

HIF-mediated innate immune responses: cell signaling and therapeutic implications

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Abstract: Leukocytes recruited to infected, damaged, or inflamed tissues during an immune response must adapt to oxygen levels much lower than those in the circulation. Hypoxia inducible factors (HIFs) are key mediators of cellular responses to hypoxia and, as in other cell types, HIFs are critical for the upregulation of glycolysis, which enables innate immune cells to produce adenosine triphosphate anaerobically. An increasing body of evidence demonstrates that hypoxia also regulates many other innate immunological functions, including cell migration, apoptosis, phagocytosis of pathogens, antigen presentation and production of cytokines, chemokines, and angiogenic and antimicrobial factors. Many of these functions are mediated by HIFs, which are not only stabilized posttranslationally by hypoxia, but also transcriptionally upregulated by inflammatory signals. Here, we review the role of HIFs in the responses of innate immune cells to hypoxia, both *in vitro* and *in vivo*, with a particular focus on myeloid cells, on which the majority of studies have so far been carried out.

Keywords: hypoxia, neutrophils, monocytes, macrophages

Introduction

Innate immune cells include both myeloid cells, such as neutrophils and circulating monocytes which differentiate into macrophages within tissues, and innate lymphoid cells (ILCs). These cells have important functions in host defense, including the phagocytosis of pathogens, the release of antimicrobial proteases, and the secretion of cytokines and chemokines that attract and activate other immune cells. Cells of the immune system encounter a range of different oxygen tensions, with lower levels in tissues than in blood, which are further reduced during inflammation. To function under hypoxic conditions, innate immune cells increase anaerobic adenosine triphosphate (ATP) generation by upregulating glycolytic enzymes and glucose transporters.^{1–5} Hypoxia also, however, has a plethora of additional effects on innate immune responses.

Low oxygen levels act on vascular endothelial cells to upregulate adhesion molecules and chemokines, attracting circulating monocytes and neutrophils.^{6–11} Neutrophils and monocytes upregulate adhesion molecules, such as intercellular adhesion molecule 1 (ICAM-1) and CD18/CD11b as they reach hypoxic regions, promoting their arrest and transendothelial or transepithelial migration toward the site of the inflammation.^{2,12} Hypoxia also acts to retain myeloid cells at the site of injury by upregulating factors that inhibit migration.^{2,3,5,13–16}

Hypoxia tends to potentiate inflammatory functions and has a prosurvival effect in neutrophils, monocytes, and eosinophils.^{4–5,15,17,18} The ability of innate immune cells to

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survive and function effectively under hypoxic conditions is crucial for host defense in tissues. Immune cell apoptosis is, however, an important means of resolving inflammation; prolonged survival and activation of myeloid cells under hypoxic conditions is likely to play a major role in chronic inflammation, contributing to diseases such as chronic obstructive pulmonary disease, asthma, and rheumatoid arthritis (RA).^{19–23}

Lack of oxygen and a reduction in oxidative phosphorylation, in favor of less efficient glycolysis, might be expected to reduce ATP availability and impair cellular functions; however, there is evidence that cardiac hypoxia inducible factor (HIF) expression can increase ATP availability despite inducing a glycolytic switch.²⁴ This may also be the case in myeloid cells, with hypoxia enhancing many effector functions. Glycolysis is critical for ATP production by myeloid and endothelial cells even under normoxic conditions.^{25,26} Inhibition of glycolysis reduces endothelial cell proliferation and impairs formation of lamellipodia, causing a defect in directional migration and vessel formation.²⁶ Potentially, therefore, glycolysis (and its upregulation by hypoxia) may promote migration of myeloid cells.

Hypoxia upregulates Fc and complement receptors on macrophages and neutrophils, thereby enhancing phagocytosis.^{27–32} It also promotes the release of antimicrobial mediators, such as β -defensin by macrophages and elastase by neutrophils.^{33–35} Monocytes and macrophages upregulate proinflammatory cytokines, such as tumor necrosis factor α (TNF α), interferon γ (IFN γ), interleukin (IL)-1 β , and several neutrophil chemoattractants in hypoxia, while downregulating chemokines associated with adaptive immunity.^{2,3,31,36–38}

Macrophages precultured in hypoxia are more effective at inducing ovalbumin-specific CD8 T-cell responses, due to the upregulation of major histocompatibility complex (MHC) class I and costimulatory molecules, but they also produce arginase, which inhibits CD4 T-cell responses.^{31,39,40}

Many effects of hypoxia are mediated by release of HIFs from suppression by the oxygen-dependent hydroxylases. These comprise prolyl hydroxylases (PHDs), which target them for degradation, and the asparaginyl hydroxylase, factor inhibiting HIF, which inhibits HIF transcriptional activity. Many immune-related genes are regulated by HIF, either directly, via binding of HIF to a HIF-responsive element (HRE) in the promoter region, or indirectly, via HIF-mediated induction of other signaling molecules and transcription factors, such as NF- κ B. Effects of HIF signaling in innate immune cells in response to hypoxia and other stimuli are summarized in Figure 1.

