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Cycling hypoxia and free radicals regulate angiogenesis and radiotherapy response

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Abstract

Hypoxia and free radicals, such as reactive oxygen and nitrogen species, can alter the function and/or activity of the transcription factor hypoxia-inducible factor 1 (HIF1). Interplay between free radicals, hypoxia and HIF1 activity is complex and can influence the earliest stages of tumour development. The hypoxic environment of tumours is heterogeneous, both spatially and temporally, and can change in response to cytotoxic therapy. Free radicals created by hypoxia, hypoxia–reoxygenation cycling and immune cell infiltration after cytotoxic therapy strongly influence HIF1 activity. HIF1 can then promote endothelial and tumour cell survival. As discussed here, a constant theme emerges: inhibition of HIF1 activity will have therapeutic benefit.

Virchow first described the abnormal structure of tumour vessels using contrast agents in human tumours as early as the mid 1800s¹. A few decades later, Goldman was among the first to suggest that angiogenesis was associated with tumour growth². A number of other investigators in the nineteenth and early twentieth centuries evaluated tumour angiogenesis, using techniques such as microangiography, vascular casting and window chamber models. Warren has reviewed this early work in great detail³. Folkman illuminated the crucial role of tumour angiogenesis by hypothesizing that tumour growth was dependent upon angiogenesis and that, if angiogenesis could be inhibited, it could maintain tumour deposits in a quiescent state⁴. These early works set the stage for the concept that tumours require angiogenesis to provide a nutrient supply. Although it is well-established that angiogenesis occurs in nearly all human solid tumours, it does not occur in an efficient manner, leading to spatial and temporal inadequacies in delivery of oxygen and other nutrients. These inefficiencies in nutrient delivery lead to a brutal cycle of unsatisfied demand by the tumour, which drives continued aberrant angiogenesis.

The transcription factor hypoxia-inducible factor 1 (HIF1) plays an important part in tumour progression by upregulating genes that control angiogenesis, meta-stasis and resistance to oxidative stress. It also controls the switch to anaerobic metabolism, which can maintain cell

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National Cancer Institute: http://www.cancer.gov/ breast cancer | cervical cancer | colorectal cancer | glioma | head and neck cancer National Cancer Institute Drug Dictionary: http://www.cancer.gov/drugdictionary/ tirapazamine FURTHER INFORMATION

M. W. Dewhirst's homepage: http://pathology.mc.duke.edu/website/webform.aspx?id=GradDewhirst SUPPLEMENTARY INFORMATION See online article: <u>S1</u> (movie) | <u>S2</u> (movie) | <u>S3</u> (figure) ALL LINKS ARE ACTIVE IN THE ONLINE PDF

In this Review we will discuss four important and interrelated aspects of tumour hypoxia. First, we will define the hypoxic response, discussing its relevance in influencing tumour cell survival, behaviour and angiogenesis. We will examine the physiological factors that contribute to hypoxia, emphasizing a perspective based on spatial and temporal dysfunction of tumour microcirculation. The question of whether hypoxia is involved in angiogenesis initiation and the role that this might have in angiogenesis remodelling during tumour growth will be discussed. Finally, we will discuss how cytotoxic therapies, such as radiation treatment, affect tumour and endothelial cell survival, with an emphasis on a regulatory axis between hypoxia and free radicals and HIF1.

Regulation of HIF1 by hypoxia

The term hypoxia is most accurately described as an oxygenation state that is below the norm for a particular tissue. The average partial pressure of O_2 (p O_2) of normal tissues usually exceeds 20 mmHg^{5,6}. When p O_2 falls below 10 mmHg, cells typically increase production of many genes regulated by the hypoxia-inducible transcription factor HIF1 (REF. 7). HIF1 consists of α and β subunits, which heterodimerize and bind to DNA at hypoxia response elements (HREs) in promoter or enhancer regions of numerous genes to upregulate their transcription^{7–9}. These families of genes initiate anaerobic metabolism, increase angiogenesis, protect cells against oxidative stress and promote a greater degree of invasiveness and motility (FIG. 1). The <u>HIF1 α </u> subunit is exquisitely regulated by hydroxylation of proline residues in its oxygen-dependent degradation domain (ODD) that target it for recognition by the von Hippel lindau (VHI) complex and subsequent proteasomal degradation under aerobic conditions^{10,11}. A family of prolyl hydroxylases (PHDs) that require molecular oxygen to perform their function are responsible for the hydroxylation of HIF1 α^7 .

Although most tissues exist at pO₂ levels exceeding 20 mmHg, there are some tissues that exist at lower pO₂. Examples of relatively hypoxic normal tissues include pericentral vein cells in the liver¹², portions of the retina (in the dark)¹³, spinous layer of the epidermis¹⁴, the thymus^{6,15}, renal tubular epithelium¹⁶, myocardium (particularly during exercise)¹⁷ and subregions of the bone marrow that are enriched for stem cells¹⁸. Hypoxia has diverse roles in the function of these normal tissues. Hepatocytes surrounding the central vein have been shown to be hypoxic^{19,20}, and their location coincides with overexpression of vascular endothelial growth factor (VEGF) and basic fibroblast growth factor (bFGF, also known as <u>FGF2</u>)²¹. In an unperturbed state, there is no angiogenesis, so the cytokines are probably bound to proteoglycans, which maintain them in an inactive state^{22,23}. However, the inactive forms can be rapidly released after injury or stress to promote angiogenesis.

Hypoxia might be involved in squamous epithelial cell maturation in the skin. A regulator of epithelial differentiation, <u>involucrin</u>, is upregulated under hypoxic conditions *in vitro* and its expression coincides with hypoxia marker drug binding in immunohistochemical evaluation of skin samples²⁴. In the bone marrow the role of hypoxia is reversed relative to what is observed in skin. Here, the most primitive stem cells are maintained in a hypoxic niche¹⁸. These cells are maintained in a quiescent state by hypoxia-regulated proteins, such as osteopontin (encoded by <u>SPP1</u>)^{18,25}. The stem cells are prone to use anaerobic metabolism, perhaps as a means to reduce genotoxic stress associated with reactive oxygen species generated from aerobic metabolism¹⁸. Oxygen electrode and hypoxia marker drug studies

have demonstrated that the thymus of mice is heterogeneously hypoxic^{6,15}, but in this tissue there is no evidence for HIF1-regulated gene expression¹⁵. Recently it was shown that lymphocyte motility in lymph nodes is affected by pO₂, with motility decreasing with oxygen content²⁶. Thus, hypoxia in the thymus might serve to keep maturing lymphocytes within the thymus until they are ready to be released. Maintenance of hypoxia in the thymus could also have a role in protecting thymocytes from genotoxic stress.

