Chapter 3 Hypoxia in Head and Neck Cancer

Mahvash Tavassoli and Yae-eun Suh

Abstract Head and neck cancer is the sixth most common malignancy worldwide and a major cause of death from cancer, with a median 5-year survival of around 50% (Suh et al. 2014). Over 90% are squamous cell carcinomas arising from the epithelial cells that line the mucosal surfaces of the head and neck. Tumour hypoxia is an important negative prognostic factor for head and neck squamous cell carcinomas (HNSCC), associated with resistance to radiotherapy and decreased overall survival (Bittner and Grosu 2013). Therefore, tumour hypoxia is an important phenomenon in the management of HNSCC and has been the focus of studies to improve treatment response for many years. This chapter aims to discuss some of the key mechanisms affected by tumour hypoxia and review strategies under investigation towards better detection and modulation of tumour hypoxia in order to improve treatment response and survival of head and neck cancer patients.

3.1 Background

Tumour heterogeneity due to genetic and epigenetic variations within the tumours is an important contributor to phenotypic diversity of tumours and resistance to therapy (Catenacci 2014). In addition, tumour microenvironment influences phenotypic diversity or plasticity within the tumour (Fukumura and Jain 2007). Hypoxia is a fundamental biological phenomenon of solid tumours, which plays an important role in changing tumour microenvironment (Wouters et al. 2002). Hypoxia leads to cellular individuality and is associated with resistance to curative therapies and reduced survival in a number of tumour types including head and neck, pancreatic, cervical and prostate cancers (Buffa et al. 2010; Eustace et al. 2013). Tissue hypoxia occurs when the oxygen level in tissues falls below

M. Tavassoli · Y. Suh (🖂)

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Department of Molecular Oncology, King's College London, London, UK e-mail: yae-eun.suh@kcl.ac.uk

M. Tavassoli e-mail: mahvash.tavassoli@kcl.ac.uk

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physiological levels resulting from the inadequate supply of oxygen to meet demand (Hockel and Vaupel 2001). It arises in tumours due to the high metabolic demand of rapidly proliferating cancer cells, and due to disruption in the blood supply. Tumours develop aberrant and chaotic microvasculature, which are structurally immature and leaky, resulting in ineffective oxygen delivery (Vaupel 2004). Fluid accumulation and increased tumour interstitial pressure restricts and compresses intratumoural blood vessels, further reducing the blood supply. Hypoxia leads to increased risk of tumour progression, metastases, resistance to anti-cancer therapies and recurrence due to a wide range of hypoxia-related consequences, resulting in metabolic and genetic changes that promote an aggressive tumour phenotype (Vaupel and Mayer 2007). Hypoxia drives tumour progression by selecting clones that can best adapt to the stress of inadequate perfusion and nutrient deprivation, which then expand and develop into a more malignant phenotype.

Hypoxia has been extensively investigated in HNSCC and its association with poor outcome is well known. However, it is difficult to assess, as intratumoural hypoxia is heterogeneous and dynamic, due to continuing tumour and vessel growth, and constant fluctuations in blood flow. Different biological effects are triggered at different levels of oxygen partial pressure (pO_2) in tissues. There is no consensus for hypoxic thresholds in tumours but at $pO_2 < 25-30$ mmHg radiosensitivity progressively decreases (Hockel and Vaupel 2001; Rademakers et al. 2008), and at $pO_2 < 10-15$ mmHg changes in gene expression under the control of the hypoxia regulated transcription factor HIF-1 are demonstrated. At $pO_2 < 10 \text{ mmHg}$ Vaupel et al. (1994) demonstrated intracellular acidosis and adenosine triphosphate (ATP) depletion in murine fibroblasts and this level represents a critical threshold for energy metabolism. There is also decreased protein synthesis and oxygen consumption to increase tolerance to hypoxia. At $pO_2 < 1$ mmHg there is increased apoptosis, reduced oxidative phosphorylation and cells switch to glycolysis (also known as Warburg effect) to maintain adequate energy levels. Lower levels of pO₂ of 0.2–1 mmHg lengthen G₁ phase of the cell cycle or arrest cells in G₁ and anoxia causes immediate cell cycle arrest (Hockel and Vaupel 2001).

Importantly, hypoxia is a transportable property. Studies performed using patient derived xenografts with differing degrees of hypoxia demonstrated that the amount of hypoxia in the xenograft derived from hypoxic tumours had similar oxygen distribution to the patient's tumour. The hypoxic tumours grew more rapidly and when these human derived xenografts were serially propagated, they maintained the hypoxic feature (Chang et al. 2011). These findings suggest that the genetics of cancer influences the microenvironment within which the tumours are formed, the microenvironment determines tumour behaviour, and together they impact cancer phenotype.

3.2 Classification

Hypoxia can be broadly divided into two basic types: chronic and acute. However, there are no clear cutoff between the two types and there are typically mixed heterogeneous patterns throughout the tumour.

3.2.1 Chronic Hypoxia

Chronic, or diffusion-limited hypoxia, was first described by Thomlinson and Gray in 1955, and is caused by consumption of oxygen by cells close to the blood vessels, leaving an inadequate supply of oxygen and nutrients to the cells further away from the vessels. A distance of approximately 180 μ m from vascularised stroma was shown to be the diffusion distance of soluble oxygen (Thomlinson and Gray 1955). It can also be caused by reduced oxygen content in the blood, such as in anaemia, which may be caused by tumour-related factors as well as anti-cancer therapy or carboxyhaemoglobin formation in heavy smokers, and by compromised perfusion of leaky microvessels. Chronic hypoxia usually occurs after several hours to days of hypoxia, long enough to induce changes in gene expression (Bayer et al. 2011).

3.2.2 Acute Hypoxia

Acute hypoxia, otherwise known as cycling, transient, intermittent, or perfusion-limited hypoxia, occurs because of fluctuations in the perfusion of tumour vasculature. This results in time periods of better or worse oxygenation of tumour areas. This type of hypoxia may be ischaemic due to transient occlusion of blood flow by blood clots or tumour cells, or hypoxaemic due to transient reduction in the oxygen content within microvessels (Harris 2002). Acute hypoxia may have greater effects than chronic hypoxia on the development of an invasive phenotype with hypoxia-reoxygenation episodes resulting in genomic instability, accelerated growth and metastases (Rofstad et al. 2007).

3.3 **Biological Consequences**

Tumour cells exposed to hypoxia activate many signalling pathways, which act in an integrated network affecting common downstream pathways. They work together to ensure adaptation to overcome the lack of oxygen and nutrients. They induce changes such as angiogenesis, glycolysis, inhibition of apoptosis and downregulation of cell adhesion molecules resulting in tumour cell detachment, all of which lead to the development and selection of more aggressive clones of tumour cells.

3.3.1 Metabolism

Glucose is the main source of energy for cells and enters the cell through a family of glucose transporters (GLUT1 to 4). Energy in the form of ATP is generated under aerobic conditions via glycolysis or the tricarboxylic (TCA) cycle. During glycolysis, glucose is metabolised to pyruvate in the cytosol to produce 2 ATPs from each molecule of glucose. Pyruvate enters the mitochondria and is oxidised by pyruvate dehydrogenase (PDH) to produce acetyl coenzyme A (CoA), which enters the TCA cycle. Acetyl CoA is metabolised through a series of reactions generating nicotinamide adenine dinucleotide hydrogen (NADH), a reducing agent used by the oxidative phosphorylation pathway in mitochondria. NADH passes electrons derived from the TCA cycle to the electron transport chain (ETC) in mitochondria, and combines with oxygen to produce water and a proton gradient that is used to generate 36 ATPs per glucose. In the mitochondria ETC, electrons transfer steps are catalysed at Complex I (NADH dehydrogenase or ubiquinone oxidoreductase) and III (coenzyme Q or cytochrome c reductase) and is transferred to O₂ at Complex IV (cytochrome c oxidase). Each step is coupled with proton translocation across the inner mitochondrial membrane, driving ATP synthesis at Complex V. Electrons can escape the ETC and be captured by O₂, forming reactive oxygen species (ROS) and resulting in oxidative stress (Sabharwal and Schumacker 2014).

Under anaerobic conditions the lack of oxygen to act as the final electron acceptor at complex IV in the ETC prevents aerobic metabolism from progressing. Pyruvate is not used in the TCA cycle, but converted to lactate in the cytosol by lactate dehydrogenase (LDH) to regenerate nictotinamide adenine dinucleotide (NAD⁺) from reduced NADH, which is required by glyceraldehyde-3-phosphate dehydrogenase (GAPDH) for additional cycles of glycolysis. PDH is phosphory-lated and inactivated by pyruvate dehydrogenase kinase (PDK) (Lu et al. 2002).

