Review

Inflammation and breast cancer

Microenvironmental factors regulating macrophage function in breast tumours: hypoxia and angiopoietin-2

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Abstract

Considerable evidence has now accumulated for tumour-associated macrophages stimulating key aspects of tumour progression, including the proliferation, survival and metastasis of tumour cells, tumour angiogenesis and suppression of the anti-tumour functions of other immune effectors at the tumour site. Tumour microenvironmental factors such as hypoxia have profound, direct effects on these cells, stimulating many of their pro-tumour functions. Hypoxia also does so indirectly by stimulating the release of the cytokine angiopoietin-2 from tumour cells and tumour blood vessels. This in turn then recruits Tie-2-expressing monocytes into tumours from the bloodstream and inhibits their production of anti-apoptotic and anti-angiogenic cytokines.

Role of tumour-associated macrophages in tumour progression

Two different approaches – the use of various transgenic mouse models and the analysis of human tumours – have demonstrated a close link between the activity of tumour-associated macrophages (TAMs) and tumour progression [1,2]. TAMs are abundant in most forms of solid tumour, where they often display a relatively immature phenotype and are positively correlated with tumour angiogenesis and/or progression (reviewed in [3]). Pollard's group crossed PyMT-MMTV mice (which spontaneously develop mammary tumours) with the transgenic op/op mouse model lacking the gene for colony-stimulating factor-1, a crucial growth factor for macrophages and their precursors from the bone marrow, namely blood monocytes. The tumours that developed in these macrophage-depleted mice showed a slower rate of progression to malignancy and formed far fewer metastases in the lungs than those in non-macrophage-depleted mice [1]. Moreover, Pollard's group recently characterised the development of the vasculature in PyMT-MMTV tumours during progression to malignancy and showed that the onset of the 'angiogenic switch' (the formation of the high-density vessel network associated with the transition to malignancy) was regulated by TAMs. Preinvasive mammary lesions in op/op mice exhibited both a delayed angiogenic switch and transition to malignancy, whereas genetic restoration of the macrophage population in tumours reversed this [4]. Although these studies suggest that TAMs have a key role in promoting tumour angiogenesis, progression to malignancy and metastasis, they have yet to be confirmed in similar studies with other macrophage-depleted, transgenic mouse tumour models.

However, these data accord well with our finding that high numbers of TAMs correlate with increased tumour angiogenesis, lymph node status and reduced survival of breast cancer patients [5]. Moreover, we showed that TAMs in breast carcinomas express numerous tumour-promoting factors such as the important mitogen epidermal growth factor [6] and the pro-angiogenic cytokine vascular endothelial growth factor (VEGF) [7]. TAMs have also been shown to release a variety of other cytokines and enzymes known to promote tumour invasion, angiogenesis and metastasis [3,8]. Recent studies indicate that when macrophages migrate into tumours they downregulate their expression of the potent anti-angiogenic cytokine IL-12 [9].

These findings have prompted investigations into how the tumour microenvironment 'educates' macrophages to perform these pro-tumour activities. Here we outline the important role of tumour hypoxia in this, both in the form of a direct effect on the expression of pro-tumour genes by TAMs, and indirectly by upregulating the pro-angiogenic cytokine angiopoietin-2 (Ang-2), which in turn has profound effects on TAM function.

Ang = angiopoietin; IL = interleukin; TAM = tumour-associated macrophage; TNF = tumour necrosis factor; VEGF = vascular endothelial growth factor.

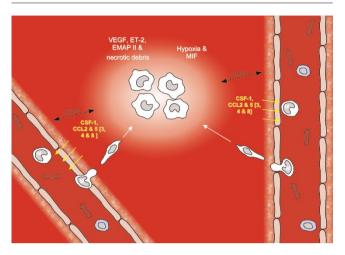## Effect of tumour hypoxia on tumour-associated macrophages

The vasculature in tumours is often disorganised, chaotic and prone to collapse. This results in the formation of multiple areas of inadequate vascular perfusion and hypoxia in solid tumours [10]. For some time it has been known that hypoxia alters the function of tumour cells, stimulating them to release pro-angiogenic factors, de-differentiate, become resistant to most forms of chemotherapy and metastasise [11]. We and others have shown that hypoxia also has marked effects on macrophage function in tumours.

A subpopulation of TAMs gather in hypoxic, avascular and/or necrotic sites in breast tumours [5,7,12], possibly as a result of the release of such macrophage chemoattractants as endothelial-monocyte-activating polypeptide (EMAP) II, endothelin-2 and VEGF by tumour cells in these sites (reviewed in [13]). Furthermore, because macrophages are phagocytes they may also be attracted into hypoxic, peri-necrotic areas along a trail of necrotic debris emanating from these areas. Indeed, we have shown recently that necrotic debris generated from the repeated freezing and thawing of a human breast tumour cell line (T47D) acts as a powerful chemoattractant for human macrophages *in vitro* and that this was mediated in part by their detection of necrotic debris by cell surface receptors called class A scavenger receptors (R Hughes, C Murdoch, S Tazzyman and CE Lewis, unpublished observations). Once they reach a hypoxic area, it seems from the work of Balkwill and colleagues [14] that hypoxia then inhibits macrophage migration, immobilising them in these areas. The interplay of these microenvironmental cues on TAM migration is illustrated in Figure 1.

