disorders, YC-1 inhibits platelet aggregation and vascular contraction by activating soluble guanylyl cyclase [146]. The precise mechanism of HIF1 $\alpha$  inhibition is not clear: NF- $\kappa$ B and Mdm2 may contribute to this inhibition. *In cellulo*, YC-1 inhibits HIF-1 $\alpha$  accumulation at the post-translational level in hepatoma cells upon hypoxia [147]. *In vivo*, YC-1 reduces the tumor growth in five independent xenograft models, and inhibits the expression of HIF1 $\alpha$  and HIF1-regulated genes [148]. Given that no serious toxicity was highlighted in mice, this drug could be a good candidate for a Phase I study.

**Heat-shock protein 90 (Hsp90) inhibitors.** Hsp90 is a chaperone protein required for the stability and/or maturation of numerous proteins, including key mediators of signal transduction and cell cycle control. Among these proteins, Hsp90 regulates HIF1 $\alpha$  activation [149]. Moreover, *in vivo*, geldanamycin (an Hsp90 inhibitor) induces HIF1 $\alpha$  protein degradation under normoxia or hypoxia [150]. In the same manner, radicicol is another Hsp90 inhibitor that inhibits DNA binding of the HIF1 complex and reduces hypoxia-

induced VEGF expression [151]. These compounds have shown preclinical efficacy in xenograft model and clinical trials [152]. Nevertheless, up to date, there is no reported data related to their impact on the HIF pathway.

**Microtubule-disrupting agents (MDAs).** Another interesting group of cytotoxic agents that target HIF1 $\alpha$  are those that induce disruption of microtubule architecture. Among these drugs, 2-methoxyestradiol (2ME2) is an endogenous metabolite of estrogen that has an anti-angiogenic and anti-proliferative activity [153]. Recent data have shown that this anti-angiogenic effect is mediated through the inhibition of HIF1 $\alpha$  expression in endothelial and cancer cells. Other agents that stabilize (taxol) or destabilize (vincristine) microtubules cause a similar effect. 2ME2 reduces tumor growth and angiogenesis *in vivo*, but the impact on HIF1 $\alpha$  expression is still unclear [154].

These data are, however, in contradiction with the results reported by Jung et al. [155] showing that certain MDAs (vinblastine, colchicine or nocodazole) increase HIF1 activation through an NF- $\kappa$ B-dependent pathway. Further studies are necessary to clarify this point and validate the potential interest of these compounds.

### Gene therapy

**Antisense therapy.** Several groups have evaluated the inhibition of HIF activity by antisense therapy. Sun et al. [156] were the first to show the efficacy of this treatment in a mouse lymphoma model. Intratumoral injection of a plasmid vector containing an HIF1 $\alpha$  antisense led to the regression of small tumors (<1 mm) and slowed the growth of large tumors (>4 mm). This effect correlated to the inhibition of HIF1 $\alpha$  and VEGF expression. The same team showed an excellent anti-tumor efficacy of the HIF1 $\alpha$  antisense therapy in combination with pVHL overexpression in a glioma model [157]. Antisense therapy used in pancreatic cancer xenograft models similarly reduced tumor size and weight [158]. Accordingly, gene transfer techniques using antisense plasmids may provide future anti-cancer therapy but, to date, this anti-HIF1 $\alpha$  strategy has not been tested in clinical trials.

**Suicide gene therapy.** This strategy is based on the administration into the tumor cells of a gene encoding for an enzyme that is able to transform a non-toxic pro-drug into a cytotoxic agent. HRE-controlled gene suicide therapy has shown interesting data *in cellulo* and *in vivo*. For example, an adenoviral vector under the control of an optimized HRE promoter has been

used to express the gene encoding the human cytochrome P450 (CYP2B6) and shown to delay tumor growth upon treatment with cyclophosphamide [159]. Several teams have used similar strategies but, for the moment, no clinical data are available.