Other mechanisms also play a role in hypoxia signaling; for example, mitochondrial-derived reactive oxygen species stabilize HIFs and also activate other transcription factors, such as activator protein-1 (AP-1), which induces CXC-chemokine ligand (CXCL) 8 in hypoxic macrophages.^{37,38,41–43} Hypoxia also reduces activity of the Jumonji histone demethylases, resulting in increased methylation of histone H3 residues in the promoter regions of chemokine ligand 2 (CCL2), chemokine (C-C motif) receptor 1 (CCR1), and CCR5, repressing their transcription by macrophages.⁴⁴

Though research so far has focused predominantly on myeloid cells, there is also some evidence of the effects of hypoxia on other aspects of innate immunity. For instance, mice with chronic hypoxia-induced pulmonary hypertension have reduced natural killer (NK) cell numbers, cytotoxicity, and IFN γ production.⁴⁵ Hypoxic NK cells downregulate the key cytotoxic mediators granzyme B and perforin and lose their ability to upregulate activating receptors in response to cytokines, thereby reducing their ability to recognize and kill infected or abnormal cells.^{46,47}

In addition to the effects of hypoxia on innate immune responses to infection, hypoxia can also have direct effects on the pathogens themselves, such as modulating the expression of virulence factors,⁴⁸ but this is beyond the scope of this review.

Roles of HIFs in neutrophil responses to hypoxia

Neutrophils are inherently adapted to low oxygen tensions, with few mitochondria and reliance on glycolysis for much of their energy demand even under normoxic conditions.²⁵

While *HIF1A* messenger RNA (mRNA) is present in freshly isolated neutrophils, HIF-1 α protein is not expressed in unstimulated cells. It can be induced by hypoxic culture, the iron chelator desferrioxamine, or the competitive hydroxylase inhibitor dimethylxalylglycine (DMOG).^{5,49} When cultured in hypoxia, neutrophils upregulate the glycolytic HIF-1 α target genes glyceraldehyde 3-phosphate dehydrogenase (*GAPDH*) and phosphoglycerate kinase (*PGK*), which may serve to provide additional glycolytic capacity for enhanced effector function.⁵ Hypoxic induction of *Pgk* expression is reduced but not abolished in *Hif1a*^{-/-} neutrophils, suggesting that both HIF-1 α and other factors (possibly HIF-2 α) are involved in its induction. ATP levels are substantially reduced in *Hif1a*^{-/-} neutrophils and macrophages, reflecting the central role of HIF-1 α in regulation of metabolic activity.⁵⁰

Expression of the cathelicidin-related antimicrobial peptide (CRAMP) is upregulated by hypoxia and in

neutrophils lacking the von Hippel-Lindau protein (VHL) (involved in the proteasomal degradation of HIF), whereas *Hif1a*^{-/-} neutrophils express lower levels of CRAMP and display reduced ability to breakdown artificial substrates for antimicrobial proteases.⁵¹ HIF-1 α also promotes extracellular bacterial killing by inducing formation of neutrophil extracellular traps.⁵² Furthermore, neutrophil hypoxic survival is mediated by HIF, with *Hif1a*^{-/-} neutrophils displaying a failure of hypoxia-mediated apoptosis suppression.^{5,49} In wild-type (WT), but not *Hif1a*^{-/-} neutrophils, hypoxia reverses the decline in nuclear factor (NF)- κ B and I κ B kinase (IKK)- α levels observed during culture and maintains DNA-binding activity of NF- κ B family members, which is essential for hypoxic survival.

HIF-2 α is also expressed by human and murine neutrophils and its expression is upregulated by hydroxylase inhibitors, bacteria or lipopolysaccharide (LPS).⁵³ The respiratory burst, phagocytosis, chemotaxis, LPS-induced receptor upregulation and constitutive apoptosis are unaffected by deletion of *Hif2a*. However, *Hif2a*^{-/-} inflammatory neutrophils derived from bronchoalveolar lavage (BAL) fluid of LPS-challenged mice express lower levels of the antioxidant enzyme catalase than WT neutrophils and display increased apoptosis in response to nitrosative stress.

While circulating leukocytes that are attracted to a site of inflammation initially upregulate HIF in response to hypoxia, cells exposed to chronic hypoxia may respond differently due to the activation of negative feedback mechanisms. For instance, hypoxia is reported to induce HIF-mediated transcription of *PHD2*, *PHD3*, and *VHL* – all negative regulators of HIF expression.^{54–57}

Although PHDs are generally understood to act upstream of HIFs, promoting their degradation, in neutrophils PHD3 may function downstream of HIF. PHD3 is upregulated during hypoxia; this is significantly impaired in *Hif1a*^{-/-} neutrophils.⁵⁴ PHD3 does not, however, appear to regulate HIF-1 α expression, since the hypoxic upregulation of HIF target genes is not affected by the deletion of *Phd3*. *Phd3*^{-/-} neutrophils fail to upregulate the anti-apoptotic protein B-cell lymphoma extra-large (Bcl-X_L), as occurs in hypoxic WT neutrophils, and exhibit reduced survival during hypoxia.