Exposure to hypoxia in these sites stimulates TAMs to acquire a pro-angiogenic phenotype. For example, it stimulates them to express VEGF [7] and the pro-invasive and pro-angiogenic enzyme matrix metalloproteinase-7 [15], as demonstrated *in vitro* and in hypoxic areas of breast tumours. Furthermore, we recently demonstrated that when human macrophages infiltrate into the hypoxic centre of human breast tumour spheroids *in vitro*, they release VEGF and significantly enhance the angiogenic potential of spheroids when they are subsequently implanted into dorsal skin window chambers in nude mice and observed after 3 days [16]. This accords with the finding that hypoxia stimulates primary human macrophages to upregulate more than 30 other pro-angiogenic genes [17]. The impressive array of pro-tumour cytokines, enzymes and cell surface receptors expressed by macrophages in hypoxia are summarised in Figure 2 and reviewed in [18].

Hypoxia-induced gene expression in tumour cells is known to involve the nuclear accumulation of the transcription factors hypoxia-inducible transcription factor (HIF)-1 and HIF-2, which bind to cognate binding sequences in or near the promoters of target genes. Macrophages upregulate both
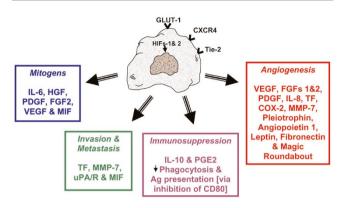
Figure 1

Mechanisms responsible for the accumulation of tumour-associated macrophages in hypoxic areas of solid tumours. A microenronment within a human tumour is shown containing two blood vessels and a hypoxic area (white) that has formed because it is more than 100 to 150 µm from either vessel – the critical distance for oxygenation in such tissues. Monocytes (M) pass through these vessels and are recruited into tumours by the release of the monocyte chemoattractants CCL2, 3, 4, 5 and 8 and colony-stimulating factor-1 (CSF-1) by the tumour. Once monocytes have moved across the tumour vasculature, many are attracted into hypoxic areas by the hypoxia-induced release of other monocyte attractants such as vascular endothelial growth factor (VEGF), endothelin-2 (ET-2) and endothelial-monocyte-activating polypeptide (EMAP) II. These innate cells may also be attracted into sites experiencing chronic hypoxia (and thus cell death) along a trail of necrotic debris emanating from these areas. Hypoxia then acts directly on macrophages to immobilise them and also via the upregulation of macrophage migration inhibitory factor (MIF) by tumour cells, which has a similar effect on macrophage migration (reviewed in [13]).

HIFs when exposed to hypoxia *in vitro* or inside human tumours [15,19]. Furthermore, high expression of one of these (HIF-2) in TAMs in breast carcinomas was correlated with increased tumour vascularity, presumably because of the upregulation by these cells of HIF target genes such as that encoding the pro-angiogenic cytokine VEGF [20]. However, the effects of hypoxia on TAMs are linked to the differentiation status of macrophages: our studies indicate that, unlike fully differentiated macrophages, their precursors, human monocytes from peripheral blood, fail to upregulate either HIF in tumour levels of hypoxia. Rather, they upregulate other hypoxia-induced transcription factors such as Ets-1, ATF-4 (activating transcription factor-4), Egr-1 (early growth response-1), C/EBPβ (CCAAT-enhancer-binding protein β) and nuclear factor (NF)-κB [21]. This finding was confirmed by a recent report showing that a human monocytic cell line required exposure to a chemical inducer of differentiation before it could upregulate HIF-1 in response to hypoxia [22]. Because several recent studies have shown that TAMs exhibit a relatively immature phenotype in tumours [23], the exact contribution of these various transcription factors to the

Figure 2

Hypoxia induces marked changes in the phenotype of macrophages. Macrophages upregulate hypoxia-inducible transcription factor (HIF)-1 and HIF-2 in hypoxia, which translocate to the nucleus to induce the expression of a wide array of target genes. Several important cell-surface receptors are upregulated in hypoxia, including the glucose receptor GLUT-1 (for increased glucose uptake as the cell switches to anaerobic glycolysis to make ATP in the absence of oxygen), the chemokine stromal cell-derived factor-1 (SDF-1) receptor CXCR4, and the angiopoietin receptor Tie-2. Hypoxia also stimulates the expression of a wide array of other pro-tumour cytokines, enzymes and receptors, grouped here according to their known function in tumours. Downregulation of a factor or tumour-associated macrophage function is indicated by an arrow [15,17,18]. Ag, antigen; COX, cyclo-oxygenase; FGF, fibroblast growth factor; HGF, hepatocyte growth factor; MIF, macrophage migration inhibitory factor; MMP, matrix metalloproteinase; PDGF, platelet-derived growth factor; PGE2, prostaglandin E_2; TF, tissue factor; uPA/R, urokinase plasminogen activator receptor; VEGF, vascular endothelial growth factor.

pro-angiogenic phenotype of hypoxic TAMs now merits further investigation.