Regulation of HIF1 by free radicals

HIF1 has been referred to as the "master regulator of oxygen homeostasis"⁷, but it is regulated by many factors aside from hypoxia, including oncogenes, growth factors and free radicals^{27–29}. Free radicals are chemical species that contain unpaired electrons; of particular importance here are oxygen-containing radicals, such as the superoxide anion (O_2^-) . The reactive oxygen molecule hydrogen peroxide (H_2O_2) , the end product of enzymatic conversion of O_2^- by superoxide dismutase (SOD), is also important in the regulation of HIF1 (REFS 28,30). Nitric oxide (NO), also a free radical, is a multifunctional biological gas that is an important cell signalling molecule at low concentrations but can be cytotoxic at high concentrations^{31,32}. There are three isoforms of nitric oxide synthase that produce NO from 1-arginine³³. Nitric oxide synthases are overexpressed in many forms of human cancer^{34,35}, in tumour cells and/or in infiltrating macrophages. The role of reactive oxygen species in regulating HIF1 activity is at least as important as hypoxia *per se* and is far more complex than hypoxia itself.

It is important to consider free radical regulation of HIF1 in the context of the oxygenation state of the cells, the concentration of each species and which species are involved. As will be discussed below, there are multiple sources of reactive oxygen and nitrogen species in tumours and these can clearly influence HIF1 activity in ways that are independent of hypoxia.

Free radical regulation of HIF1 under aerobic conditions

Increased levels of <u>SOD1</u> in pulmonary artery smooth muscle cells has been shown to increase levels of reactive oxygen species and stabilize HIF1 α , and this effect is dependent upon the activity of catalase. When catalase is overexpressed, stabilization of HIF1 α disappears, indicating that H₂O₂ is required for the response³⁶. <u>Angiotensin II</u> and <u>thrombin</u> can stimulate generation of free radical species and mimic the effects of hypoxia to upregulate HIF1 α levels and activity. It has recently been shown that angiotensin II mediates this effect by increasing the levels of H₂O₂, which depletes intracellular stores of ascorbate, an important co-factor involved in PHD activity³⁷. Thus, H₂O₂, the product of O₂⁻ dismutation by sOD, is one candidate likely to be responsible for stabilization of HIF1 α under aerobic conditions.

NO can influence the levels of HIF1 α in complex ways. Mateo and moncada reported that NO concentration has a strong influence on whether HIF1 is stabilized under aerobic conditions³⁸. Human HEK-293 cells were stably transduced with an inducible nitric oxide synthase (iNOs), under control of the tetracycline promoter. By varying the amount of tetracycline in media, they were able to control NO concentrations and increased HIF1 α protein levels were observed when NO concentrations exceeded 5 µm. There are two potential mechanisms for this effect. Exposure to NO has been shown to nitrosylate thiols in the HIF1 α protein leading to HIF1 α stabilization under aerobic conditions³⁹. Recently, the specific target has been identified to be a cysteine residue in the ODD of HIF1 α , which blocks its binding to the VHI complex for subsequent degradation²⁷. NO can promote binding of HIF1 to HREs in HIF1 target genes by binding to an adjacent HIF1 ancillary sequence (HAs) and hence acting as a transcriptional co-activator⁴⁰. It has also been

reported that NO can upregulate the rate of HIF1 α synthesis by activating the phosphatidylinositol 3-kinase (PI3K)–mAPK (mitogen-activated protein kinase) pathway^{41,42}.

Free radical regulation of HIF1 under hypoxic conditions

Schumacker and Chandel have theorized that the oxygen sensor in cells is not the PHD but rather the formation of reactive oxygen species by mitochondria under hypoxia that lead to stabilization of HIF1 $\alpha^{28,30}$. Evidence for this mechanism of stabilization of HIF1 is multifold. First, ρ^0 cells, chemically depleted of functional mitochondria, cannot increase HIF1a levels in response to hypoxia²⁸. Second, overexpression of catalase blocks hypoxic stabilization of HIF1a, whereas administration of H2O2 to cells increases HIF1 expression²⁸. Recently, a genetic approach has been taken to further test the hypothesis that reactive oxygen species are important in hypoxic signalling for HIF1a stabilization and hypoxia signalling. It has been reported that mitochondrial complex III may be important in this pathway. Cells in which complex III activity has been knocked down through RNA interference-mediated reduction of expression of one of the protein subunits of this complex show decreased reactive oxygen species formation and decreased HIF1 α expression levels²⁹. When H_2O_2 is added back, the knockdown cells regain the ability to stabilize HIF1a under hypoxia. This result suggests that electron flow through complex III is required for hypoxic stabilization of HIF1 α . Cytochrome c, which is responsible for shuttling electrons between complex II and complex III, might be central to this sensing mechanism. Mouse embryonic fibroblasts depleted of cytochrome c are completely incapable of eliciting an increase in HIF1 α in response to hypoxia, but these same cells show strong induction of HIF1 α after exposure to H₂O₂ (REF. 43). Similarly, cells in which mitochondrial transcription factor A (TFAm) is knocked down, rendering them incapable of making mitochondria, cannot produce reactive oxygen species nor can they stabilize HIF1 α under hypoxic conditions³⁰. Finally, overexpression of manganese superoxide dismutase (MnSOD, also known as SOD2), the mitochondrial sOD, has been shown to inhibit HIF1a levels, downregulate VEGF and inhibit tumour growth^{44,45}.

The effects of NO in controlling HIF1 activity under hypoxic conditions are complex. The initial report indicated that NO inhibits stabilization of HIF1 under hypoxia⁴⁶. The underlying mechanism for this effect does not appear to be mediated by nitrosylation of the cysteine residue in the ODD, although deletion of the ODD eliminates the effect of NO in downregulating HIF1 α protein levels⁴⁷. As detailed above, under aerobic conditions, NO nitrosylates this residue to prevent degradation of HIF1 (REF. 27). Using reporter genes that were targeted to either mitochondria or cytosol, Hagen demonstrated that NO inhibits respiration by inhibiting cytochrome *c* oxidase activity under hypoxic conditions, thereby permitting increased PHD activity as oxygen is redistributed to the cytosol⁴⁸. Recently it has been reported that NO elicits a feedback regulatory mechanism between HIF1 and PHD2 (also known as <u>EGLN1</u>)⁴⁹. When cells are first exposed to NO, HIF1 α levels transiently increase, stimulating increased expression of PHD2. Increased levels of PHD2 can then feed back to inhibit HIF1 α levels. In summary, the regulation of HIF1 by NO under hypoxic conditions centres on modification of the ODD, but might involve redistribution of residual oxygen from the mitochondria to the cytosol.