Roles of HIFs in monocyte and macrophage responses to hypoxia

Hypoxic macrophages upregulate HIF-1 α and HIF-2 α proteins in a phosphatidylinositol 3-kinase (PI3K)/protein kinase B (AKT)-dependent manner.^{3,58} In contrast to neutrophils, macrophages from *Phd3*^{-/-} mice exhibit higher

expression of HIF-1 α and NF- κ B than WT, produce more cytokines, and exhibit enhanced chemotactic and phagocytic responses, suggesting that in macrophages, PHD3 is acting upstream of HIF-1 α to degrade HIF-1 α protein.⁵⁹

As in neutrophils, HIF-1 α has an important role in regulating macrophage metabolism under hypoxic conditions. *HIF1A*-null or small interfering RNA (siRNA)-treated macrophages have reduced basal expression and hypoxic induction of the glycolytic genes *PGK* and *GLUT1*, whereas the loss of VHL mimics hypoxic induction.^{3,50} Release of lactate, a marker of glycolytic activity and inflammatory activation, is significantly lower in *Hif1a*^{-/-} macrophages and significantly higher in *Vhl*^{-/-} cells.⁵⁰ HIF signaling is, in turn, influenced by metabolism. For instance, the tricarboxylic acid cycle intermediate, succinate, which is generated during aerobic respiration and upregulated in response to LPS stimulation, inhibits PHD activity, resulting in HIF-1 α stabilization and *Iilb* transcription in macrophages.⁶⁰

Hypoxic upregulation of the adhesion molecule ICAM-1 is inhibited by siRNA targeted against *HIF1A* or *HIF2A*, while induction of β -integrins in monocytes is abrogated by mutation of the HRE in the promoter of CD18, the subunit common to the four known β 2 integrin heterodimers.^{3,12} *Hif1a*^{-/-} macrophages display defects in chemotaxis, in their ability to penetrate the extracellular matrix, and in homotypic adhesion, a process of self-aggregation that amplifies the tissue recruitment of leukocytes.^{50,61} *Hif2a*^{-/-} macrophages have reduced expression of the chemokine receptors macrophage colony-stimulating factor receptor and CXC chemokine receptor 4 and the adhesion molecule fibronectin-1, resulting in a 50% reduction in migratory and invasive capacities.⁶²

Phagocytosis and intracellular bactericidal activity are also regulated by HIFs. Hypoxia induces p38 in macrophages and p38 inhibition reverses HIF-1 α stabilization and enhancement of phagocytosis.³² HIF-1 α induces thrombospondin-1 and the scavenger receptor CD36, which are involved in the recognition of apoptotic cells, therefore hypoxia enhances phagocytosis of both bacteria and apoptotic neutrophils – effects that are abrogated in the presence of *HIF1A*-directed siRNA.^{32,63} Killing of intracellular bacteria, such as streptococci and *Pseudomonas aeruginosa*, is enhanced by iron chelators and in *Vhl*^{-/-} macrophages, but impaired in *Hif1a*^{-/-} macrophages, possibly due to HIF-1 α -mediated inducible nitric oxide synthase (iNOS) induction.^{50,51,61}

Hypoxia also induces the expression of IFN γ , which promotes phagocytosis of opsonized cells by upregulating surface levels of the CD11b/CD18 complement receptor and the CD16/CD32 Fc γ receptor.³¹ The *Ifng* promoter

contains a functional HRE and hypoxic induction of IFN γ and the associated upregulation of opsonic receptors are abolished by mutation of this site or by deletion or siRNA knockdown of *Hif1a*. *Hif1a*^{-/-} macrophages secrete less TNF- α in response to LPS or group A streptococci (GAS) and fail to upregulate TNF- α when LPS is administered under hypoxic conditions.^{50,51} HIF-1 α also binds to a highly polymorphic region of the promoter encoding solute carrier family member 1 (Slc11a1), a phagocyte-specific solute carrier that induces expression of cytokines, chemokines, and MHC class II molecules.⁶⁴ Allelic variation affects the ability of HIF-1 α to induce *SLC11A1* expression and has been associated with susceptibility to various inflammatory diseases. HIF-2 α also appears to be involved in cytokine and chemokine transcription, since the hypoxic upregulation of *Il1b*, *Il12*, *Cxcl2* and *Cxcl6* is impaired in *Hif2a*^{-/-} macrophages while hypoxic expression of *CXCL8* is reduced in macrophages treated with siRNA targeting *Hif2A*.^{3,62}

In contrast to its many proinflammatory effects, hypoxia also promotes repair mechanisms and macrophage suppression of T-cells in a HIF-dependent manner.⁶⁵ Induction of the proangiogenic factors vascular endothelial growth factor (VEGF) and adrenomedullin is significantly reduced by siRNA targeting either *HIF1A* or *HIF2A*.³ *Hif1a*^{-/-} macrophages have reduced basal expression and hypoxic induction of VEGF, whereas deletion of VHL induces VEGF expression under normoxic conditions.^{3,50} Plasminogen activator inhibitor-1 (PAI-1), which promotes thrombosis by inhibiting plasminogen-mediated fibrinolysis, is also activated by HIF-1 α .⁶⁶

HIFs can also act as negative regulators of gene expression. For instance, HIF-1 α (but not HIF-2 α) binds to the HRE of the histone deacetylase-2 (*HDAC2*) promoter independently of the HIF-1 β subunit and reduces transcription by inhibiting binding of Polymerase II.⁶⁷ Hypoxia, therefore, reduces mRNA and protein levels of HDAC2, a suppressor of NF- κ B-mediated transcription, resulting in increased inflammation and corticosteroid resistance.