Cancer cells have a high rate of glucose consumption even in the presence of oxygen, known as the Warburg effect or aerobic glycolysis (Warburg et al. 1927). ATP production via glycolysis is much faster but less efficient and cancer cells avidly consume glucose to meet their increased energy and biosynthesis needs. A constant high rate of glycolysis in tumour cells uses up the NAD⁺ pool and glycolysis cannot be sustained unless NAD⁺ is regenerated, which is achieved by increasing lactate production.

3.3.2 Regulation of Hypoxia Driven Pathways

The underlying mechanisms that promote hypoxia and influence the level of hypoxia in tumour are currently unclear. There are two fundamentally different concepts for hypoxia driven pathways. One concept describes that hypoxia provides an environment for the selection of mutant cancer cells that tolerate hypoxic conditions (Graeber et al. 1996). Further mutations allow the cell to develop tolerance

mechanisms, giving rise to a population of cells resistant to the hypoxic microenvironment. One such mutation is the p53 gene, which has been shown to play an important role in tolerance to hypoxia; when p53 mutant cells were mixed with wild type cells and grown under hypoxic condition, only the p53 mutant cells could be selectively grown (Sermeus and Michiels 2011). The second mechanism is adaptation of normal physiological responses to hypoxia, for example by inducing factors such as hypoxia-inducible factors (HIFs) (Wouters and Koritzinsky 2008). There are a large number of oxygen sensitive enzymes and signalling pathways that are present in all cells, and hypoxia reduces the activity of enzymes that use oxygen as co-factor and leads to changes in cellular signalling.

The best understood pathway is hypoxia-inducible factor (HIF) pathway, which is stabilised due to the inhibition of the PHD (Prolyl Hydroxylase Domain oxygen sensor) enzymes (Rabinowitz 2013). The two other major cellular pathways which have been shown to be influenced by hypoxia include the kinase mammalian target of rapamycin (mTOR) pathway, which is an important moderator of metabolic signals, and the unfolded protein response (UPR), a cellular stress response related to the endoplasmic reticulum (Wouters and Koritzinsky 2008). Although they are activated independently a number of studies suggest that HIF, mTOR and UPR mediated responses to hypoxia act in an integrated fashion, both influencing each other and the common downstream pathways that affect gene expression, metabolism, cell survival, tumourigenesis, tumour growth and therapy resistance.

Below we briefly review the importance of these pathways in relation to hypoxia.

3.3.2.1 Hypoxia-Inducible Factor-1 Signalling Pathway

Tumour cell responses to hypoxia are initiated through activation of the hypoxia-inducible factor (HIF) family of transcription factors. The HIF family comprises the HIF-1 α , HIF-1 β , HIF-2 α , and HIF-3 α subunits, the most important of which is HIF-1 (Marxsen et al. 2004) HIF-1 is a heterodimeric protein consisting of an oxygen-regulated α subunit and a constitutively expressed oxygen-independent β subunit. The β subunit of HIF is a constitutive nuclear protein present in normoxic cells.

Under conditions of normal oxygen HIF-1 α is rapidly degraded by the Von-Hippel-Lindau (VHL) tumour suppressor protein via the ubiquitin-proteasome pathway. The α subunits of HIF are hydroxylated at conserved proline residues P402 and P564 by propyl hydroxylase (PHD), which use oxygen as a co-substrate, resulting in recognition and ubiquitination by the VHL E3 ligase, labelling them for degradation by the proteasome. HIF-1 α accumulation is also prevented under normoxia by factor inhibiting HIF-1 (FIH), which hydroxylates the transactivation domain of HIF-1 α . This inhibits the binding of CREB binding protein (CBP) and p300 transcription co-factors to the HIF-1 complex (Fig. 3.1) (Semenza 2000).



Fig. 3.1 HIF-1 activation pathway: **a** Under normoxia HIF-1 α is hydroxylated by propyl hydroxylase and is degraded by VHL via the ubiquitin-proteasome pathway. **b** Under hypoxia propyl hydroxylase and FIH are inhibited and HIF-1 α is stabilised, allowing heterodimerisation with HIF-1 β and subsequent binding to the hypoxia-responsive elements in target genes. This results in the upregulation of proteins that promote cell survival. HIF-1 α can expression can also be upregulated by EGFR and PI3 K signaling pathways. Figure adapted from Rademakers et al. (2008). Reproduced with permission from Elsevier

Hypoxia inhibits hydroxylation of HIF-1 α by PHD and FIH, which results in stabilisation and accumulation of HIF-1 α . HIF-1 α translocates to the nucleus where it heterodimerises with constitutively expressed HIF-1 β , and binds to the hypoxia-responsive elements (HREs) in the promotors of target genes. Genes that help overcome hypoxia, such as those involved in cell survival under oxidative stress, erythropoiesis, angiogenesis, glucose metabolism and pH regulation are upregulated, resulting in the upregulation of more than 100 proteins that promote survival and increased aggressiveness of hypoxic tumour cells (Semenza 2000).

Vascular endothelial growth factor (VEGF) is upregulated by HIF-1 α in hypoxia and is the strongest inducer of angiogenesis, stimulating the proliferation of endothelial cells and the formation of new blood vessels. VEGF also acts as a tumour cell survival factor, inhibiting cell apoptosis by inducing the anti-apoptotic protein Bcl-2 (Ferrara et al. 2003).

Under hypoxia HIF-1 α mediates a switch from oxidative to glycolytic metabolism (Lu et al. 2002). HIF-1 α activation affects cellular glucose metabolism by inducing the transcription of genes involved in increased glucose uptake, such as the glucose transporters GLUT1 and GLUT3, and stimulation of glycolytic enzymes that breakdown intracellular glucose. HIF-1a induces lactate dehydrogenase A (LDHA), increasing the conversion of pyruvate to lactate. In addition, HIF-1a activates pyruvate dehydrogenase kinase (PDK), which inactivates pyruvate dehydrogenase (PDH), actively shunting pyruvate away from the mitochondria. This reduces flux through the TCA cycle, reducing delivery of NADH to the ETC. Mitochondrial function is therefore reduced, decreasing oxygen demand in hypoxic cells. In addition HIF-1a induces BCL2/adenovirus E1B 19 kDa interacting protein 3 (BNIP3) expression, which triggers mitochondrial autophagy. decreasing mitochondrial mass and oxygen consumption. ¹⁸HIF-1a also diverts pyruvate to be phosphorylated by hexokinase to glucose-6-phosphate, which is used in anabolic pathways such as the pentose phosphate pathway, by upregulating hexokinase. This pathway generates the precursors of nucleotides and amino acids required for the rapid tumour cell growth and proliferation. Hypoxia also triggers the generation of reactive oxygen species (ROS) by the ETC, which is released into the cytosol where they inhibit PHDs, resulting in stabilisation of HIF-1 α (Chandel et al. 2000). Glycolytic metabolism results in acidosis of the tumour microenvironment due to increased lactate and carbon dioxide levels, which are actively expelled from the tumour cells into the extracellular matrix. HIF-1 α upregulates monocarboxylate transporter 4 (MCT4), which removes lactate from the cell, carbonic anhydrase 9 (CA9), which catalyses the conversion of carbon dioxide released during the pentose phosphate pathway to carbonic acid, and sodium-hydrogen exchanger (NHE1), which maintains an alkaline intracellular pH and acidic extracellular pH. Acidosis induces the secretion of matrix-degrading hyaluronidase and metalloproteinases by tumour-associated fibroblasts, creating a tumour microenvironment favourable for invasion and migration (Fang et al. 2008). In addition it reduces anti-tumour immunity, inhibiting the activity of dendritic cells and T cells, and the efficacy of drug uptake by tumours, further driving tumour progression (Balamurugan 2015).

HIF-1 α activation has many other downstream effects including the expression of several proteins important in epithelial-mesenchymal transition, preparing tumour cells for invasion and migration, such as SNAIL, SLUG, TWIST (Yang et al. 2008), and inhibition of E-cadherin (Krishnamachary et al. 2006). It has also been implicated in the regulation of cancer stem cells, through activation of NOTCH1, and tumour-mediated immune suppression through attenuation of T cell function (Doedens et al. 2010). HIF-1 α is also involved in the activation of tumour-associated inflammatory signalling in conjunction with nuclear factor-kappa B (NF- κ B) (Bruning et al. 2012), which activates genes that promote cell proliferation and cell survival, and STAT3, which also works with HIF1 α to activate VEGF (Jung et al. 2005). In addition, miRNAs are also activated, with direct binding of HIF-1 α to a HRE on the proximal miR-210 promotor (Huang et al. 2009).

Factors other than hypoxia can also activate HIF-1 α , such as pyruvate and lactate, indicating a positive feedback mechanism. HIF-1 α expression can also be upregulated under conditions of normoxia, such as through activation of the

receptors of the tyrosine kinase family including EGFR and HER2/erbB2, activation of the PI3K-AKT-mTOR pathway, and genetic mutations including VHL inactivation, loss of PTEN, p16^{INK4a} and p53 tumour suppressors (Span and Bussink 2015).