In summary, the roles that free radical species have in controlling HIF1a levels and HIF1 transcriptional activity are complex and interrelated. We have presented this story in four categories, segregating hypoxia and normoxia and reactive oxygen species and nitric oxide. However, it is likely that all of these pathways and factors are active simultaneously in tumours, with micro-regional and temporal variations. However, as discussed below, the

role of reactive oxygen species in regulating HIF1 α might overshadow the effect of hypoxia *per se* on HIF1 α expression.

In <u>cervical cancer</u> it has been shown that there is good correlation between the expression of HIF1 targets and the hypoxia marker drug, pimonidazole⁵⁰, but in other reports there has been poor correlation between HIF1 or its regulated cytokine VEGF⁵¹ and localization of hypoxia marker drugs^{52,53}. Given the complexities involved in regulating HIF1 α , as outlined above, it comes as no surprise that there is generally a lack of correlation between hypoxia and HIF1 levels in tumours. In <u>head and neck cancer</u> it has recently been reported that HIF1 α expression levels tracked with expression of eNOs (also known as nitric oxide synthase 3 (<u>NOS3</u>))⁵⁴, indicating that NO levels might independently regulate HIF1 α levels. As will be discussed below, the complexities of oxygen transport and angiogenesis in tumours will also probably contribute to heterogeneity in the regulation of HIF1 activity and consequently correlations between HIF1 α expression and hypoxia.

Origins of hypoxia in solid tumours

The presence of hypoxia in tumours is a well-established source of resistance to radiation therapy and chemotherapy⁵⁵. In the case of radiation therapy, this is known to be related to the effect of oxygen in stabilizing DNA damage from radiation in a form that is not easily repaired. The activity of many chemotherapeutic drugs is also altered at low oxygen concentration, at which the percentage of cycling cells is relatively low and drug delivery may be compromised⁵⁵. Overexpression of HIF1 α , driven by hypoxia and free radical species, will also contribute to therapy resistance. Thus, it is important to understand the underlying mechanisms that contribute to failure to deliver adequate oxygen and other nutrients to tumour cells. There are several methods to measure hypoxia in tumours (BOX 1).

Deficiencies in oxygen transport

The oxygenation state of any location in a tissue is the result of a balance between delivery and consumption of oxygen. There are several excellent reviews on this subject^{5,56,57}. The deficiencies of oxygen transport can be summarized as having eight dominant features: first, a relatively sparse arteriolar supply^{5,58}; second, inefficient orientation of microvessels^{59,60}; third, low vascular density; fourth, extreme variations in microvessel red blood cell flux (many tumour vessels do not carry red blood cells, whereas large diameter shunt vessels contain a relative abundance of red blood cells⁶¹); fifth, the limited arteriolar supply can lead to pathologically low vascular pO₂ in regions distant from the arteriolar source (longitudinal oxygen gradient)^{58,62,63}; sixth, hypoxic red blood cells stiffen, increasing blood viscosity, which contributes to sluggish flow; and seventh, large diameter shunts can divert blood away from the tumour bed. Consequently, networks of microvessels can exhibit low pO₂, leading to relatively large regions of tumour that are dominated by microregions of pathologically low pO₂^{58,62,64–66} (FIG. 2).

The seven features discussed above influence the overall oxygenation state of a tumour, but an important feature of tumour hypoxia is that it is unstable. The eighth dominant feature of tumour hypoxia is that it is cyclical, a phenomenon proposed almost 30 years ago^{67-69} . This feature of tumour pathophysiology has often been referred to as "intermittent" or "perfusion limited" hypoxia^{70,71}. We believe that a more precise definition might be 'cycling hypoxia' because this feature demonstrates clear patterns of periodicity across many tumour types⁷⁰. Two of the first studies on this subject used dorsal skin fold window chambers in rats (BOX 2), which allowed direct visualization of the developing vasculature of tumours under a microscope in the living animal. Reinhold observed fluctuations in redox ratio in tumour tissues. The redox ratio is derived from the ratio of fluorescence of endogenous molecular

species involved in respiration and thereby reflects oxygenation status⁶⁸. Yamaura observed vascular stasis in microvessels in skin fold window chamber tumours near regions of tumour regrowth after radiation therapy. He surmised that tumour cells near such vessels would become hypoxic during vascular stasis. If stasis happened during radiotherapy, it could be responsible for hypoxic radioresistance⁶⁹. Martin Brown was the first to demonstrate that intermittent hypoxia could actually be a cause for radioresistance. He selectively killed hypoxic EmT6 tumour cells in mice with a hypoxic cytotoxin and 24 h later found reemergence of a radiobiologically significant hypoxic fraction⁶⁷.

Kinetics

There is strong evidence that the kinetics of cycling hypoxia are complex, with at least two dominant timescales. One has a frequency of a few cycles per hour and the second exhibits variations on a timescale of hours to days. These two frequencies superimpose on each other to yield a complex timescale for fluctuations. An analogy is the relationship between the tides and waves in the ocean. Both occur at the same time, but the waves have a higher frequency than the tides. Also, the tides influence how high the waves wash up onto the shore. In the same way, the degree of hypoxia is influenced by the overall oxygenation state of large tumour regions (analogous to tides), superimposed on higher frequency instabilities (analogous to waves) that are caused primarily by fluctuations in microvessel red blood cell flux.

The higher frequency cycles were described first. Chaplin and co-workers defined a dominant periodicity for flow instability and radioresistance of around 20 min for a variety of different preclinical tumour models^{72–76}. In their first experiments, the investigators administered a dye that binds to the nuclei of cells surrounding micro-vessels to animals either simultaneously with or sequentially with radiation treatment⁷⁷. Following irradiation, the tumours were excised and sorted by perfusion dye intensity using flow cytometry. The brightest cells and dimmest cells were assumed to be most proximal and distal to microvessels, respectively. The fraction of tumour cells surviving was determined for each cohort of cells (classified by dye staining intensity). When the dye and irradiation were given simultaneously, the dimmest cells were the most radioresistant, but when there was a 20 min time delay between dye and irradiation, there was no relationship between dye staining intensity and radiosensitivity. The conclusion was that the cells that were hypoxic at time zero were no longer hypoxic 20 min later.