Hif1a mRNA has a shorter half-life than *Hif2a* mRNA and, following intrapulmonary LPS challenge, *Hif1a* is rapidly upregulated but declines after 24 hours, whereas *Hif2a* expression is delayed but maintained during the resolution phase.^{53,68} HIF-1 α and HIF-2 α bind to the same promoter sequence and the expression of many hypoxia-inducible genes, including *VEGF*, *IL1B*, *CXCL8*, *GLUT1*, *ADM* (adrenomedullin), *CXCR4*, and *STAT4*, can be impaired by siRNA targeted against either *HIF1A* or *HIF2A*.³ Some target genes, however, such as the adenosine A2a receptor (*ADORA2A*), are only

regulated by one HIF and not the other, which is thought to be due to interaction with different coactivators.^{3,69}

Roles of HIF-1 α and HIF-2 α in macrophage polarization

Although both HIF-1 α and HIF-2 α are induced by hypoxia, other stimuli such as cytokines also influence their relative expression levels. For instance, Takeda et al⁶⁸ reported that HIF1- α mRNA and protein are induced during M1 polarization of macrophages (by IFN γ or LPS), whereas HIF-2 α is induced during M2 polarization (by IL-4 or IL-13) and is decreased by stimulation with LPS or IFN γ . *Hif1a*^{-/-} macrophages infiltrating tumor spheroids in vitro develop a more M2-like profile than their WT counterparts, with lower cytotoxicity and decreased production of TNF α , iNOS, and IL-6, suggesting that HIF-1 α is important for proinflammatory responses.⁷⁰ The work of Imtiyaz et al⁶² indicates that HIF-2 α also has important proinflammatory functions, since myeloid-specific *Hif2a* deletion inhibits in vitro secretion of the proinflammatory cytokines IL-1 β , IL-6, and IL-12 in response to a combined IFN γ /LPS stimulus and reduces serum levels of IL-1 β , IL-12, TNF α , and IFN γ following in vivo LPS stimulation.

The relative levels of HIF-1 α and HIF-2 α may play an important role in the regulation of nitric oxide (NO) metabolism by controlling the levels of both iNOS and arginase (which competes with iNOS for the L-arginine substrate required for NO production). *Hif2a*^{-/-} macrophages exhibit normal hypoxic induction of iNOS (*Nos2*); however, both the basal expression and hypoxic induction of arginase 1 (*Arg1*) are attenuated.^{62,68} Conversely, *Hif1a* deletion dramatically reduces hypoxic iNOS induction, but it has a smaller effect on arginase expression.⁶⁸ Similar data was obtained by Kobayashi et al,⁷¹ who observed an increased iNOS response to LPS in cells with a *Vhl/Hif2a* double knockout (in which HIF-1 α is stabilized while HIF-2 α is absent), while *Vhl/Hif1a* double knockout mice produced more arginase in response to IL-13 stimulation. In accordance with the altered ratio of iNOS/arginase, mice lacking myeloid HIF-1 α exhibited reduced NO production 6 hours after LPS treatment, while those lacking HIF-2 α exhibited increased NO after 24 hours.⁶⁸ Takeda et al⁶⁸ also demonstrated decreased and increased levels of in vitro NO production in IFN γ -stimulated *Hif1a*^{-/-} and *Hif2a*^{-/-} thioglycollate-elicited peritoneal macrophages respectively. However, when Imtiyaz et al⁶² stimulated bone marrow-derived macrophages (BMDMs) with a combination of IFN γ and LPS, no difference was observed in NO production between *Hif2a*^{-/-} and WT. The discrepancy between the

studies could be explained by many factors, including the sourcing of macrophages from different tissues, use of a different stimulus, and the apparent dependency of the *Hif2a* phenotype on the level of L-arginine availability.⁶⁸

Mice with a myeloid-specific *Phd2* haploinsufficiency exhibit greater infiltration of mannose receptor, C type 1 (MRC1)⁺ proangiogenic/wound-healing M2-like macrophages, which release arteriogenic factors, following femoral artery ligation.⁷² In comparison to WT macrophages, *Phd2*^{+/-} macrophages isolated at baseline express higher levels of M2 genes, such as arginase (*Arg1*), *Cxcr4*, platelet-derived growth factor beta (*Pdgfb*), neuropilin 1 (*Nrp1*), found in inflammatory zone 1 (*Fizz1*) and transforming growth factor beta (*Tgfb*) and lower levels of the M1 genes *Il12*, *Nos2*, *Il1b*, *Il6*, and *Tnfa*. However, following ligation, WT macrophages from the ischemic musculature upregulate M2 genes, reaching a similar level to that of *Phd2*^{+/-} macrophages, indicating hypoxia is releasing M2 genes from PHD2-mediated suppression. Given the previous work indicating an association of HIF-1 α with iNOS and HIF-2 α with arginase expression,⁶⁸ this data might suggest that PHD2 expression favors an M1 response by preferentially targeting HIF-2 α for destruction.

Modulation of HIF signaling by infection and inflammatory stimuli

While hypoxia regulates HIFs largely at the posttranslational level, other factors such as cytokines and infectious agents regulate its transcription, resulting in a synergistic effect when the stimuli are combined.⁷³⁻⁷⁸

The *HIF1A* promoter contains a binding site for NF- κ B transcription factors.⁷⁷ Both the basal transcription and induction of *HIF1A* (but not *HIF2A*) in response to stimuli, such as bacteria, TNF- α , or hypoxia, are dependent on NF- κ B. Reduced HIF-1 α accumulation and target gene induction are observed in IKK β -deficient macrophages and in cells treated with siRNA targeting members of the NF- κ B family.^{76,77} Hypoxia, in turn, augments NF- κ B expression, phosphorylation, and target gene induction in macrophages and neutrophils.^{3,5,79-81} Hypoxic induction of mRNA encoding IKK α , IKK β , and p65 is impaired in *Hif1 α* ^{-/-} cells, while the increase in p65 phosphorylation is ablated by the deletion of *Hif2a*.^{3,5} IKK β and NF- κ B1 are also directly regulated by PHDs and factor inhibiting HIF respectively.^{80,82} HIF-1 α also binds to the *Tlr4* promoter and upregulates TLR4 expression, thereby increasing the macrophage response to LPS.⁸³