Angiopietin-2 and tumour-associated macrophage functions

The cytokines Ang-1 and Ang-2 regulate processes such as angiogenesis by binding specifically to the receptor tyrosine kinase Tie2/Tek on endothelial cells. Ang-1 acts as a Tie-2 agonist to promote and stabilise mature vessels by promoting interactions between endothelial cells, pericytes, basement membrane and surrounding extracellular matrix. Conversely, Ang-2, the predominant form of angiopoietin in malignant tissues, has been shown to be a functional antagonist of Ang-1 and binds competitively to Tie-2, antagonising the stabilising effect of Ang-1, resulting in an overall destabilisation of existing vessels (reviewed in [24]). In the presence of VEGF these destabilised vessels undergo angiogenic changes and sprout to form new vessels. Thus, angiogenesis is controlled by a dynamic balance between vessel stabilisation and growth, mediated by VEGF, Ang-1 and Ang-2. It has been shown that breast carcinomas express higher levels of Ang-2 than of Ang-1 and that this is correlated with high levels of VEGF expression and tumour angiogenesis [25].

However, Ang-2 has recently been shown to have agonistic functions and to be capable of activating Tie-2 to stimulate endothelial cell migration and tubule formation *in vitro* [26]. Thus, Ang-2 seems to possess both agonist and antagonist functions when acting on endothelial cells.

Until recently, expression of Tie-2 was thought to be restricted to endothelial cells, but in 2005 De Palma and colleagues showed that a subpopulation of murine blood monocytes expressing Tie-2 are recruited into spontaneous murine and orthotopic human xenograft tumours and have a crucial role in stimulating tumour angiogenesis [27]. We and De Palma's group have now extended these studies to show that Tie-2+ monocytes are also abundant in human peripheral blood and exist in a range of human tumours [28,29]. Moreover, Ang-2 was seen in both studies to act as a powerful chemoattractant for these Tie-2+ monocytes *in vitro* and is therefore highly likely to recruit Tie-2+ monocytes from the bloodstream into tumours [28,29].

Our studies also demonstrated that hypoxia stimulates Tie-2 expression by human monocytes and macrophages [28], suggesting that hypoxia may modulate the response of these cells to Ang-2. Interestingly, several recent studies have shown Ang-2 to be upregulated by tumour cells in hypoxic areas of human tumours [30]. This means that it is highly likely that TAMs would be exposed to both hypoxia and Ang-2 in such areas. This is important because we found that exposure to hypoxia and Ang-2 had marked inhibitory effects on the release of IL-12 by human Tie-2+ monocytes. This suggests that when monocytes are recruited into tumours and exposed to both Ang-2 and hypoxia it inhibits their ability to mount an anti-angiogenic response. This, together with their hypoxia-induced pro-angiogenic functions, would ensure rapid angiogenesis in (and hence re-oxygenation of) the avascular, hypoxic site.

Moreover, the combined action of Ang-2 and hypoxia also inhibited the release of TNF-α by such cells [29]. This is important because high-dose TNF-α is known to promote the apoptosis of both tumour and endothelial cells [31], so its downregulation near newly formed angiogenic blood vessels could enhance tumour and endothelial cell survival and thus promote metastasis and angiogenesis, respectively. Further-more, high concentrations of TNF-α inhibit Ang-2 synthesis by endothelial cells [32], so our data suggest that this response of TAMs to Ang-2 may contribute to the high levels of Ang-2 reported in breast tumours [25].

Conclusion

There is now undeniable evidence that macrophages drive tumour angiogenesis and progression in certain murine mammary tumour models, and this correlates well with studies of their function in human breast tumours. Many TAMs are found in hypoxic areas of such tissues where unequivocal evidence has now been provided to show that

hypoxia has profound effects on their function, stimulating them to produce a wide array of pro-tumour growth factors, cytokines and enzymes. Furthermore, hypoxia-induced cytokines such as Ang-2 produced with the tumour microenvironment seem to amplify the effects of hypoxia on TAMs. It is hoped that present attempts to unearth the signalling pathways mediating the powerful effects of hypoxia and Ang-2 on these cells will highlight new targets for new anti-cancer strategies.

Competing interests
The authors declare that they have no competing interests.

Acknowledgements

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