Durand administered two perfusion marker dyes, either simultaneously or separated in time by periods of up to 30 min^{71,78,79}. He observed regions of dye mismatch in frozen sections of tumour samples when the dyes were administered with a time delay. He also observed differences in perfusion marker dye fluorescence intensity, concluding that variations in blood flow rate may be sufficient to induce changes in the level of hypoxia⁷⁹. Skin fold window chamber tumour studies showed that perivascular pO₂ and red blood cell fluxes were temporally synchronized^{61,80,81}. Theoretical predictions were that fluctuations in vascular pO₂ would influence the pO₂ of cells residing several cell layers away from the microvessel. Recently, this prediction was confirmed experimentally, demonstrating for the first time that fluctuations in red blood cell flux are primarily responsible for cycling hypoxia in tumours⁸¹. Implantable microsensor studies have shown that red blood cell flux and pO₂ fluctuate a few cycles per hour in several different tumour lines^{82–84}. Similar results have been reported using laser Doppler flowmetry and fibre-optic oxygen sensors in human and canine tumours, respectively^{83,85}.

Benne with performed multiple administrations of a hypoxia marker drug to obtain steady state levels of the drug systemically over 8 h. Using flow cytometric analysis of cells taken from the tumour and an antibody to detect the presence of drug–protein adducts, they found

that the hypoxic fraction increased following chronic administration to a level higher than administration of a single bolus of a similar marker drug. This result suggested that intermittent hypoxia is not relegated to specific subregions within a tumour⁸⁶. When they administered a second hypoxia marker drug 1 h after the last administration of the first drug, they found a substantial portion of cells that were labelled with one drug but not the other. They observed variation in the fraction of cells experiencing these events in two different tumour lines, but the results suggest that relatively large subregions of tumours can experience transient periods of hypoxia and reoxygenation.

Spatial characteristics of pO₂ instability

Cardenas-Navia *et al.* used phosphorescence lifetime imaging to serially monitor vascular pO_2 in 2–3 mm regions of skin fold window chamber tumours over periods of 60–90 min. The oxygenation state of all microvessels continuously changed in three different tumour lines, with significant differences in the spatial relatedness of the instability⁸⁷ (see supplementary information <u>S1 & S2</u> (movies)). Baudelet *et al.* used serial measurements of blood oxygen level detection (BOLD) magnetic resonance imaging (MRI) to identify rather macroscopic (a few millimetres in diameter) regions that exhibited synchronized fluctuating signals related to the degree of haemoglobin saturation. Sorg measured haemoglobin saturation in networks of tumour microvessels, demonstrating that fluctuations were temporally coordinated in regions of a few hundred microns in dimension⁸⁸. These results indicate that small vascular networks are probably cycling together⁸⁹. Similar results have been reported comparing serial dynamic contrast enhanced (DCe) MRI measurements in human tumour xenografts⁹⁰. These preclinical studies suggest that cycling hypoxia involves relatively large tumour regions that are connected by common microvascular networks.

Mechanisms underlying cycling hypoxia and implications

The higher frequency hypoxic cycles are caused by variations in red blood cell flux within tumour microvessels^{80,81,85}. Fluctuations in feeding arteriolar diameter have been shown to match temporal changes in microvessel red blood cell flux in the same tumour⁶¹. Similarly, fluctuations in BOID mRI signals have been observed more frequently in tumour regions that contain blood vessels with smooth muscle⁹¹. Thus, some of the fluctuations in red blood cell flux might be due to vasomotion of resistance vessels. Using serial observations of tumour microvessels in window chambers, Patan identified vascular intussusception, a process in which microvessels remodel to split into parallel smaller vessels⁹². This process occurred over periods of tens of minutes, which is consistent with the time frame for the high frequency cycling hypoxia. Flow resistance (R) is inversely proportional to vessel radius to the fourth power (R = $8\eta/\pi r^4$; where η = viscosity and r = vessel radius)⁹³. Thus, as a microvessel splits, it will reduce flow in the two vascular segments disproportionally to the split in diameter, leading to redistribution of flow within the network. It is important to note that even small perturbations in microvessel diameter in a small proportion of vessels within a network can profoundly influence haemodynamics within an entire network^{94,95}. Uneven distribution of red blood cells at microvessel bifurcation points would be more exaggerated in tumours than normal tissues because hypoxic red blood cells are less deformable, which increases blood viscosity96.

Using window chamber tumours, we have observed dynamic changes in microvascular structure on a daily basis. Sometimes, the vascular network changed so dramatically that the same microvessels could not be observed serially⁹⁷. These results suggest that vascular remodelling might be responsible for the slower frequency of cycling hypoxia. Recent ¹⁸F-misonidazole (a hypoxia marker drug) positron-emission tomography (PET) studies in human patients with head and neck cancer have verified that large regions of tumours can experience the slower frequency cycling hypoxia⁹⁸. In these studies, serial PET studies

The primary consequence of cycling hypoxia is upregulation of HIF1 activity to a level that supersedes that of chronic hypoxia. Peng *et al.* examined HIF1 expression in the carotid body following 10 days of repeated 15 s exposures to 5% O₂, followed by breathing air. They observed significant upregulation of HIF1 α levels in carotid bodies of mice⁹⁹. Similar results were seen *in vitro* with the PC12 cell line¹⁰⁰. The mechanism of enhanced HIF1 activity is multifactorial, stemming from alteration in the rate of HIF1 α synthesis, regulated by mTOR (also known as FRAP1), and phosphorylation of the HIF1 binding cofactor CBP (also known as p300 and CREBBP)¹⁰¹. Repeated exposure of endothelial cells to hypoxia and reoxygenation can lead to much greater upregulated by increased phosphorylation of HIF1 α ^{102,103}. Cycling hypoxia can also be expected to increase levels of free radicals, as a result of hypoxia–reoxygenation injury. Activation of the xanthine oxidase pathway during hypoxia–reoxygenation injury results in release of H₂O₂, secondary to lipid peroxidation. As discussed above, H₂O₂ can stabilize HIF1 α .

Features of cycling tumour hypoxia

The model that we propose incorporates vascular geometry, radial diffusion of oxygen from vessels, distribution of red blood cells at bifurcation points and instabilities in red blood cell flux, all of which can lead to dynamic changes in the oxygenation of dependent tumour cells (FIG. 3). These features represent the higher frequency waves of cycling hypoxia. The location, shape and steepness of the oxygen gradient, which is influenced by vascular density and proximity to the arteriolar source, will vary according to how the structure of the larger vascular network changes over time. Remodelling of the networks occurs on a slower timescale and would result in the slower tidal cycling of hypoxia. As both chronic and cycling hypoxia can influence HIF1 transcriptional activity, it is expected that there will be phenotypical changes in cells as a consequence. One of the primary consequences might be alterations in angiogenesis, as most pro-angiogenenic factors are regulated by HIF1.