**RNA interference.** RNA interference (RNAi) is an alternative strategy to inhibit the HIF1 signaling pathway. This innovative technology is particularly interesting since RNAi triggers sequence-specific gene silencing. This approach is well controlled *in cellulo* but the conditions of administration are not yet well established *in vivo*. Mizuno et al. [160] reported the impact of HIF1 $\alpha$  silencing on a xenograft model of pancreatic and hepatobiliary carcinoma cells stably transfected with a plasmid producing siHIF1 $\alpha$ , and showed that this silencing significantly reduced tumor growth. More recently, Gillepsie et al. [161] showed, in a mouse model of glioma, the reduction of tumor growth and HIF1 activity following intratumoral injection of siHIF1 $\alpha$ . All these results are very promising and suggest the efficacy of this strategy but, to date, no systemic administration has been evaluated.

### Ischemic cardiovascular diseases

Many angiogenic factors (VEGF, FGF, PDGF) have been tested in preclinical studies, but in clinical practice the results are either modest or disappointing. Activating HIF, which induces angiogenesis *via* the expression of many angiogenic factors, might be more effective than targeting a single factor. Thus, the development of HIF stimulants may be helpful for the treatment of ischemic disease. However, all the strategies that improve systemic angiogenesis might induce side effects in patients with cancer, angioma, arthritis, retinopathies or atherosclerotic plaque progression. Consequently, we have to be very careful with the use of such agents. Two main strategies have been developed to activate HIF by either inhibiting the endogenous degradation pathway or by overexpressing the protein.

### Inhibiting HIF degradation as a therapeutic approach in ischemia

**PR39.** This molecule was identified as a macrophage-derived peptide that induces HIF1 $\alpha$  by inhibiting the proteasome [162]. In a pig model of chronic myocardial ischemia, the intracardial delivery of an adenoviral construction (Ad)-PR39 improves myocardial blood flow and collateral formation by a mechanism implicating HIF1 $\alpha$  overexpression and induction of VEGF, VEGFR-1 and 2, syndecan and FGFR-1 [163].

**Dibenzoylmethane.** This product is a natural dietary compound and iron chelator, and inhibits HIF1 $\alpha$  degradation and induces VEGF secretion under normoxia [164]. This agent might be of interest in ischemic disease but, to date, no clinical or preclinical data are available.

**siPHDs.** Another innovative strategy is the RNAi to silence PHD2 to increase HIF1 transcriptional activity. Indeed, intraperitoneal injection of siPHD2 in a mouse model of myocardial ischemia reperfusion injury induces up-regulation of cardiac iNOS expression [165]. This is translated clinically by a higher ventricular function and a lower infarct size than seen in the control group.

**DMOG.** Dimethylxalylglycine (DMOG) is a small molecule that inhibits the oxoglutarate-dependent dioxygenase and thus induces HIF1 $\alpha$  stabilization. In a mouse model of hind limb ischemia, administration of DMOG increased VEGF levels and capillary density [166]. Other PHDs inhibitors have shown similar results: P4-H, L-Mim, 3,4-DHB and S956711 [167].

#### Ectopic expression of HIF as a therapeutic approach in ischemia

A constitutively active form of HIF1 $\alpha$  has been constructed associating the DNA-binding and dimerization domains from HIF1 $\alpha$  and the transactivation domain from herpes simplex virus VP16 protein (HIF-1 $\alpha$ /VP16). In preclinical studies, the administration of this strong constitutive transcriptional activator improved angiogenesis in a rabbit hind limb ischemia model (increasing the blood flow and the collateral vessel development) and in a rat myocardial infarction (reduction of the infarct size and increasing capillary density) [168].

Another recombinant replication-deficient adenovirus Ad2/HIF-1 $\alpha$ /VP16 has been evaluated in a Phase I for critical limb ischemia patients with promising results and particularly with no major adverse effect [169].

**AdCA5** is another replication-defective recombinant adenovirus encoding a constitutive form of HIF1 $\alpha$ . In preclinical studies, ocular injection of AdCA5 in mice induced neovascularization of retinal vessels *via* the expression of multiple angiogenic factors [170]. In the same way, this construction injected in a rabbit model of limb ischemia stimulates the recovery of limb perfusion by increasing capillary density and luminal area of arteries [171].

All these results support our particular interest in the understanding of the molecular mechanisms underlying the hypoxia-signaling cascade to improve our

basic knowledge and also to identify new prognostic and/or predictive clinical markers, as well as therapeutic targets against the pathologies in which hypoxia is implicated.

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