3.3.2.2 mTOR Signalling

The adaptive response to hypoxia involves inhibition of energy-intensive cellular processes including protein translation. This effect is mediated in part through a decrease in the kinase activity of mammalian target of rapamycin complex 1 (mTORC1) (Vadysirisack and Ellisen 2012). The mTOR pathway responds to microenvironment signals such as availability of growth factors, energy (ATP) and nutrients including lipids and amino acids. These factors influence upstream kinase signalling pathways that all converge upon mTOR and the outputs of mTOR drive changes in the metabolic rate of the cells (Wouters and Koritzinsky 2008). Activation of mTOR requires oxygenation as one of its inputs and low oxygenation due to hypoxia inhibits signalling through this pathway (Magagnin et al. 2008).

Mild to moderate hypoxia levels suppress mTOR through a HIF-1 independent pathway involving the activation of tuberous sclerosis protein 1(TSC1)-TSC2 complex via adenosine monophosphate-activated protein kinase (AMPK) or via transcriptional regulation of REDD1. Hypoxia reduces the phosphorylation and activation of 4EBP1, an important negative regulator of protein synthesis, by mTOR (Magagnin et al. 2008; Rouschop et al. 2011). mTOR is a principle regulator of autophagy under nutrient depleted conditions, which is regulated by BNIP3 and AMPK during hypoxia. In addition, mTOR interacts with HIF-1 α , promoting the rate of HIF-1 α mRNA translation. In tumour cells hypoxia has been shown to be less effective at inhibiting mTOR through mechanisms such as constitutive mTOR activity or loss of function of TSC2 or PTEN, the negative regulator of this pathway (Wouters and Koritzinsky 2008).

The shut off of protein synthesis through mTOR inhibition is important for survival of hypoxic condition. Paradoxically mTOR is an oncogene and many known negative regulators of mTOR such as TSC1, TSC2, LKB1 and PTEN signalling have been shown to function as *bona fide* tumour suppressors, yet hypoxic cells need to shut down this pathway to survive metabolic stress.

3.3.2.3 The Unfolded Protein Response (UPR)

Messenger RNA translation, post-translational modifications and protein folding takes place in the endoplasmic reticulum (ER). The folding of these proteins in ER requires disulphide bind formation which is an oxidation reaction (Nagelkerke et al. 2013).

Under conditions of stress, such as severe hypoxia, misfolding of proteins occur which accumulate in the ER, resulting in ER stress. This activates the unfolded protein response, which aims to relieve this stress and increase cell survival by activating the ER stress sensors such as protein kinase R-like endoplasmic reticulum kinase (PERK). PERK activation results in inhibition of global mRNA translation and protein synthesis to reduce the load of misfolded proteins and lower energy demands. It also upregulates the expression of genes involved in resistance to oxidative stress and amino acid metabolism to promote cell survival, and enhances HIF-1 α transcriptional activity. ER-associated degradation and autophagy occur to remove misfolded proteins, further reducing ER stress (Pereira et al. 2014). ER stress also modulates the expression of VEGF, stimulating angiogenesis and promoting cell survival of rapidly growing cells. In addition VEGF signalling can activate the UPR in endothelial cells in the absence of ER stress via mTOR signalling, further promoting angiogenesis (Wang et al. 2012). The UPR is upregulated in tumours and is an important survival response in conditions of prolonged hypoxia, increasing the threshold for apoptosis. Tumours cells with abrogated PERK activity show significantly reduced clonogenic survival and decreased ability to tolerate moderate to extreme hypoxia, with higher levels of apoptosis in hypoxic areas (Bi et al. 2005).

3.3.3 Micro RNAs

Several studies have demonstrated through genome-wide miRNA profiling that miRNA expression in tumour cells is altered in response to hypoxia (Hebert et al. 2007; Hua et al. 2006; Kulshreshtha et al. 2007; Guimbellot et al. 2009; Bruning et al. 2011; Crosby et al. 2009). However due to differences in tumour cell lines, variations in exposure to hypoxia severity and duration, as well as differences in microarray hybridisation platforms resulting in different ranges of miRNAs screened, there has been limited overlap in the patterns of miRNA up- and downregulation. Next generation sequencing techniques have revealed a greater number of previously not identified miRNAs, highlighting the complexity of miRNA response to hypoxia (Camps et al. 2014).

MiR-210, however, has been robustly and consistently induced by hypoxia, in both normal and cancer cells. It has also been shown to be overexpressed in a variety of solid tumours and patient plasma or serum, including HNSCC, where it is generally associated with poor clinical outcome (Wang et al. 2014; Gee et al. 2010). MiR-210 is maintained at low levels in normoxia. Under hypoxic conditions miR-210 is induced independently and directly by HIF-1 α through interaction with its promotor sequence, and targets genes involved cell cycle regulation, mitochondrial function, apoptosis, angiogenesis and metastases. MiR-210 promotes stabilisation of HIF-1 α and inhibit PHDs preventing the degradation of HIF-1 α (Chang et al. 2013).

MiR-210 targets the MYC antagonist MAX network transcriptional repressor (MNT), resulting in MYC activated cell cycle progression, and the cell cycle regulator E2F transcription factor 3 (E2F3), promoting proliferation and inhibiting apoptosis (Zhang et al. 2009). Apoptosis is also inhibited by targeting apoptosis-inducing factor mitochondrion-associated 3 (AIFM3), reducing caspase-dependent apoptosis, and caspase-8-associated protein 2 (CASP8AP2), an integral protein in the cleavage and activation of apoptosis initiator caspase-8 (Kim et al. 2009). MiR-210 has been found to bind to two sites in the 3'UTR of RAD52 mRNA, resulting in degradation. RAD52 is involved in the repair of double stranded DNA breaks and therefore miR-210 suppression of RAD52 increases genetic instability (Crosby et al. 2009). MiR-210 has a repressive effect on the mitochondrial ETC, aiding the metabolic shift to glycolysis in hypoxia. It targets and suppresses several integral components of the mitochondrial ETC such as cytochrome oxidase assembly protein-10 (COX-10), succinate dehydrogenase subunit D (SDHD), compromising electron transport and promoting glycolysis (Puissegur et al. 2011). In addition miR-210 suppresses iron-sulphur cluster scaffold proteins (ISCU), which catalyse the assembly of iron-sulphur clusters that are critical for enzymes in the TCA cycle and for the function of ETC complexes I, II and III (Chan et al. 2009). Angiogenesis is stimulated by the suppression of ephrin-A3 (EFNA3), which stimulates normoxic endothelial cell to form capillary-like structures (Fasanaro et al. 2008). In addition vacuole membrane protein 1 (VMP1), which inhibits cell migration and invasion, is suppressed, inducing a more metastatic phenotype (Ying et al. 2011).

MiR-210 is also reported to have a tumour suppressor role, inducing cell cycle arrest through fibroblast growth factor receptor-like 1 (FGFRL1) (Zuo et al. 2015) and apoptosis through suppression of anti-apoptotic Bcl-2 expression (Chio et al. 2013). As hypoxia can influence both cell death and survival the role of miR-210 is likely to depend on the cell type, extent and duration of exposure to hypoxia, and reflect the adaptation of tumour cells to a hostile environment.

3.4 Clinical Consequences in HNSCC

3.4.1 Resistance to Radiotherapy

Radiosensitvity rapidly decreases when the pO_2 in a tumour is less than 25–30 mmHg (Hockel and Vaupel 2001). In 1953 Gray et al. (1953) determined the radiation dose required to achieve the same biological effect is 2.8–3 times higher in the absence of oxygen than presence. When ionising radiation is absorbed in tissue, free radicals are produced as a result of ionisations either directly in the DNA itself, or indirectly in other molecules such as water. Free radicals break chemical bonds and initiate a chain of events that results in DNA damage. Oxygen molecules are able to react with the free radicals and create a stable change in the chemical composition of the DNA damage. Oxygen therefore enhances the damage by

"fixing" or making the damage permanent. In the absence of oxygen the target is chemically restored and the damage is repaired. In order for oxygen to consolidate radiation-induced damage it needs to be present at the time of irradiation or within a few milliseconds (Howard-Flanders and Moore 1958). Other hypoxia-induced mechanisms also contribute to radioresistance including glycolytic metabolism and lactate accumulation, which act as antioxidants and scavenge free radicals (Brizel et al. 2001; Sattler et al. 2010).

The changes in hypoxia heterogeneity during a course of fractionated radiotherapy (RT) can affect the response. During the early stages of treatment cell density decreases as the well-oxygenated tumour cells are killed. This results in reduced intratumoural pressure and increased vascular density leading to tumour reoxygenation (Bussink et al. 2000). This reoxygenation does not increase HIF-1 α degradation, but the increase in ROS formation stabilises HIF-1 α and increases the expression in a hypoxia-independent manner (Harada et al. 2009). This, together with accelerated repopulation that occurs at the later stages of RT, means that the entire cumulative dose of RT needs to be delivered.