Hypoxia and angiogenesis

There is considerable evidence that upregulation of the HIF1-regulated protein VEGF and other factors such as angiopoeitin 2 (<u>ANGPT2</u>) promote the initiation and promulgation of angiogenesis. Other angiopoeitins also have a role^{104–115}. The 'angiogenic switch' describes a collection of circumstances under which angiogenesis can be initiated, which is determined by a balance of proand anti-angiogenic factors. Importantly, hypoxia has been thought of as a primary trigger that tips the balance toward angiogenesis, as it will increase HIF1 activity and hence VEGF levels^{115–117}. As many cytotoxic treatments change tumour oxygenation, it might be expected that treatments may also influence the angiogenic response of a tumour.

Oncogenic regulation of angiogenesis has also been suggested, either alone or working in combination with hypoxia, to stimulate angiogenesis. For example, upregulation of <u>HRAS</u> and epidermal growth factor receptor (<u>EGFR</u>)^{118,119} and inactivation of the tumour suppressor gene <u>TP53</u> (REFS 120,121), can increase rates of synthesis of HIF1 α to a level that overwhelms the capacity of the degradation pathway to eliminate it¹²². The increase in HIF1 synthesis appears to be regulated by the PI3K–AKT pathway¹²³.

VEGF upregulation is a trigger for initiation of angiogenesis from dormant metastases and the angiogenesis inhibitor <u>thrombospondin 1</u> might inhibit VEGF activity to promote dormancy¹²⁴. Hypoxia, on the other hand will increase HIF1 activity to upregulate VEGF

while downregulating thrombospondin, to create a pro-angiogenic environment¹¹⁷. Some metastatic tumours replace normal parenchymal cells with tumour cells thereby co-opting normal vasculature. For example, human <u>breast tumours</u> that have metastasized to liver show this phenotype. They have been reported to exhibit little evidence for hypoxia (indicated by expression of the endogenous hypoxia marker protein carbonic anhydrase IX) and express low levels of VEGF^{125,126}. Hepatic metastases of <u>colorectal cancer</u>, by contrast, show high levels of carbonic anhydrase IX and increased VEGF and clearly exhibit angiogenesis¹²⁵.

In the case of primary tumours, growth beyond a few millimetres is unusual in the absence of angiogenesis. As an example, consider the case of intraductal breast cancer (ductal carcinoma *in situ*). These lesions are multifocal with typical diameters of 200–300 μ m. Bos *et al.* examined HIF1 α expression in ductal carcinoma *in situ* lesions and found that HIF1 α and VEGF expression and angiogenesis were not apparent in hyperplasia. The frequency of tumours with positive HIF1 expression increased proportionally with tumour grade; VEGF expression and angiogenesis tracked with HIF1 expression¹²⁷. These data are consistent with HIF1 driving VEGF production and angiogenesis in these early lesions. However, it is not known whether hypoxia-induced stabilization of HIF1 α has a dominant role in the initiation of angiogenesis because of the multiple regulatory mechanisms of HIF1 α expression and activity that could be involved (such as the presence of free radicals or oncogenic regulation).

Holash *et al.* performed elegant experiments using an intracranially transplanted C6 glioma model to examine whether there was evidence for hypoxia-driven angiogenesis initiation¹⁰⁹. The experiments were performed by transplanting the tumour cells into the brain of recipient rats and then examining the tissues histologically at set time points after transplant. They observed vessel cooption in the first week after tumour transplantation, which was characterized by histological evidence of tumour cells infiltrating around pre-existing normal vessels. Immunohistochemical data showed breakdown of vascular basement membrane and vessel regression during week 2, and by week 4 there was clear evidence of vasodilation and new vessel formation interspersed with islands of necrosis. *In situ* hybridization showed increased *Angpt2* mRNA expression before the upregulation of VEGF. This was also observed in a rat mammary carcinoma and in lung metastases from lewis lung carcinoma. They hypothesized that vascular shutdown, caused by ANGPT2mediated de-differentiation of microvessels, in the absence of VEGF (and hence angiogenesis), led to hypoxia. Thus, in their model, a 'hypoxic crisis' was responsible for the initial upregulation of VEGF and onset of angiogenesis.

We used the mouse dorsal skin fold window chamber model to address the same question. Window chambers were implanted with murine and human tumour cell lines, stably transfected with two reporter genes: red fluorescent protein under control of a constitutive CmV promoter (RFP_c), and green fluorescent protein under control of an HRE derived from the *VEGF* promoter region (GFP_h) ⁹⁷. Angiogenesis occurred before detection of GFP_h, typically by 1–2 days. The hypoxic cytotoxin <u>tirapazamine</u> delayed the detection of GFP_h, but had no effect on angiogenesis initiation. Both tumour lines expressed VEGF at low levels under normoxia, but several hours of hypoxia significantly increased VEGF expression *in vitro*. In contrast to Holash's theory, we did not observe vascular stasis before angiogenesis onset, but angiogenesis was accelerated when HIF1 α was upregulated. Thus, we term our model the 'acceleration model'. We propose that HIF1 α upregulation accelerates tumour angiogenesis, as opposed to its having a role in the initiation stages. There is a normal tissue corollary to this, based on wound healing. In a rat dermal skin wound model, hypoxia was not involved in the initiation of angiogenesis; rather, it was associated with the period of wound involution after reepithelialization¹⁴. In an analogous

fashion, HIF1 upregulation and increases in VEGF in response to tumour hypoxia may be important in vascular remodelling during continued tumour growth¹²⁸. In summary, results from *in vivo* window chamber and wound healing models are not consistent with the vascular crisis theory of Holash, and instead point to a role for hypoxia in angiogenesis acceleration and remodelling (FIG. 4). Because the results from the studies presented above came from different tumour models, more efforts are needed in a broad range of tumour models to further elucidate the role of hypoxia in the initiation and/or acceleration of tumour angiogenesis.