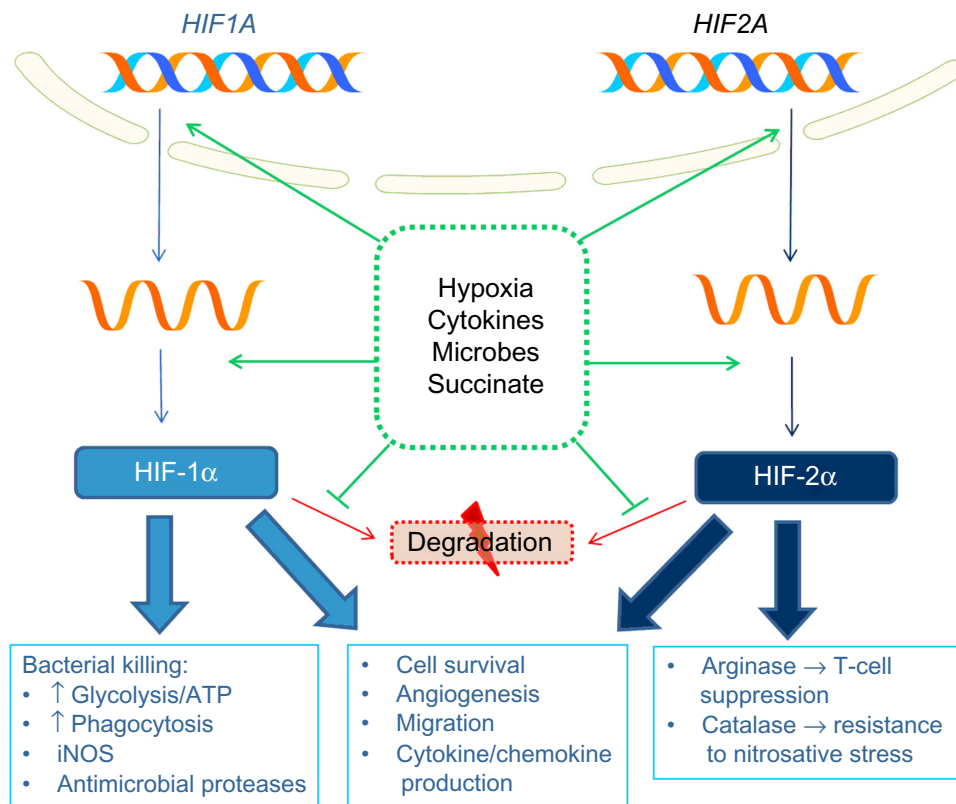


Figure 1 Role of HIFs in innate immune responses to hypoxia.

Abbreviations: HIF, hypoxia inducible factor; ATP, adenosine triphosphate; iNOS, inducible nitric oxide synthase.

In addition to activating NF- κ B-mediated transcription, LPS induces a mammalian target of rapamycin-dependent increase in HIF-1 α translation in neutrophils and promotes succinate-mediated HIF-1 α stabilization in macrophages, as detailed previously.^{52,60} Oxygen consumption by bacteria, resulting in cellular hypoxia, has also been proposed as a mechanism for the upregulation of HIFs during infection.^{84,85}

In vivo models

Immune responses *in vivo* may be influenced by HIF expression in both leukocytes and tissue cells, and systemic modulation of HIF expression affects both. Myeloid-specific knockouts of HIF signaling components, however, have permitted elucidation of the innate immune cell-intrinsic effects of hypoxia and HIFs.

Administration of the hydroxylase inhibitor DMOG delays inflammatory resolution following tail transection in zebrafish larvae, due to decreased apoptosis of neutrophils and their retention at the site of injury during the resolution phase.⁸⁶ The effects of DMOG were abrogated by injection with a *hif1b* morpholino or by expression of a dominant negative form of the zebrafish *HIF1A* homolog, *hif1ab*. In contrast, mutation of the proline hydroxylation site targeted by PHDs, to create a systemic or myeloid-specific dominant active form of HIF-1 α b, recapitulated the effect of DMOG, indicating a role for HIF-1 α in delayed inflammation resolution.

In a murine trauma/hemorrhagic shock model, in which gut ischemia/reperfusion injury leads to both local and systemic inflammatory responses, *Hif1a*^{+/-} mice were protected against mucosal gut injury and subsequent lung damage, with reduced expression of TNF α , IL-1 β , and iNOS in the intestinal mucosa and IL-1 β in the lung, further demonstrating a role for HIF-1 α in promoting inflammation *in vivo*.⁸⁷

Cramer et al⁵⁰ crossed mice with floxed *Hif1a*, *Vegf*, or *Vhl* with *LysM-cre* mice to generate strains with myeloid-specific deletions of each of these genes, providing important insights into the role of HIF in innate immunity. In a model of acute skin inflammation, leukocyte infiltration, myeloperoxidase production, and edema were reduced in *LysM-cre/Hif1a* mice and increased in *LysM-cre/Vhl* mice. Likewise, in an arthritis model, edema, synovial infiltration, pannus formation, and cartilage destruction were profoundly reduced in *LysM-cre/Hif1a* mice.