3.4.2 Resistance to Chemotherapy

Some chemotherapeutic agents, such as etoposide, are dependent on the presence of oxygen for maximum cytotoxic effect (Wozniak et al. 1984). The blood vessels in hypoxic tumours are disordered and leaky, preventing efficient delivery of chemotherapy drugs. Hypoxic cells are distant from blood vessels and therefore not adequately exposed to drugs (Tannock 1998). Alkylating agents, such as cisplatin, and anti-metabolites, such as 5-FU, act during DNA synthesis by damaging DNA and initiating apoptosis. Cellular proliferation and DNA synthesis decreases as a function of distance from blood vessels and hypoxia selects for cells that have lost sensitivity to p53 mediated apoptosis, reducing the cytotoxic effect. Hypoxia upregulates genes involved in drug resistance, such as multidrug resistant 1 (MDR1) leading to the increased expression of P-glycoprotein, which is associated with tumour resistance to chemotherapeutics (Comerford et al. 2002). The acidic tumour microenvironment creates a pH gradient between the tumour cell and extracellular space, inhibiting the accumulation of drugs that are weak bases, such as doxorubicin (Skovsgaard 1977).

3.5 Therapeutic Approaches to Overcome Hypoxia Radioresistance in HNSCC

Hypoxia represents a potential target for therapy. Various strategies have been used to overcome hypoxia in HNSCC, summarised in Table 3.1. A meta-analysis of 32 clinical trials of various strategies of hypoxic modification demonstrated a small but

Method	Example	
Increasing oxygen availability	Hyperbaric oxygen ARCON	
	Blood transfusion, erythropoeitn	
Radiosensitisers	Nimorazole	
Hypoxia cytotoxin	Tirapazamine	
Increasing radiotherapy dose	Dose painting	
Other	Anti-angiogenic agents with RT HIF-1 targeting	

Table 3.1 Summary of methods investigated to overcome hypoxic radioresistance

significant improvement in locoregional control (LRC), disease specific control (DSS) and overall survival (OS) in HNSCC treated with primary RT (Overgaard 2011). Hypoxia stratification is required to identify patients at risk of treatment failure but also patients with normoxic tumours, who do not gain benefit from additional treatment, and can avoid additional toxicity.

3.5.1 Improving Haemoglobin (Hb) Levels

Many studies have demonstrated that low haemoglobin level prior to treatment associated with significant reduction in survival and increase in locoregional failure in HNSCC after RT or chemoradiation (Fortin et al. 2008; Lee et al. 1998). Over 1000 patients pooled from two studies demonstrated that high Hb was significantly associated with higher OS and the administration of blood during therapy increased Hb level. However blood transfusions had no impact on LRC, DSS or OS (Hoff et al. 2011). Exogenous erythropoietin administration to stimulate red blood cell production and correct anaemia has been investigated. However this did not improve tumour control or survival as demonstrated in 2 meta-analyses, but suggested an increase in mortality (Bohlius et al. 2009; Lambin et al. 2009).

3.5.2 Increasing Oxygen Availability

Hyperbaric oxygen is the administration of 100% oxygen at higher than normal atmospheric pressure. This elevated pressure results in increased pO_2 in tissues and has been delivered in hyperbaric oxygen chambers with RT. This strategy has been shown to significantly improve tumour control and mortality, but also causes an increased rate of severe radiation tissue injury and chance of oxygen toxic seizures during therapy (Bennett et al. 2008).

Carbogen (95% oxygen and 5% carbon dioxide) breathing before RT did not improve tumour control (rubin 1979 carbogen), but combined with accelerated radiotherapy and nicotinamide (ARCON, accelerated radiotherapy with carbogen and nicotinamide) has been shown to result in high LRC in advanced larynx and oropharynx SCC (Kaanders et al. 2002). This combination aims to limit clonogenic repopulation during therapy by reducing the overall duration of RT by delivering more than one fraction a day, decrease diffusion-limited hypoxia with inhalation of carbogen, and decrease perfusion-limited hypoxia by administering nicotinamide, a vasoactive vitamin B_6 analogue. A more recent study compared accelerated RT with ARCON in larynx SCC and did not demonstrate a benefit. However, a subgroup of patients had their hypoxic status assessed with pimonidazole IHC, which stains regions of hypoxia, and significant improvement in regional control and disease free survival (DFS) was found in the high pimonidazole staining group treated with ARCON (Janssens et al. 2012).

3.5.3 Radiosensitisers

Cellular sensitivity to RT can be improved by increasing the apparent cellular pO_2 levels using radiosensitisers. Nitromidazoles are chemical compounds that mimic the radiosensitising effect of oxygen by inducing and stabilsing free-radical mediated double-stranded DNA breaks. The benefit of the addition of a nitroimidazole compound to RT was carried out in many studies, without clinical benefit (Lee et al. 1995; Van den Bogaert et al. 1995). However these studies used the older generation of the drug which is limited by neurotoxicity. Overgaard (1994) carried out a meta-analysis of over 7000 patients treated with nitroimidazole-based radiosensitisers in various tumour types of unknown hypoxia status and found an improvement in LRC and OS, with the greatest benefit in HNSCC. This was followed by the DAHANCA 5 study which randomized over 400 HNSCC patients to receive RT with nimorazole or RT with placebo. LRC and DSS were significantly higher with the addition of nimorazole (Overgaard et al. 1998). A phase II study investigated accelerated RT with nimorazole and cisplatin in 227 patients with locally advanced HNSCC and recently reported a high 5-year OS of 75% (Bentzen et al. 2015).

3.5.4 Hypoxic Cytotoxins

This group of drugs, of which tirapazamine is the most widely investigated, have a direct cytotoxic effect on hypoxic cells. Tirapazamine (TPZ) is a prodrug that requires one-electron reduction to a radical by intracellular reductases. In normoxia the unpaired electron in the radical is rapidly transferred to molecular oxygen, forming superoxide and regenerating the initial prodrug. In hypoxic conditions the radical accumulates and induces DNA damage, in part due to poisoning of topoisomerase II (Brown 1993).

TPZ combined with cisplatin and RT in HNSCC has shown to be well tolerated in phase I and II studies (Lee et al. 1998; Rischin et al. 2001). Rischin et al. carried out a phase II randomised trial of this combination versus cisplatin and RT and demonstrated a trend towards improved 3-year survival with acceptable toxicity (Rischin et al. 2005). However, further studies have not shown a benefit in survival or quality of life from the addition of TPZ (Rischin et al. 2010). Rischin et al. also carried out a substudy of 45 patients who had their tumour hypoxic status assessed by ¹⁸F-fluoromisonidazole, a hypoxic radiotracer, PET scans. Hypoxic patients performed poorly in the absence of TPZ augmentation, whereas the rate of complete response in the hypoxic group receiving TPZ was nearly as high as the rate for normoxicchemoRT patient. Patients with hypoxic tumours were more likely to develop distant metastatic disease compared with normoxic patients (Rischin et al. 2006).

3.5.5 Hypoxia Image-Guided Radiotherapy

Intensity modulated radiotherapy (IMRT) is used in the treatment of HNSCC. The intensity of the radiation beam can be modulated to reduce doses to normal structures without compromising the doses to the tumour. Tumour recurrence after compete response to radiotherapy has been shown to occur predominantly in high radiation dose regions, implying that tumours have radioresistant subvolumes, potentially due to hypoxia, within this region (Hendrickson et al. 2011). IMRT dose painting is a method of assigning different dose levels to structures within the same treatment fraction, resulting in the potential for higher total doses to selected targets (Galvin and De Neve 2007). Dose painting requires functional imaging such as positron emission tomography to direct regions for dose escalation. Intensification of intratumoural subvolumes of hypoxia has been investigated as a strategy to improve outcome in RT planning studies and radiobiological modelling studies only as this requires accurate and stable identification of hypoxia to guide target volume delineation.

3.5.6 Targeted Therapies

3.5.6.1 Targeting Angiogenesis: VEGF Receptors

High VEGF expression levels has been associated with worse overall survival in HNSCC (Kyzas et al. 2005). As tumour vasculature is abnormal, increased VEGF expression and microvascular density does not lead to increased blood flow and oxygen delivery. Targeting VEGF is a strategy to improve oxygenation by normalising the tumour vasculature and thereby increase response to RT. RT can lead

to stimulation of angiogenesis through upregulation of VEGR and therefore anti-angiogenic therapy has been tested with RT (Koukourakis et al. 2001). Bevacizumab is a humanised monoclonal antibody against VEGF approved for use in colorectal cancer, and has been investigated in phase II studies concomitant with chemoRT in HNSCC with encouraging results (Fury et al. 2012). It inhibits endothelial cell growth and function, disrupting the formation of capillary networks, and has been shown to enhance RT response in HNSCC xenograft models (Hoang et al. 2012). Anti-angiogenic agents can improve tumour oxygenation by reducing the number of oxygen consuming endothelial and tumour cells, and the number of inefficient blood vessels. This also results in less leaky vessels and decreased interstitial pressure, improving perfusion (Willett et al. 2004).