HIF1 regulation after angiogenesis initiation

To further establish the interrelationship between angiogenesis and oxygen delivery we used optical methods to directly measure haemoglobin saturation of microvessels in the same tumours in which we measured GFP_h expression. In the earliest stages of tumour growth, GFP_h was found in regions relatively devoid of vasculature (hypoxic). As tumour growth continued, however, we often observed relatively high expression of GFP_h in the immediate vicinity of microvessels⁶³. We knew that these vessels were relatively well-oxygenated because Hb_{sat} levels indicated sufficient O₂ supply (see supplementary information <u>S3</u> (FIGURE)). Therefore, reactive oxygen species and/or NO might be stabilizing HIF1 α in perivascular tumour cells. This scenario could further accelerate angiogenesis and vascular remodelling as a result of high perivascular VEGF concentrations.

Effects of radiation therapy on HIF1 activity

It is well-established that hypoxic tumours will reoxygenate after treatment with radiation and some forms of chemotherapy 129-133. The effect is probably caused by reduction in oxygen consumption rate as a result of death of the more radiosensitive oxygenated tumour cells in addition to an increase in perfusion^{130,134,135}. We used the RFP_c and GFP_h cell lines described above to evaluate the effects of radiation therapy on HIF1 activity¹³⁶. It was expected that reporter gene expression would decrease after radiation treatment, commensurate with improved oxygenation. However, the opposite effect was observed: GFP_h reporter activity actually increased up to 72 h after treatment. This was confirmed by showing that HIF1 α protein levels increased, along with several HIF1-regulated proteins. Two independent mechanisms were found to be responsible for this effect. First, increased free radicals after radiation: if the accumulation of these radicals was inhibited using a manganese porphyrin sOD mimetic, the stabilization of HIF1a was prevented and the tumour microvasculature was sensitized to radiation damage. Antisense knockdown or pharmacological inhibition of HIF1 synthesis after radiation treatment was shown to also induce significant growth delay in several tumour lines by our group^{136–138}. In a separate study, Williams et al. showed prolonged tumour growth after radiotherapy in a tumour line that is null for HIF1 β (also known as ARNT), and therefore unable to render a HIF1 response¹³⁹. The second mechanism of HIF1 α stabilization after reoxygenation was related to stress granules¹⁴⁰. These are protein-mRNA complexes that form in cells during periods of stress and prevent the cell from using energy for protein translation. Stress granule formation coincided with hypoxia in vivo and in vitro and disaggregated during reoxygenation, releasing HIF1-regulated transcripts. When stress granule formation was blocked by using an antisense against one of the key proteins involved in forming the stress granule, HIF1-mediated gene expression was reduced¹³⁶.

Recently, in collaboration with Chuan li's group, we found another pathway for HIF1 α stabilization after radiation therapy. A second peak of HIF1 α stabilization occurs at 6–7 days after treatment, coinciding with macrophage infiltration into tumours²⁷. In this case, NO production, mediated by infiltrating macrophages, was responsible. Inhibition of macrophage function or blockade of NO synthesis during and after radiation treatment

prevented upregulation of HIF1 α and prolonged growth delay after radiation treatment. The specific target of NO in this case was a single cysteine residue in the ODD of HIF1, which prevented HIF from being recognized by the VHl complex (FIG. 1).

Conclusion

It is important to note that HIF1 has complex effects on tumour cell biology and these effects might have a role in optimizing strategies for HIF1 inhibition when combined with radiation or chemotherapy. On one hand, blockade of HIF1 activity severely compromises the ability of tumour cells to undergo anaerobic glycolysis¹³⁸. This will reduce the proliferation rate of hypoxic cells and promote necrosis or apoptosis in cells that are deprived of both oxygen and glucose. This effect is attractive, as it provides a clear mechanism for killing relatively radioresistant and chemoresistant hypoxic cells. On the other hand, cells that contain wild-type p53 are more prone to undergo apoptotic death after radiation if the HIF1 response is intact¹³⁸. Anti-tumour effects of HIF1 inhibition may be preserved in such cases if HIF1 inhibition is initiated after cytotoxic therapy. Alternatively, it has recently been reported that inhibition of angiogenesis can lead to tumour cell apoptosis when combined with ¹³¹I radiotherapy in a tumour line that does not contain a functional p53 (REF. 141).

The response of HIF1 to cytotoxic therapy appears to be predicated on the presence of hypoxia before treatment. Both stress granules and free radical species are important in stabilizing HIF1 α and hence induce HIF1-mediated gene expression after treatment, potentially reducing the sensitivity to such agents. The upregulation of HIF1 activity in this setting promotes endothelial cell survival, while creating pleiotropic effects on tumour cells that counterbalance promotion of apoptosis versus cell survival. The axis can be tipped toward cell death by judicious scheduling of HIF1 inhibition with radiotherapy or by using anti-angiogenic therapies in combination with radiotherapy (FIG. 5).

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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Glossary

Microangiography This method permits visualization of microvessels in a tissue following injection of an X-ray contrast agent and exposure to X-rays; X-ray film or digital imaging are used to visualize the vessels.

Vascular casting With this method, a cast of microvessels as they actually exist in tissues is preserved to permit visualization. Vasculature is filled with a polymer that sets inside the vessels. The tissue is then digested away to leave just the cast of the microvessels.

At a glance

- This Review discusses four important and interrelated features of tumour hypoxia: the hypoxic response, the factors that influence tumour hypoxia, the role of hypoxia in the initiation of angiogenesis (angiogenic switch) and how hypoxia influences treatment responses.
- The hypoxia response, driven primarily by the heterodimeric transcription factor hypoxia-inducible factor 1 (HIF1) influences cell survival, behaviour and angiogenesis.
- Several pathophysiological factors contribute to the development of tumour hypoxia, which is typified by heterogeneity in oxygenation in space and in time.
- Conflicting theories exist with respect to whether hypoxic stabilization of HIF1 is the primary feature of the angiogenic switch. There is clear evidence that HIF1 upregulation is associated with angiogenesis acceleration as opposed to angiogenesis initiation.
- The appearance of perivascular (oxygenated regions) HIF1 expression during angiogenesis acceleration might be the result of increased levels of reactive oxygen species, associated with proliferation and/or instability in flow and hypoxia–reoxygenation injury.
- Cytotoxic therapies, such as radiation therapy, improve tumour oxygenation but also increase HIF1 levels and transactivation of target genes through mechanisms associated with stress granule depolymerization and the production of free radicals. The upregulation of HIF1 in these circumstances protects tumour and endothelial cells from damage by the cytotoxic therapy.

Shunt vessels These are large-diameter vessels that directly connect between feeding and draining vessels at the periphery of a tumour. These can shunt flow around the main body of the tumour, thereby starving the tumour of nutrients.