HIF-1 α and VEGF upregulation correlates with increased disease severity in a mouse model of allergic rhinitis and inhibition of HIF using 2-methoxyestradiol attenuates

VEGF production and inflammation.⁸⁸ In an OVA-induced murine model of allergic airway disease, inflammatory cell infiltration, levels of Th2 cytokines (IL-4, IL-5, and IL-13), vascular permeability, fibrosis, goblet cell mucus production, OVA-specific IgE, and airway hyperresponsiveness were all significantly reduced by inhibition of HIF-1 α or VEGF.⁸⁹⁻⁹¹ Similar effects were observed in mice with a systemic *Hif1b* deletion or a myeloid-specific *Hif1a* deletion.^{89,92,93}

Activation of HIFs may be important for antibacterial responses. Following subcutaneous GAS administration, *LysM-cre/Hif1a* mice lost more weight than WT mice and developed larger necrotic skin lesions, containing approximately 1,660-fold more bacteria.⁵¹ Bacterial counts in blood and spleen were also increased, indicating greater systemic spread of infection. Similarly, HIF-1 α -mediated induction of CRAMP in keratinocytes protects against induction of skin lesions by GAS and approximately three-fold more bacteria were found in the skin ulcers of mice with a keratinocyte-specific deletion of *Hif1a*.⁹⁴

Conversely, treatment with the HIF agonist mimosine reduced lesion size by 50% following subcutaneous *S. aureus* challenge.⁹⁵ However, mice infected intraperitoneally with a mutant strain of *S. aureus* (SCV) that does not activate HIF-1 α survive, whereas those infected with the WT strain have 100% mortality except when given the HIF-1 α inhibitor 17-N-allylamino-17-demethoxygeldanamycin (17-DMAG).⁸⁴ This demonstrates the fine balance between controlling infection and avoiding excessive and deleterious inflammation.

Mice with a myeloid-specific deletion of *Hif2a* exhibit greater survival, reduced proinflammatory cytokine production and less severe hypothermia and cardiac dysfunction during LPS-induced endotoxemia.⁶² Deletion of *Hif2a* also decreases macrophage infiltration in a thioglycollate-induced model of peritonitis and neutrophil infiltration in the 12-O-tetradecanoylphorbol-13-acetate-induced model of acute skin inflammation.

Overexpression of the dominant active zebrafish *HIF2a* ortholog, *hif2aa*, did not affect neutrophil recruitment in the tail injury model described previously, but it did delay the resolution of neutrophilic inflammation.⁵³ Conversely, in a neutrophil-mediated LPS-induced model of acute lung injury, mice with a myeloid-specific *Hif2a* deletion displayed normal recruitment of neutrophils to the lung, but the clearance was more rapid, with reduced neutrophil counts after 48 and 72 hours, increased neutrophil apoptosis, and reduced markers of lung injury.⁵³

Although the majority of evidence supports a role for HIFs predominantly in promoting inflammation, they may

have anti-inflammatory roles under certain circumstances. For instance, renal F4/80⁺ cell (macrophage) accumulation in a unilateral ureteral obstruction-induced kidney injury model is increased in *Vhl*^{-/-} and *LysMcre/Vhl* mice and decreased in *Hif*^{-/-} and *LysMcre/Hif* mice (in which both *Hif1a* and *Hif2a* are deleted), suggesting that HIFs are inhibiting migration toward the site of injury.⁷¹

HIF signaling also has a complex role in tumor biology, including effects on the innate immune response, particularly tumor-associated macrophages (TAMs), which preferentially accumulate in hypoxic regions of tumors.⁹⁶ Roda et al⁹⁷ demonstrated opposing roles of HIF-1 α and HIF-2 α in tumor angiogenesis in a malignant melanoma model. In *LysM-cre/Hif1a* mice, granulocyte-macrophage colony-stimulating factor treatment induced soluble VEGF receptor, but not VEGF, resulting in a greater reduction in tumor growth and vascularization. In contrast, myeloid *Hif2a* deletion resulted in increased expression of VEGF, but not soluble VEGF receptor, and abrogated the granulocyte-macrophage colony-stimulating factor-induced reduction in tumor growth and angiogenesis observed in control mice. In a murine model of breast cancer, targeted deletion of *Hif1a* in myeloid cells reduced tumor growth, despite *Vegf* expression and vascularization remaining unchanged, perhaps due to the release of T-cells from HIF1- α -dependent suppression by macrophages.⁶⁵ Hypoxia within the tumor microenvironment also confers on myeloid-derived suppressor cells (MDSC) the ability to suppress T-cell proliferation and IFN γ production and promotes their differentiation into macrophages expressing *Il10*, *Il12*, *Nos2*, *Arg1*, and *Il6*.⁹⁶ Stabilization of HIF by desferrioxamine treatment mimics the enhancement in suppressive capacity, whereas *Hif1a*^{-/-} MDSC are poor suppressors of T-cell proliferation.