3.5.6.2 Targeting HIF-1

Targeting HIF-1 is an attractive therapeutic strategy to overcome the effects of hypoxia. Drugs that inhibit HIF-1 activity are already in clinical use, such as the topoisomerase I inhibitor topotecan and mTOR inhibitors which decrease HIF-1 synthesis, the anthracycline doxorubicin which suppresses HIF-1 DNA binding, and the proteasome inhibitor bortezomib, which reduces HIF-1 transactivation (Semenza 2010). Newer drugs have been developed, such as PX-478 that inhibits HIF-1 by inhibiting HIF-1 mRNA translation and HIF-1 dequbiquitination and is being investigated as a phase I trial. In HNSCC tumour models this drug provided significant radiosensitisation of hypoxic cell lines and xenografttumours (Schwartz et al. 2009).

3.6 Methods to Detect Hypoxia in HNSCC

Accurately detecting and quantifying hypoxic tumours is essential in identifying patients who have aggressive, treatment resistant disease and has been one of the limiting factors for translating hypoxia-modification strategies into routine practice.

3.6.1 Clinical Studies Using Eppendorf polarographic Needle Electrodes

Approximately 50–60% of human tumour contain hypoxic and/or anoxic tissue regions that are heterogeneously distributed within the tumour (Vaupel et al. 2007). Many clinical studies, mainly in HNSCC and cervix cancers, using Eppendorf polarographic needle electrode histography have directly demonstrated that tumour

hypoxia predicts for decreased local control, increased disease recurrence and reduced overall survival. This technique involves inserting an electrode into multiple sites in a tumour and measuring the pO_2 at several points per needle track and data from all tracks form a histogram. In head and neck cancers the pO_2 measurements between the primary tumour and metastatic lymph nodes have shown little difference, and nodal measurements have been used to represent the hypoxic status of the patient (Becker et al. 1998).

In head and neck cancer, Gatenby et al. (1988) in 1988 measured the oxygen tension in 31 lymph node metastases from HNSCC patients, and demonstrated a significant relationship between low mean pO₂ measurements and poor response to radiotherapy. The volume of tumour containing low oxygen levels was also found to be important. Nordsmark et al. (1996) measured the pretreatment oxygenation status initially in 34 lymph nodes and 1 primary using the Eppendorf polarographic needle electrode histography and evaluated the tumour oxygenation status as the percentage of pO₂ values \leq 2.5 mmHg (hypoxic fraction 2.5, HF2.5). The median HF2.5 of 15% was used to define hypoxia and these patients had significantly poorer locoregional control (LRC). This was confirmed by the same group in a further 35 patients (Nordsmark and Overgaard 2000). Brizel et al. (1997, 1999) assessed 63 patients for pretreatment tumour oxygenation using the primary site or metastatic lymph node. A preliminary study had shown significantly worse disease free survival (DFS) at 12 months for patients with a median $pO_2 < 10$ mmHg, and this was used to define hypoxic tumours. Whether the measurement was taken at the primary or nodal site did not affect the DFS and hypoxic tumours had significantly worse 3 year LRC, DFS and overall survival (OS), independent of tumour stage. However, a small study of 25 patients did not find an association between pretreatment oxygen levels and LRC or OS (Adam et al. 1999).

Stadler et al. (1999) identified the importance of the hypoxic subvolume, defined as the percentage of pO_2 values <5 mmHg multiplied by the total tumour volume. Fifty-nine HNSCC primaries and nodes were assessed and the median pO_2 was 16 mmHg, with no difference between the sites. On multivariate analysis the hypoxic subvolume and the HF5 were significant prognostic factors for survival. This was confirmed in a further 125 patients, where hypoxic tumour volume was a strong and independent prognostic factor for survival (Dunst et al. 2003). Rudat et al. (2000) evaluated the repeatability and predictive value of pO_2 Eppendorph electrode histography in HNSCC. High variability was seen in patients who had two repeated independent measurement of the same tumour. In 41 patients with follow up data, locally advanced HNSCC the fraction of $pO_2 \leq 2.5$ mmHg was a significant prognostic factor for survival.

A joint analysis of prospectively collected Eppendorf pO_2 measurements from multiple centres was performed, which consisted of 397 patients with HNSCC (Nordsmark et al. 2005). Median tumour pO_2 was 9 mmHg and multivariate analysis demonstrated that HP2.5 greater than population median (19%) was associated with poorer OS at 3 years, providing strong evidence that tumour hypoxia has a significant role in HNSCC. Table 3.2 summaries the clinical data from HNSCC.

No. of patients	Outcome	Ref.
31	Mean pO ₂ 20.6 mmHg in complete responders versus 4.7 mmHg in non-responders to radiotherapy	Gatenby et al. (1988)
35	Median pO ₂ 14 mmHg Median HF2.5 > median (15%) significantly worse LRC	Nordsmark et al. (1996)
31	Median pO ₂ 12 mmHg, median HF2.5 30% Median HF2.5 > median (15%) significantly worse LRC	Nordsmark and Overgaard (2000)
63	Median pO_2 4.8 mmHg for 24 primary sites Median pO_2 4.3 mmHg for 39 nodes Median pO_2 < 10 mmHg significantly worse LRC, DFS, OS	Brizel et al. (1999, 1997)
41	Median pO_2 10 mmHg HF2.5 > median (21%) worse OS	Rudat et al. (2000)
397	Joint analysis of prospectively collected data in HNSCC Median tumour pO_2 9 mmHg HP2.5 > median (19%) associated with poorer OS	Nordsmark et al. (2005)
25	Mean pO ₂ 20.2 mmHg No prognostic impact on outcome	Adam et al. (1999)
59	Median pO_2 16 mmHg Hypoxic subvolume > 6 cm3, HF5 > median (30%) worse overall survival	Stadler et al. (1999)
125	Median pO ₂ 9.4 mmHg Hypoxic volume prognostic factor for OS	Dunst et al. (2003)

Table 3.2 Summary of Eppendorf electrode histography studies in HNSCC

Table 3.3 Summary of methods used to detect hypoxia

Method	Example	
Direct detection	Eppendorf oxygen electrode histography	
Exogenous markers	2-nitroimidazole—pimonidazole, EF5	
Endogenous markers		
Tumour protein levels Circulating markers Gene expression signatures	HIF-α1, CA9, osteopontin Osteopontin, miR-210 Hypoxia metagene, 15-gene classifier	
Non-invasive imaging	DCE MRI, BOLD MRI PET— ¹⁸ F-FMISO, ⁶⁴ Cu-ATSM	

The use of Eppendorf oxygen electrode histography has been crucial to demonstrate the adverse effects of hypoxia on outcome in the clinical setting. It has the advantage of directly measuring absolute pO_2 values and has prognostic potential. However, it has not been incorporated into routine clinical practice as it is an invasive procedure restricted to accessible tumours. It also requires an experiences operator but still exhibits large inter-observer variability (Nozue et al. 1997), and only provides information on the areas sampled. In addition it cannot differentiate between tumour and normal tissues (Vaupel et al. 2007). Other methods that have been investigated are summarised in Table 3.3.

3.6.2 Exogenous Markers of Hypoxia

The most commonly used exogenous markers of hypoxia are 2-nitroimidazole compounds such as pimonidazole and pentafluorinatedetanidazole (EF5) (Evans et al. 2000; Raleigh et al. 1998). They are irreversibly bioreduced by cellular nitroreductases and bind to thiol-containing proteins in viable hypoxic cells, forming stable adducts (Varghese et al. 1976). These adducts can then be detected by IHC staining with specific antibodies. IHC staining can quantify hypoxia by visually or digitally estimating the fraction of stained cells and provides hypoxic measurements with high spatial resolution. Binding is commonly seen adjacent to regions of necrosis and at a distance from blood vessels, in keeping with diffusion-limited hypoxia (Evans et al. 2000).

2-nitroimidazoles bind to tissue at a pO_2 level of <10 mmHg and are more sensitive at lower pO_2 levels than microelectrodes (Raleigh et al. 1999). The amount of bound marker is dependent on oxygen, the accumulation rate of individual hypoxic cells and intact nitroreductase enzymes in viable hypoxic cells (Ljungkvist et al. 2007). Differences in the activity of the enzymes can affect level detected and the accumulation rate of pimonidazole is dependent on pH (Dennis et al. 1985). The main disadvantage to exogenous markers of hypoxia is that they have to be administered intravenously prior to tissue biopsy or surgical resection, to allow fixation of the adducts that form in hypoxic regions. Pimonidazole binding has shown good correlation with Eppendorf electrode measurements in mouse models (Raleigh et al. 1999), but not in the clinical setting in cervix cancers (Nordsmark et al. 2003). Neither pimonidazole nor Eppendorf electrode pO₂ were not prognostic for outcome in 127 cervix cancer patients (Nordsmark et al. 2006).