Redox ratio This ratio, which is derived from the relative abundance of two naturally fluorescent coenzymes, fAD and NADH, is related to the metabolic activity of a tissue.

Hypoxic cytotoxin Hypoxic cytotoxins are drugs that are selectively toxic to hypoxic cells.

Laser Doppler flowmetry This method measures velocity of red blood cells in tissue. When a laser illuminates tissue the light strikes red blood cells that are moving. The reflected light undergoes a detectable change in shift in frequency (Doppler shift) that is related to red blood cell velocity.

Blood oxygen level detection magnetic resonance imaging (BOLD MRI.) This is an MRI method that is sensitive to the difference in magnetic properties of deoxyhaemoglobin versus oxyhaemoglobin.

Dynamic contrast enhanced MRI (DCe MRI.) This is a method to measure perfusion/ permeability of an MR contrast agent as it enters and leaves a tissue following bolus intravenous injection. Kinetic analysis permits derivation of parameters related to vascular permeability and perfusion.

Vasomotion Arterioles in the peripheral circulation exhibit fluctuations in diameter that control perfusion of dependent tissues.

Bifurcation point This term refers to a branch point in the microcirculation, in which flow splits from one vessel to two or more daughter vessels.

Carotid body The carotid body is a collection of pH-, partial pressure of O_2 (pO₂)- and pCO₂-sensitive chemoreceptor cells located on the carotid artery wall. When stimulated they send signals to the central nervous system to regulate respiratory and heart rates.

Dormancy This is a state of tumour growth at a time when cell loss is equal to cell proliferation and before the onset of angiogenesis.

Stress granules Cell stress initiators, such as nutritional deprivation or hypoxia, cause a general downregulation of protein translation, involving prevention of mRNA entry into ribosomes, forming stress granules (in complex with RNA-binding proteins) in the cytoplasm. Stress granules rapidly disaggregate upon removal of stress, permitting subsequent protein translation.

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Box 1 | Methods to measure hypoxia

There are a number of methods to measure hypoxia in tissues; all have advantages and disadvantages. Three key methods that are relevant to the discussions in this paper are below (for reviews see REFS 142–147).

Hypoxia marker drugs

These drugs are 2-nitroimidazoles; after entering cells they undergo 1–2 electron reduction in cells and the reduced drug is a highly reactive free radical that binds to macromolecules, including proteins¹⁴⁸. However, when oxygen is present, the drug is oxidized and reverts back to its original state, allowing it to diffuse out of the cell and eventually into the circulation. The rate of protein binding of the reduced drug increases exponentially with a decrease in partial pressure of O₂, particularly below 10 mmHg^{144,149}. The presence of drug–protein adducts can be detected immunohisto-chemically using antibodies specific for the drug protein adduct. ¹⁸F-labelled versions of these drugs are also being developed for positron-emission tomography imaging^{150,151}.

Hypoxia marker proteins

As HIF1 upregulates the synthesis of many proteins, it has been proposed that identification of such proteins in tissues could be markers of hypoxia. Literally dozens of such proteins have been studied at the preclinical level and in clinical trials¹⁵². Some of the more promising endogenous markers include carbonic anhydrase IX (<u>CA9</u>), plasminogen activator inhibitor 1 (PAI1, also known as <u>SERPINE</u>), osteopontin and <u>lysyl oxidase</u>. Combinations of markers might prove to be better predictors of clinical outcome than any single marker¹⁵³.

Oxygen electrodes and optical probes

Oxygen can be measured in any aqueous media using polarography. In principle, two electrodes are placed into the medium and a polarizing voltage of -0.7 volts is applied across them. This voltage corresponds to the binding energy of outer shell electrons of oxygen. The electrons are captured by the cathode and the current generated is linearly proportional to oxygen concentration. In practice, the cathode is embedded into a needle that can be introduced into tissues and the anode is placed on the body surface. This technique has been used extensively in preclinical and clinical studies; the presence of hypoxia is an independent predictive factor for poor prognosis in many different tumour types¹⁵⁴. Optical probes have also been developed: these are implanted into tissues and contain a fluorochrome that emits fluorescent light with a certain decay rate when illuminated. The rate of fluorescent light decay is proportional to the oxygen concentration in the region of measurement. These probes yield data similar to that of the oxygen electrode⁶.

Box 2 | Multiparametric imaging with the window chamber model

Window chamber models in rodents permit multiparametric analysis of tumour microcirculatory function and gene expression in the living animal. Examples of endpoints include vascular geometry, red cell flux, flow velocity, vascular permeability, leukocyte-endothelial cell interactions, tumour-endothelial cell interactions and haemoglobin saturation. Typically more than one of these parameters is measured in the same animal in the same experiment, which permits study of relationships between measured parameters. Because these measurement methods are optical and therefore noninvasive, the endpoints described can be measured repeatedly on the same animal, providing a powerful method for serially assessing the effect of therapeutic treatment or other manipulations. Shown in this example are three parameters that are particularly related to the evaluation of oxygen transport. In part **a**, two reporter genes are shown, which are simultaneously expressed in the same tumour cell line. Red indicates constitutively expressed red fluorescent protein. This reporter gene permits identification of all tumour cells within the field of view. Green indicates hypoxia-inducible factor 1 (HIF1)-regulated green fluorescent protein. This permits identification of tumour cells in which HIF1 is transcriptionally active. In part b, haemoglobin saturations of tumour microvessels are imaged. Orange to red represents 70-90% saturated. Light blue represents 10–30% saturated. Figure part \mathbf{c} shows the appearance of microvessels using transillumination. Images reproduced, with permission, from REF. 156 © Elsevier Ireland Ltd (2007).





Figure 1. Features of hypoxia-inducible factor (HIF1) regulation

HIF1 transcriptional activity regulates numerous genes that control a variety of cellular functions, including anaerobic metabolism, angiogenesis stimulation and mechanisms for resistance to therapy. HIF1 transcriptional activity requires formation of a heterodimer consisting of HIF1 α and HIF1 β . The heterodimer binds to hypoxia response elements (HREs) in promoter regions of its target genes, where it activates transcription. Whereas HIF1 β is constitutively expressed, HIF1 α protein levels are subject to a number of points of regulation. HIF1a consists of the following regulatory domains: bHLH (basic-helix-loophelix), PAS (Per-ARNT-Sim), NTAD (N-terminal transactivation domain), CTAD (Cterminal transactivation domain) and ODD (oxygen-dependent degradation domain). The rate of synthesis of HIF1 α is controlled by activation of the phosphatidylinositol 3-kinase (PI3K)-Akt and Ras pathways by a variety of stimuli. HIF1a protein is rapidly targeted by the von Hippel-Lindau protein (VHL) complex for proteasomal degradation under normoxic conditions, following an oxygen-dependent prolyl hydroxylation of proline residues in the ODD. Activity of the prolyl hydroxylases (PHDs) is influenced by protein kinase C (PKC), PTEN and reactive oxygen species (ROS). Nitric oxide (NO) has variable effects on the stability of HIF1 α , depending on the oxygenation state of the cell. Once HIF1 α forms a heterodimer with its partner, HIF1^β, the transcriptional activity is further regulated by cofactors, such as CREB binding protein (CBP) or factor-inhibiting HIF (FIH). ARD, acetyl transferase.