The role of hypoxia and HIFs in macrophage polarization in the context of tumor biology remains subject to debate. Laoui et al⁹⁸ suggest that hypoxia is not an important driver of macrophage polarization, as it induces HIF-1 α and HIF-2 α simultaneously, therefore promoting the expression of factors associated with both M1 and M2 responses. Tumors of PHD2^{+/-} mice, in which tumor oxygenation is increased, display a similar relative abundance of MHCII^{hi} and MHCII^{lo} TAM subsets (corresponding to M1 and M2 phenotypes, respectively) to that seen in WT mice, and the expression of hypoxia-regulated genes associated with both M1 and M2 responses is reduced in TAM infiltrating PHD2^{+/-} tumors.

Data is also beginning to emerge regarding the role of other components of the HIF signaling pathway, such as PHDs. For example, in an LPS-induced acute lung injury

model, where rapid neutrophil influx is followed by resolution, *Phd3*^{-/-} and WT mice recovered equally well in normoxia. With coexisting hypoxia, however, *Phd3*^{-/-} mice had lower BAL neutrophil counts and higher neutrophil apoptosis.⁵⁴ Increased apoptosis and reduced neutrophil numbers were also observed in bowel submucosa following dextran sulfate sodium-induced colitis, a neutrophil-dominated inflammatory model that involves tissue hypoxia. In a fatal model of abdominal sepsis, however, *Phd3*^{-/-} mice exhibited increased macrophage proinflammatory cytokine production and decreased survival, due to an overwhelming innate immune response.⁵⁹

HIF-hypoxia signaling in human infectious and inflammatory disease

Hypoxia and HIF activation are features of many human diseases, including infection, cancers, and a variety of inflammatory conditions. Immune cells are an important feature of these conditions, as are the tissue cells, and their relative and context-specific expression of the HIF pathway components may modify clinical outcomes. In skin biopsy samples from patients with a variety of cutaneous infections (including Gram-positive and Gram-negative bacteria, parasites, and viruses), a strong nuclear HIF-1 α signal is detected in keratinocytes, dermal capillaries, neutrophils, lymphocytes, and macrophages, which is absent in uninfected biopsies.⁸⁴ Biopsies from damaged intestinal mucosa of patients with inflammatory bowel disease exhibit increased HIF-1 α expression in lamina propria macrophages, compared to those from undamaged sections.⁶³ The level of HIF-1 α in damaged mucosa correlates with the expression of the scavenger receptor CD36, suggesting that HIF-1 α might promote macrophage phagocytosis of microbes and/or apoptotic cells within the colon.

HIF-1 α is also found in central hypoxic regions of human atherosclerotic plaques and colocalizes with Unc5b and netrin-1, proteins that are induced in macrophages by hypoxia and low-density lipoprotein, promoting cell survival and inhibiting migration away from the plaque.¹⁵ siRNA knockdown of *HIF1A* in U937 monocytic cells inhibits foam cell formation and reduces the expression of 57 of the 70 atherosclerotic genes that are induced by oxidized low-density lipoprotein, including *PTGS2*, *IL1B*, *ICAM1*, and *VCAM1*, suggesting that myeloid HIF-1 α may have an important role in atherosclerosis.⁹⁹

HIF-2 α expression is increased in peripheral blood neutrophils of inflammatory arthritis patients compared to those from healthy controls.⁵³ Strong nuclear HIF-2 α and

cytoplasmic VEGF immunostaining is also found in both neutrophils and neovasculature in lesions of patients with pyoderma gangrenosum, an inflammatory skin condition thought to involve a defect in resolution of neutrophilic inflammation following an inflammatory stimulus.¹⁰⁰

Although hypoxia might be expected to activate HIF equally in all cell types present in diseased tissue, this is not always the case. HIF-2 α is expressed in neutrophils infiltrating the airway of chronic obstructive pulmonary disease patients, but not in the epithelium.⁵³ Likewise, HIF-2 α protein is expressed in TAMs in the majority of uterine cervical cancer specimens, whereas <10% of specimens exhibit tumor cell HIF-2 α expression.¹⁰¹ TAMs infiltrating lung tumors also exhibit strong HIF-2 α immunostaining in the absence of HIF-2 α expression in the tumor cells themselves.¹⁰² The frequency of clusters of HIF-2 α ⁺ TAM in breast carcinomas correlates with increased density of microvessels and appears to be associated with a reduced overall survival rate.¹⁰³

In cervical cancer patients, the overall macrophage count in tumor biopsies does not affect the chance of disease-free survival; however, the higher the percentage of tumor-associated macrophages that express HIF2 α , the greater the risk of local recurrence and the lower the probability of disease-free survival following radiotherapy.¹⁰¹ In many types of cancer, including breast, ovarian, colon, and pancreatic adenocarcinomas, HIF-1 α and HIF-2 α are expressed in the tumor cells themselves.¹⁰² This might induce factors, such as chemokines, that attract monocytes and MDSC and promote their differentiation into immunosuppressive macrophages that inhibit T-cell responses, as evidenced by murine models. Urothelial carcinomas with higher expression of HIF-1 α in tumor cells exhibit higher TAM infiltration and these patients have a reduced disease-free survival rate.¹⁰⁴

HIF1A, *HIF2A*, *PHD2* and *PHD3* transcripts are increased in peripheral blood neutrophils of RA patients.^{53,54} HIF-1 α is highly expressed in synovial fibroblasts and infiltrating macrophages in the inflamed RA joint, where it enhances TNF- α - and IL-1 β -driven activation of the p38 and extracellular signal-regulated kinase signaling pathways, resulting in increased IL-33 expression.^{105,106} IL-33, in turn, upregulates HIF-1 α expression in synovial fibroblasts.¹⁰⁵ Synovial fibroblasts derived from osteoarthritis or RA patients subcutaneously engrafted into immunodeficient mice induce significantly greater myeloid cell infiltration and angiogenesis than fibroblasts derived from healthy individuals.¹⁰⁷ This effect is dependent on both HIF and VEGF, since it can be

reversed by *Hif1a* siRNA or by pharmacological inhibition of either HIF-1 α or VEGF.