The prognostic value of 2-nitroimidazoles has been shown in HNSCC. High pimonidazole binding in 43 biopsies was significantly associated with LRC and DFS at 2 years. This association was lost in the subgroup treated with ARCON, suggesting a predictive role for pimonidazole (Kaanders et al. 2002). Pretreatment hypoxia using EF5 binding was investigated in 22 HNSCC patients and severe hypoxia, defined as maximum 30% EF binding which approximated to a pO₂ of 0.76 mmHg, was associated with shorter event free survival at 3 years (Evans et al. 2007).

3.6.3 Endogenous Markers of Hypoxia

Endogenous markers are genes or gene products that are specifically upregulated under hypoxic conditions and can be measured on tumour specimens. HIF-1 α , GLUT1 and 3, CA9 and osteopontin are detected by antibody staining and immunohistochemistry and can be assessed on formalin fixed paraffin embedded (FFPE) archival material, allowing correlation with outcomes. However the techniques of tissue processing and staining analysis have not been validated between different laboratories, limiting comparisons and clinical applications. Samples are usually from small biopsy specimens and therefore not representative of the hypoxia heterogeneity of entire tumour. In addition their expression may be affected by the type of hypoxia, for example HIF-1 α is induced and repressed rapidly and reflect acute changes in hypoxia, whereas CA9 is slow to accumulate (Sobhanifar et al. 2005). In general, studies using these markers show conflicting results, and the association between these markers and oxygen status of tissue is weak.

3.6.3.1 HIF-1a

HIF-1 α is overexpressed in a wide variety of tumours. In HNSCC many studies have demonstrated an association with poor survival and poor response to chemoRT (Aebersold et al. 2001; Koukourakis et al. 2002), and has been found to be an independent prognostic factor in HNSCC. However there are also studies that show no association with prognosis (Kyzas et al. 2005; Wachters et al. 2013) and in one study of in surgically resected cohort HNSCC, HIF-1 α expression was associated with significantly better DFS and OS (Beasley et al. 2002).

3.6.3.2 Glucose Transporters (GLUT)

These glucose transporters are present in normal tissue but upregulated in tumours due to increased glucose requirements in anaerobic glycolysis (Rademakers et al. 2011). In HNSCC GLUT-1 has been associated with poor treatment outcome in multiple retrospective series (Mineta et al. 2002; Oliver et al. 2004). In 58 patients treated with ARCON GLUT-1 was associated with an increased rate of distant metastasis and worse OS but GLUT-3 was associated with better LRC (Jonathan et al. 2006). Co-expression of HIF-1 α and GLTU-1 significantly correlated with an increased risk of tumour-related death (Eckert et al. 2011).

3.6.3.3 Carbonic Anhydrase 9 (CA9)

CA9 is a transmembrane enzyme and a downstream target of HIF-1. It catalyses the reversible hydration of carbon dioxide to carbonic acid and is involved in pH regulation. Elevated expression has been demonstrated at $pO_2 < 20$ mmHg (Wykoff et al. 2000). Overexpression of CA9 has been found in different types of cancer including HNSCC. Kaanders et al. (2002) showed that the expression of CA9 in HNSCC demonstrated some overlap in the distribution pattern as pimonidazole, with increasing signal at greater distance from blood vessels, but

overall the correlation was found to be weak and not associate with outcome. Greater CA9 expression was seen at shorter distances from the vessels, suggesting that upregulation may occur at pO_2 levels higher than that required for pimonidazole (Kaanders et al. 2002). CA9 shows conflicting results as a prognostic marker. Strong CA9 expression was found to be related to poor complete response rate after treatment with chemoRT (Koukourakis et al. 2001), but positivity was associated with better LRC and freedom from distant metastases in 58 patients treated with ARCON (Jonathan et al. 2006), or no association with LRC (Nordsmark et al. 2007). In the DAHANCA 5 study patients 320 samples were available for CA9 staining and the expression of CA9 did not correlate with any of the tumour or patient characteristics, was not a prognostic marker and did not correlate with nimorazole treatment (Eriksen et al. 2007).

3.6.3.4 Osteopontin (OPN) in Tissue and Plasma

Osteopontin (OPN) is an extracellular matrix-associated integrin-binding glycoprotein protein induced by hypoxia, initially identified in non-collagenous bone matrix. Binding of OPN to cell surface receptors on tumour cells activates integrins and matrix metalloprotein signalling pathways, increasing the risk of tumour invasion and migration (Chien et al. 2009). It is also involved in angiogenesis, promotes cell survival through PI3K/AKT and JAK/STAS3 signalling and regulation of NF-KB. It is upregulated through AKT activation, independent of HIF-1 under hypoxia (Ahmed et al. 2011). High OPN levels in both tissue and plasma have been correlated with low tumour pO_2 in HNSCC and outcome (Nordsmark et al. 2007). A retrospective study of the DAHANCA 5 trial patients demonstrated that high OPN plasma levels was associated with higher locoregional tumour failure and disease specific mortality (Overgaard et al. 2005). Patients with high OPN levels who received nimorazole had better outcomes compared with patients with high OPN levels who did not receive nimorazole. No effect was seen in the low/intermediate OPN group, suggesting that high concentrations of OPN could predict clinically relevant hypoxia and identify patients who may benefit from hypoxic modification. Lim et al. (2012) investigated the predictive potential of OPN in patients who received tirapazamine but did not find that high levels were associated with poor prognosis and found no interaction between OPN and treatment. The difference in results compared with the DAHANCA study may be due to differences in treatment protocols, but also lack of standardisation for OPN thresholds between the studies.

3.6.3.5 Gene Expression Signatures of Hypoxia

As hypoxia is influenced many biological pathways, a single marker is incapable of adequately describing this complex heterogeneous response. To improve hypoxia

specificity combining several markers in gene expression signatures of hypoxia have been investigated. Different methods have been used to derive clinically applicable hypoxic gene signatures and are summarised in Table 3.4.

In vitro derived gene sets have been described by culturing tumour cell lines under normoxic and hypoxic conditions, then assessing the differences in gene expression. Koong et al. (2000) used FaDu HNSCC and SiHa cervical SCC cell lines to identify 9 genes exhibiting greater than 3-fold induction under hypoxic conditions when using the expression of VEGF as a cut off point for assessing hypoxia-induced genes. An 84 upregulated gene signature was generated by culturing normal and transformed keratinocytes at various different oxygen levels, which could be grouped into 6 functional groups, including as metabolism/ transport, angiogenesis, tissue remodelling, apoptosis, proliferation/differentiation and gene expression (Denko et al. 2003).

Chi et al. (2006) evaluated the gene expression in response to hypoxia in several primary cell lines in vitro using cDNA microarrays and found a wide variation in the response between cell and tissue types. However, a 253 gene signature was identified which were concordant with gene expression data from a distinct subset of renal tumours, allowing classification by hierarchical clustering into 2 groups with high or low expression of the hypoxia response genes. The high expression group consisted of clear cell renal carcinomas, which typically have loss of function al VHL proteins, and the low group were other histological subtypes or normal samples. Similar classification grouped breast cancers into ductal adenocarcinomas (high expression) and fibroadenomas or normal samples (low expression). In addition, the gene signature was a strong predictor of clinical outcome in independent breast and ovarian cancer datasets.

Human mammary epithelial cells were culture under hypoxic conditions at early and late time points to assess the time dependency of hypoxia-regulated gene expression (Seigneuric et al. 2007). The early response gene signature was characterised by genes related to growth, apoptosis, insulin and oestrogen receptor signalling, whereas the late response was characterised by genes involved in angiogenesis, glucose transport, proliferation, metastasis and apoptosis, and were similar to the genes identified by Chi et al. The early but not late signature was prognostic in univariate analysis, but not maintained in multivariate analysis in a breast cancer dataset. Sorensen et al. (2010) exposed 4 head and neck cell lines to different oxygen concentrations at normal or low pH, and analysed the gene expression analysed to identify genes upregulated by hypoxia, independent of pH. Hypoxia induced pH independent genes were selected if they were common in 3 of the 4 head and neck cell lines, resulting in a 27 gene signature.