Figure 2. Seven points of regulation of tumour oxygenation

Seven features are often present in tumours, contributing in a multifactorial and interrelated fashion to hypoxia. **a** | Intravascular partial pressure of O_2 (pO₂) drops as oxygen is unloaded from haemoglobin in red blood cells (RBCs) as they traverse distally from feeding arterioles; this is termed a longitudinal oxygen gradient. The distance that oxygen can diffuse radially from a vessel is dependent on how much oxygen is present in the vessel. The extent and severity of hypoxia surrounding blood vessels will become more severe as intravascular pO₂ drops. $\mathbf{b} \mid$ At the extreme, red blood cells lose virtually all oxygen and there is no oxygen available to diffuse out into the tumour. c | Shunt flow diverts blood around the tumour, stealing nutrients from the tumour bed. \mathbf{d} | Low vascular density creates hypoxia in extravascular tumour tissue because of limitations on the diffusion distance of oxygen once it leaves the blood vessel. e | Blood vessels that are haphazardly oriented will be less efficient in supplying adequate oxygen to all regions of tissue. **f** | Oxygen consumption rate is the most dynamic feature of oxygen transport in tumours. Small changes in demand for oxygen create large changes in the extent and severity of hypoxia, because higher demand not only limits the diffusion distance of oxygen, it also more severely depletes vessels of oxygen, thereby exacerbating longitudinal oxygen gradients¹⁵⁵. \mathbf{g} Intravascular hypoxia reduces red blood cell deformability, increasing blood viscosity, which in turn reduces flow rate⁹⁶. Relative oxygen concentrations are colour-coded. The direction of blood flow is from the red vessel(s) to the blue vessel(s) as indicated by the grey arrows. These diagrams provide a general pictorial depiction of these basic features. In reality, within any tumour there are microregional variations in oxygenation, with oxygen concentration decreasing radially from microvessels (FIG. 3).



Figure 3. Composite model for the effect of cycling tumour hypoxia on radial diffusion of oxygen In this figure, the effect of four different types of change in red cell flux (the primary determinant of oxygen delivery) on the location and severity of hypoxia are depicted within a tumour microregion. a | The baseline case depicts balanced red cell fluxes in all microvessels. The hypoxic zone, defined in this discussion as regions where the partial pressure of O_2 (pO₂) <10 mmHg, is centrally located. **b** | If the red cell flux drops by an equal proportion in all microvessels, the size and severity of the hypoxic zone will increase, but it will still be centrally located. Spontaneous fluctuations in red blood cell flux within small networks such as this have been shown to change the oxygen gradient, creating a switch from **a** to **b**, on a timescale of 1–3 cycles per hour⁸¹. **c** | Vasodilation of a downstream vessel steals red blood cells from the bottom segment, because flow resistance is much less in the dilated vessel. This change will shift the hypoxic zone downward. This may occur during the process of ongoing angiogenesis. d | Vascular remodelling (here indicated by intussusception) changes the flow resistance in the split downstream vessel, diverting flow upstream. This will reduce the hypoxic zone and shift the gradient upward. If the pO₂ of the blood entering the microregion is higher overall, then the effects of these shifts will be less apparent on the tumour cells in terms of severity of hypoxia. If the pO_2 of the blood entering the microregion is lower, then the severity will be greater. Vascular remodelling and/or variations in shunt flow can also alter the extent and severity of hypoxia from day to day, by changing the oxygen delivery to the tumour microregion.



Figure 4. Models for role of hypoxia in initiation and acceleration of angiogenesis

The vascular crisis model was hypothesized by Holash *et al.* to explain how hypoxia leads to angiogenesis initiation. It is characterized by an early overexpression of angiopoietin 2 (ANGPT2) in the absence of vascular endothelial growth factor (VEGF), which leads to vascular regression and hypoxia. Once hypoxia occurs, VEGF is upregulated, promoting initiation of angiogenesis¹⁰⁹. The acceleration model was hypothesized by Cao *et al.*⁹⁷. In this model, hypoxia is not responsible for initiation of angiogenesis; rather it is driven by non-hypoxia-mediated mechanisms, such as VEGF upregulation by oncogenes. Once angiogenesis is initiated, proliferation of tumour cells occurs, creating hypoxia. Hypoxia-inducible factor 1 (HIF1) is then upregulated to accelerate angiogenesis. Both models incorporate the concept of cooption before angiogenesis and both emphasize the importance of hypoxia in robust and disregulated angiogenesis. The two models converge on a cycle of unchecked angiogenesis, keyed by instability in perfusion, reactive oxygen species, hypoxia and persistent HIF1 activity. Further work is needed to establish how accurate these models are in both primary and metastatic tumour models.



Figure 5. Mechanisms for hypoxia-inducible factor 1 (HIF1) upregulation and consequences after radiation therapy

a | Untreated tumours often have hypoxic subregions. In hypoxic regions, stress granules form containing HIF1-mediated transcripts, sequestering the transcripts from being translated into protein. **b** | After radiation treatment, better oxygenated cells die and there is an increase in perfusion, leading to reoxygenation of the previously hypoxic cells. Stress granules disaggregate, releasing HIF1-regulated mRNAs, which can then go on to be translated into protein. Reoxygenation also causes hypoxia–reoxygenation injury, causing byproduction of reactive oxygen species (ROS), which stabilize HIF1 α , even in the presence of improved oxygenation. **c** | Macrophages are attracted to the dying tumour cells, become activated and release nitric oxide (NO), which also stabilizes HIF1 α . **d** | The increase in HIF1 activity increases vascular endothelial growth factor (VEGF) levels, promoting endothelial cell survival, angiogenesis and tumour cell survival and proliferation. This is highly simplified. Reoxygenation does not completely eliminate hypoxia and there are oxygen gradients within tumours as opposed to categorical differences between aerobic and hypoxic cells.