Chronically low levels of oxygen may result in the down-regulation of HIFs under certain circumstances. For example, pO₂ is reduced in the skin of systemic sclerosis patients, yet HIF-1 α expression in epidermal keratinocytes is lower than that of controls, perhaps due to increased PHD levels, following induction of negative feedback loops.^{54,55,57,108}

Patients with mutations in components of HIF signaling pathways are beginning to shed light on their role in humans. Neutrophils from erythrocytosis patients with a gain of function mutation in *HIF2A* express higher levels of the target genes *PAI1* and *PHD3* and undergo lower rates of apoptosis *ex vivo*; however, phagocytosis and respiratory burst are unaffected.⁵³ Neutrophils derived from patients who are heterozygous for a mutation in the *VHL* gene exhibit lower rates of apoptosis and enhanced bacterial phagocytosis under normoxic conditions, resembling the phenotype normally seen in hypoxia, indicating the importance of proteasomal degradation in HIF regulation in human neutrophils.¹⁰⁹

Future perspectives and therapeutic potential

Given the large number of inflammatory diseases in which both hypoxia and HIF upregulation occur, and the important role of HIF in determining inflammatory responses, targeting the HIF pathway may be a promising avenue for therapies, although a number of important issues require addressing.

Chronic hypoxia is a feature of many human diseases, yet most research so far has focused on acute responses. One important area for future study will be comparing the effects of acute versus chronic hypoxia and investigating the role of negative feedback mechanisms such as HIF-mediated induction of *PHD* and *VHL* transcription in chronic activation of the HIF/hydroxylase pathway.

More detailed studies of the effects of hypoxia on circulating leukocytes compared with resident immune cells derived from different tissues are also warranted. For instance, specialized populations of macrophages exist in tissues, which may well differ from each other and from circulating monocytes in their responses to hypoxic and inflammatory stimuli. Given that certain organs, such as the lungs, have a higher basal level of oxygen than others, macrophages derived from these tissues may exhibit greater responses when exposed to hypoxia, due to lower HIF-mediated transcription of negative regulatory proteins. Furthermore, the study of hypoxic responses and HIF signaling in the innate immune system has largely focused on monocytes,

macrophages, and neutrophils, whereas research into other cell types, such as ILCs, eosinophils, and basophils, has been somewhat limited.

Efficacy of HIF-1 α inhibition in dampening asthmatic inflammation has recently been demonstrated in vivo, suggesting that inhibition of HIF signaling could be beneficial in chronic inflammatory diseases.^{89–91} However, given the importance of HIF-1 α for antimicrobial responses, care must be taken to ensure that treatments directed at the HIF pathway do not result in susceptibility to infection.

Pharmacological augmentation of HIF-1 α activity by mimosine has been proposed as a means of boosting innate immune responses and promoting bacterial clearance during infection, since mimosine was found to enhance myeloid cell bactericidal activity and to inhibit the development of necrotic skin lesions following *S. aureus* challenge in mice.^{51,95}

Augmentation of either HIF-1 α or HIF-2 α also enhances angiogenesis and wound healing, which may be beneficial in diabetes or ischemic disease.^{110,111} However, it is not clear whether this can be done safely in humans without inducing excessive inflammation, which could exacerbate tissue damage or induce shock.

HIF-1 α is upregulated more rapidly than HIF-2 α and appears to be more important for the acute response to infection, whereas HIF-2 α expression dominates during the resolution of inflammation.^{53,68} Therefore, the targeting of HIF-2 α might be useful in dampening chronic inflammation while preserving acute responses. The HIF-2 α Per-Arnt-Sim domain contains a cavity that will accommodate small molecule ligands which modulate its ability to form active heterodimers with HIF-1 β , and thus provides an avenue for potential pharmacological manipulation.¹¹² However, further investigation into the regulation of HIF-2 α and the immune functions it modulates (and how these differ from HIF-1 α) is required to determine the likely effects of modifying its expression or activity.

Suppression of downstream targets of HIF might also be a means of delivering therapies that target specific aspects of the hypoxic response while preserving those that are protective. PHD3 has been suggested as a therapeutic target in neutrophils, owing to its specific effects on hypoxic survival in this cell type. Evidence from animal models indicates that the inhibition of PHD3 would preserve neutrophil antimicrobial function, but encourage the resolution of neutrophilic inflammation in hypoxic tissues once the infection is cleared.⁵⁴ However, it is important to consider the impact of any potential therapy targeting PHD3 on other cell types,

as its role in macrophages appears to be different to that in neutrophils.

While modulation of the HIF pathway has had beneficial effects in animal models, further work is required to determine the most appropriate way of manipulating the pathway to promote resolution of infections and to treat inflammatory diseases without causing deleterious side effects. Nonetheless, the crucial role of HIFs in controlling innate immune responses, in response to hypoxia as well as other factors, makes this pathway a promising target for further investigation and potential therapies.

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Disclosure

The authors report no conflicts of interest in this work.

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