Winter et al. (2007) developed a hypoxia metagene from 59 HNSCC fresh frozen samples by clustering around the mRNA expression of 10 well-known hypoxia-regulated genes, such as CA9, GLUT1 and VEGF. Strongly correlated upregulated genes appearing in >50% of clusters defined a signature comprising of 99 genes, which was found to be an independent prognostic factor for RFS in an independent head and neck cancer and breast cancer dataset. The same group derived a common hypoxia metagene consisting of 51 genes by selecting genes that

Source of signature	No. of genes in signature	Prognostic validation	Ref.
FaDu, SiHa cell lines in vitro under hypoxia and normoxia	9	No	Koong et al. (2000)
Normal cervical and dermal keratinocytes, normal stromal fibroblasts, SiHa, C33a, FaDu cells	84	No	Denko et al. (2003)
Mammary and renal tubular epithelial cell lines under hypoxia or normoxia for 12 h	253	2 breast cancer, 1 ovarian cancer datasets— prognostic for OS and RFS on univariate analysis; breast dataset—prognostic for OS on multivariate analysis	Chi et al. (2006)
Mammary epithelial cell line under hypoxia; early (1–6 h) or late (12–24 h)	Early 15 Late 93	Early signature prognostic for DSS in breast cancer dataset on univariate analysis only; late signature not prognostic	Seigneuric et al. (2007)
59 HNSCC biopsy specimens Clustering around known hypoxia regulated genes	99 hypoxia metagene	Independent prognostic factor in head and neck for RFS and breast for OS and metastasis free survival	Winter et al. (2007)
In silico generation of hypoxia co-expression networks using 3 head and neck and 5 breast cancer studies	51 common hypoxia metagene	Prognostic in breast, 2 lung, head and neck datasets	Buffa et al. (2010)
FaDu, UTSCC5, UTSCC 14, UTSCC15 cell lines under hypoxia and either normal or acidic pH	27	No	Sorensen et al. (2010)
UTSCC33, FaDu and SiHatumourxenografts— hypoxic regions identified using FAZA hypoxia tracer; 58 HNSCC fresh frozen biopsies with oxygen electrode measurements	15	FFPE samples from DAHANCA5 trial— independent prognostic factor; more hypoxic tumours treated with nimorazole had reduced incidence of locoregional failure	Toustrup et al. (2011)
DU145, HT29, MCF7 cell lines exposed to normoxia and different rimes of severe hypoxia	7 temporal and 2 general signatures	Not prognostic in breast cancer dataset	Starmans et al. (2012)

 Table 3.4
 Summary of hypoxia gene expression signatures

were consistently co-expressed with previously validated hypoxia-regulated genes (Buffa et al. 2010). By applying more training sets and co-expression networks a reduced metagene had prognostic significance in 4 independent datasets of breast, lung and head and neck cancers, and outperformed larger published signatures. The top 26 genes from this signature were retrospectively assessed in FFPE samples from laryngeal and bladder cancer patients treated with ARCON or CON (carbogen and nicotinamide) respectively (Eustace et al. 2013). Tumours were categorised into high and low hypoxia groups, and laryngeal tumours in the high hypoxia group showed greater benefit from ARCON than the low hypoxia group. The hypoxia signature did not predict benefit from CON in bladder cancer.

In contrast to previous studies on hypoxia gene expression, Toustrup et al. (2011) developed a classifier based on the hypoxic status of tumours. The hypoxia induced pH independent gene profile developed by Sorensen et al. was validated in vivo in a xenograft study using ¹⁸F-FAZA as an exogenous hypoxia radiotracer in autoradiographic studies. Hypoxic, non-hypoxic and mixed heterogeneous tumour areas defined by ⁸F-FAZA positive and negative regions were demarcated and dissected. All the genes investigated were significantly upregulated in hypoxic tumour areas compared with non-hypoxic areas, and all but 3 were upregulated in samples from mixed heterogenous versus non-hypoxic areas. To identify the most informative genes, a training set of 58 hypoxia-evaluated HNSCC FFPE biopsies were analysed for gene expression. The oxygenation status of these tumours had previously been evaluated in accordance to the relative number of oxygen electrode measurements less than 2.5 mmHg in their metastatic cervical lymph nodes. A 15-gene expression classifier was generated containing 15 of the in vitro identified hypoxia-responsive genes, which could best discriminate between 'more' and 'less' hypoxic human HNSCCs. This was evaluated in an independent data set, where patients with HNSCC were randomised to receive either hypoxic modification with nimorazole or placebo concomitant with radiotherapy. Patients with 'more' hypoxic tumours defined by the classifier had significantly higher cumulative incidence of locoregional failure at 5 years compared with those with less hypoxic tumours. Within the 'more' hypoxic group, patients treated with nimorazole had significantly reduced incidence of failure compared with placebo, whereas in the 'less' hypoxic group there was no significant difference in outcome, suggesting the classifier potentially has predictive value. Further subgroup analysis demonstrated that this benefit was only found in HPV-negative 'more' hypoxic tumours and not HPV-positive, whose outcome was unaffected by hypoxic modification (Toustrup et al. 2012). The same frequencies of 'more' and 'less' hypoxic tumours were in HPV-negative and positive groups as assessed by the classifier. More recently the classifier was used to assess 55 patients recruited in a randomised trial accelerated RT with or without nimorazole (Hassan Metwally et al. 2015). RT alone in 16 patients with 'more' hypoxic tumours was associated with higher locoregionaltumour failure.

More recently further investigation into the time dependent gene expression changes in response to hypoxia was carried out using prostate, colon and breast cancer cell lines exposed to hypoxia at 8 different time points between 0 and 24 h (Starmans et al. 2012). Seven different signatures consisting of induced genes with distinct temporal profiles and 2 general hypoxia signatures were generated, but none were prognostic in a large breast cancer cohort. In contrast previously published in vivo derived signatures showed clear prognostic power, suggesting the importance of the tumour microenvironment in the response to hypoxia.

Classification of hypoxia gene expression has the potential to represent prognostic and predictive markers in cancer and has been demonstrated retrospectively to be applicable to unselected FFPE biopsy samples, and therefore translatable into clinical practice. However, the 2 smaller gene signatures (15 and 26 gene signatures) which have demonstrated predictive value, have only 4 genes that overlap, suggesting that this method alone may not identify all hypoxic tumours. Prospective validation in clinical studies is required before further conclusions can be made.

3.6.4 Magnetic Resonance Imaging (MRI)

3.6.4.1 Dynamic Contrast-Enhanced MRI (DCE MRI)

In this technique fast repeated images are acquired before, during and after the rapid administration of a small hydrophilic gadolinium-based contrast agent, which diffuses through blood vessel walls and distributes into the extracellular space. The change in signal intensity reflects tumour perfusion, vessel permeability and the volume of extracellular space (Turkbey et al. 2010). It does not correlate directly with tissue pO_2 levels but a small study in HNSCC found positive correlation between imaging parameters of poor perfusion and pimonidazole staining (Newbold et al. 2009), and with hypoxia defined by ¹⁸F-fluoromisonidazole PET uptake in nodal metastases (Jansen et al. 2010).

3.6.4.2 Blood Oxygen Level-Dependent MRI (BOLD MRI)

This is an indirect measure of visualising pO_2 in blood vessels and surrounding tissue based on the paramagnetic properties of deoxyhaemoglobin, which is related to tissue oxygenation. It does not provide quantitative information on oxygen concentration and the signal can be influenced by many factors such as blood flow and pH. However, studies have shown correlation between this technique and polargraphic electrode pO_2 measurements and pimonidazole staining in prostate cancer, suggesting BOLD MRI may provide complementary information related to tissue oxygenation (Chopra et al. 2009).

3.6.5 Positron Emission Tomography (PET)

PET is a molecular imaging technique that can visualise and quantify tumours and their microenvironment. Different biological tumour characteristics can be imaged depending on the radiotracer used. Several hypoxia-specific PET radiotracers have been developed and extensively investigated as it is a potentially useful non-invasive technique for identification, quantification of hypoxia and repeated measurements after intervention. The metabolic activity demonstrated by the commonly used radiotracer ¹⁸Fluorine-fluororodexoxyglucose (¹⁸F-FDG) is indirectly related to the proliferative activity and oxygenation status of tumours, but cannot reliably distinguish hypoxic tumours. A good hypoxia tracer should be able to detect pO_2 levels that are clinically relevant to therapy, and be able to distinguish between normoxia, hypoxia, anoxia and necrosis. The molecule should be small and lipophilic making it highly membrane permeable leading to rapid uptake into cells and rapid clearance from normoxic cells to allow a high tumour to background contrast. It should also have a good dosimetry profile with simple radiolabelling and production, and low radiation dose to the patient. Two main classes of hypoxia imaging radiotracers are in available in the clinical trial setting: one based on the nitroimidazole compounds and one based on a complex of copper with diacetyl-bis (N⁴-methylsemicarbazone) (ATSM) ligands. A meta-analysis of published hypoxia imaging studies, which included PET and other imaging modalities of hypoxia, showed a uniform tendency for poor response to radiotherapy in tumours showing a higher uptake despite the heterogeneity of the image acquisition, data analysis and treatments (Horsman et al. 2012).

3.6.5.1 F-Fluoromisonidazole (¹⁸F-FMISO)

¹⁸F-FMISO is a derivative of the nitroimidazole group of compounds and is the most widely studied hypoxic radiotracer. It is moderately lipophilic and enters cells by passive diffusion across the cell membrane. The nitro group of the imidazole ring structure (R-NO₂) is reduced by intracellular reductases to R-NO₂⁻, which is reoxidised in the presence of oxygen and the tracer can flow back into the extracellular compartment. In conditions of hypoxia R-NO₂⁻ can be further reduced with progressive production of R-NH₂ compounds that bind covalently to intracellular molecules and the tracer becomes trapped. This occurs at pO₂ < 10 mmHg and therefore ¹⁸F-FMISO detects clinically relevant hypoxia. The process requires viable cells with functional nitroreductases and the tracer does not accumulate in necrotic cells (Prekeges et al. 1991).

3.6.5.2 Clinical Studies of ¹⁸F-FMISO in HNSCC

Rajendran et al. (2006) reported the largest study ¹⁸F-FMISO, which involved 73 patients with HNSCC. Uptake was seen in 85% of patients and tumour to blood ratio and hypoxic volume showed a trend to be an independent prognostic measure. Rischin et al. (2006) demonstrated that pretreatment ¹⁸F-FMISO was effective in determining hypoxic regions and tirapazamine was effective in patients with hypoxic tumours as assessed by ¹⁸F-FMISO. Another group found that an standardised uptake value (SUV) > 2 and tumour to muscle ratio (TMR) > 1.6 at 4 h after injection were associated with disease recurrence after RT (Eschmann et al. 2005). The same group also investigated the changes in ¹⁸F-FMISO uptake during RT by performing a scan pretreatment and repeating at 30 Gy. The mean SUV and TMR significantly decreased during RT, indicating RT-induced reoxygenation. Lee et al. (2009) also performed a baseline and midtreatment scan and found 90% of 20 patients had hypoxia before treatment and only 2 had detectable hypoxia on their midtreatment scan, which did not correlate with outcome. Zips et al. (2012) carried out a baseline scan and 3 further scans during RT. The imaging parameters from scans at week 1 and 2 of RT strongly correlated with progression free survival, suggesting the prognostic value of imaging hypoxia at the start of RT rather than at baseline. Smaller studies have showed variable results, with ¹⁸F-FMISO showing borderline significance for stratifying patients into treatment outcome groups in 12 patients, and high SUV_{max} but not TMR correlating with poor DSS after RT or surgery in 17 patients (Kikuchi et al. 2011).

¹⁸F-FMISO is not a universal hypoxia tracer in cancer. It has shown minimal activity in pancreatic cancer patients (Segard et al. 2013) and high background uptake in the normal rectum in patients with colorectal cancer, as well as tracer diffusion through the bowel wall, making images difficult to interpret (Roels et al. 2008). The low uptake of ¹⁸F-FMISO in target tissue and slow clearance of unbound ¹⁸F-FMISO from normoxic areas results in images of poor contrast. The relatively short half-life of 110 min hampers late imaging that could enhance contrast between hypoxic and normoxic tissues. This has led to the development of other tracers with improved pharmacokinetics.

3.6.5.3 ¹⁸F-Fluoroazomycin Arabinofuranoside (¹⁸F-FAZA)

This second generation 2-nitroimidazole is more hydrophilic than ¹⁸F-FMISO and therefore has faster clearance from normal tissues and higher tumour to background ratios. Souvatzoglou et al. (2007) carried out an 11 patient pilot study in HNSCC and reported a higher contrast with non-target tissues compared to ¹⁸F-FMISO, with an average TMR of 2.0 at 2 h post injection. Mortensen et al. (2012) performed static ¹⁸F-FAZA imaging in 40 oropharynx SCC patients. A hypoxic volume could be identified in 25 patients, with a median TMR of 1.5. The distribution of hypoxia among HPV positive and negative tumours was not significantly different and there was a significant difference in DFS in patients with non-hypoxic tumours compared

with hypoxic tumours. Tumours were ranked according to the expression of genes included in a 15-gene hypoxia classifier but no correlation was found between the hypoxic status as assessed by the classifier and ¹⁸F-FAZA imaging, despite the gene profile being validated using ¹⁸F-FAZA autoradiography in human tumourxenografts (Toustrup et al. 2011).

3.6.5.4 Copper(II)-Diacetyl-Bis (N⁴-Methylthiosemicarbazone) (Cu-ATSM)

Cu-ATSM was developed to overcome some of the limitations of ¹⁸F-FMISO imaging. It is an uncharged lipophilic molecule of low molecular weight and high membrane permeability. It has fast tumour uptake and clearance form normoxic tissues, allowing rapid imaging after injection and higher hypoxic-to-normoxic contrast. It is also simpler to synthesise and radiolabel compared with ¹⁸F-FMISO, but the longer half-life results in a higher radiation dose to patients (Lewis et al. 2001).

The hypoxia selectivity of Cu-ATSM is thought to be due to the bioreduction of Cu(II)-ATSM to Cu(I)-ATSM, which occurs in both normoxic and hypoxic cells, generating unstable anionic Cu(I)-ATSM. Cu(I) slowly dissociates from the ATSM ligand in cells with low oxygen concentration, becoming irreversibly trapped within the cell due to its negative charge. In the presence of oxygen Cu(I)-ATSM oxidised back to neutral Cu(II)-ATSM and diffuses out of the cell. The hypoxia selectivity therefore is a competition between dissociation and reoxidation of the reduced copper complexes, which is dependent on the oxygen concentration (Dearling et al. 2002). The precise cellular retention mechanism of Cu-ATSM is still unclear and uptake and retention has been associated with a number of other factors, such as over-reduced states caused by mitochondrial dysfunction (Yoshii et al. 2012), the expression of multidrug resistance protein (Liu et al. 2009) and Cu metabolism itself (Hueting et al. 2014). In addition preclinical studies have suggested hypoxia specificity may be dependent on tumour type (O'Donoghue et al. 2005; Yuan et al. 2006) and therefore concerns have been raised regarding the use of Cu-ATSM as a hypoxia radiotracer. However, small clinical studies in a number of tumour types have demonstrated correlation between uptake and poor prognosis, warranting further investigation.

3.6.5.5 Clinical Studies of Cu-ATSM in HNSCC

In 2001 Chao et al. (2001) used ⁶⁰Cu-ATSM in a radiotherapy planning study to demonstrate the feasibility of Cu-ATSM-guided IMRT to overcome hypoxia tumour resistance. Minagawa et al. (2011) assessed tumour hypoxia using ⁶²Cu-ATSM as a predictor of response in 17 patients. After 2 years of follow up SUV_{max} but not TMR was significantly different in patients with or without residual/recurrent tumour, and a SUV_{max} > 5 was a possible cut off threshold for

poor outcome. There was no significant difference in ¹⁸F-FDG between patients with and without residual/recurrence tumour. Sato et al. (2014) also evaluated the prognostic potential of ⁶²Cu-ATSM in 25 patients with head and neck cancer. ⁶²Cu-ATSM SUV_{max} > 3.6 was associated significantly worse progression free survival (PFS), and a TMR > 3.2 was associated with significantly worse progression free survival and cause specific survival (CSS). ¹⁸F-FDG uptake parameters did not distinguish patient with good and poor outcome. ⁶⁴Cu-ATSM in head and neck cancers was used to assess the efficacy of pretherapy ⁶⁴Cu-ATSM scanning as a prognostic factor of response to therapy in 11 patients (Grassi et al. 2014). Patients received a ⁶⁴Cu-ATSM and ¹⁸F-FDG PET scan before commencing treatment and response was determined by an ¹⁸F-FDG PET 3 months after therapy. The sensitivity and specificity of ⁶⁴Cu-ATSM SUV_{max} in predicting complete response to therapy were 100 and 50% respectively.

3.7 Conclusions

Tumour hypoxia remains one of the major causes of treatment failure in head and neck cancer. Hypoxia promotes a more aggressive tumour phenotype and is considered an important therapeutic target. Current methods of hypoxia identification are inefficient and therefore it is difficult to accurately assess hypoxia in tumours; accurate detection of hypoxia is important to allow stratification of patients for tailored treatment, and to reduce unnecessary side effects of intensive treatment modalities for less hypoxic tumours. In recent years, research in this field has substantially increased our understanding of the molecular pathways that contribute to tumour hypoxia and key pathways such as HIF, mTOR, UPR, and many more upstream and downstream biological targets of tumour hypoxia have been identified. Several of hypoxia-regulated genes are under investigation for drug development. In addition, a number of mRNA and microRNA signatures of tumour hypoxia have been published. So far, the clinical applications of such biomarkers have been limited due to inconsistencies and heterogeneity between studies. Approaches exploiting tumour hypoxia both for development of anti-cancer drugs that are activated in a hypoxic environment and therapeutic approaches that more directly target tumour hypoxia are under development.

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