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Tumor angiogenesis and lymphangiogenesis: tumor/endothelial crosstalk and cellular/microenvironmental signaling mechanisms

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Abstract

Tumor angiogenesis and lymphangiogenesis are key features of tumor progression and metastasis. The role of tumor cells-derived factors in the promotion of associated angiogenesis and lymphangiogenesis is much studied and, no doubt, very important for the understanding of cancer progression. This review aims to present and discuss the work done on the pro-angiogenic and lymphangiogenic cellular interactions within the tumor microenvironment and the signaling pathways that regulate this cross talk. Such multifactor studies are critical for the development of future therapeutic approaches for cancer because they take into account the complexities of cellular interactions within the tumor microenvironment.

Keywords

Angiogenesis; angiogenesis stimulators; lymphangiogenesis; lymphangiogenesis stimulators; tumor microenvironment; endothelial cells; tumor cells; extracellular matrix

Introduction

It is well known that tumors require a microvasculature development in order to grow and metastasize (Hanahan and Folkman, 1996). The cancer cells migration to distant tissues occurs through blood and lymphatic vessels(He et al., 2005). The main metastatic route for oral squamous cell carcinoma is the lymphatic one (Tobler and Detmar, 2006). Blood and lymphatic vessels in tumor tissue are major components of the tumor microenvironment. These vessels are newly formed from pre-existing host vessels stimulated by pro-blood-angiogenic and pro-lymph-angiogenic (pro-blood/lymph-angiogenic) factors expressed in tumor cells (Onimaru et al., 2011).

Conflict of interest

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Angiogenesis plays a important role not only in the tumor growth and its blood supply, but also in the tumoral metatstasis. The angiogenic processes are mediated mostly by cellular stimulation of diverse growth factors and secreted molecules. The main elucidated signaling mechanism involves the vascular endothelial growth factor-A (VEGF-A), produced by tumor cells, and the consequent proliferation activity of vascular endothelial cells (Tammela et al., 2005). Although these processes seem simple, in fact there are many complex mechanisms involving tumor cells, endothelial cells, as well as other stromal cells and microenviromental molecules working together in several signaling pathways.

In morphological terms, angiogenesis can present itself as a sprouting (Eilken and Adams, 2010), intussusseptive event (Burri et al., 2004), or even the tumor cells or other types of cells can turn into "endothelial-like" cells (Gottfried et al., 2007; Maniotis et al., 1999). In addition, both tumor and endothelial cell scan present different phenotypes in particular organs, tumor types and subtypes (Jung et al., 2002).

The lymphatic endotelial cells are distinct from vascular endotelial cells regarding their biological, functional and structural aspects, and also in terms of their protein secretion and expression. These cells can play an important role in the pathogenesis of cancer, changing the tumoral microinvironment.

Moreover, with the advances in research it becomes more clear the complexity of growth factors and receptors involved in angiogenic and lymphangiogenic processes. Several studies have shown that the growth factors which are more implied in lymphangiogenesis, such as vascular endothelial growth factor-C and -D(VEGF-C and VEGF-D), and its correspondent receptor vascular endothelial growth factor receptor-3(VEGFR-3), can also participate in angiogenesis. On the other hand, pro-angiogenic factors (ex: VEGF-A and angiopoetin-2) can also participate in lymphangiogenesis (Scavelli et al., 2004). Furthermore, recently there were discovered crosstalk mechanisms between tumor and endothelial cells (Issa et al., 2009; Kaneko et al., 2007; Neiva et al., 2009), which will be further elucidated in this review. With the research involving crosstalk mechanism, the literature started to point out to the existence of different intercellular crosstalk mechanisms involving tumor angiogenesis and lymphangiogenesis. Vascular endothelial cells, in a particular tumor microenvironment, can interact not only with tumor cells but also with immune cells, fibroblasts, pericytes and the extracellular matrix (Jung et al., 2002).

Furthermore, it has been shown that tumor and stromal cells can be responsible for compensatory pro-angiogenic signaling against anti-angiogenic therapies, and that these treatment approaches are still much directed against one or few pro-angiogenic factors (Abdollahi and Folkman, 2010). Specific anti-lymphangiogenic therapy in cancer is still not applied in clinical trials. However, several studies in animal models seem to be promising in inhibiting the lymphatic spread of cancer cells (Stacker and Achen, 2008). A recent review has focused on the cell-dependent molecular mechanisms that regulate tumor blood/lymph-angiogenesis, discussing the features tumor-associated blood/lymph-angiogenesis separately, and concerning the role of blood/lymph-angiogenic cascades in tumor microenvironment (Onimaru et al., 2011). In the present mini review we take in account this aspects and also the interactions among all the cells in the tumor microenvironment, pointing out the inter-cellular and matrix pro-angiogenic and lymphangiogenic mechanisms have been recognized as a result of interconnected factors, didactically separated into topics in this review.

Initial considerations

Hanahan and Weinberg published two very important articles entitled "The Hallmarks of Cancer" (Hanahan and Weinberg, 2000) and "Hallmarks of Cancer: the Next Generation"

(Hanahan and Weinberg, 2011). These works show that there are tumor checkpoints, and that they can occur in different sequences within particular types of cancer (Hanahan and Weinberg, 2000). Such hallmarks are now eight: resisting cell death, deregulating cellular energetics, sustaining proliferative signaling, evading growth supressors, avoiding immune destruction, enabling replicative immortality, activating invasion and metastasis and inducing angiogenesis; mediated by two enabling characteristics: tumor-promoting inflammation and genome instability and mutation (Hanahan and Weinberg, 2011). Not all cancers need to activate all of these hallmarks and microenvironmetal and/or genetic alterations, but no cancer can develop itself without any of them. Taking this discussion to the study of signalling mechanisms involved in angiogenesis and lymphangiogenesis, it can be considered some cellular and environmental checkpoints as well.

1. Signaling pathways in tumor angiogenesis and lymphangiogenesis

The most important molecular pathway involved in both tumor angiogenesis and lymphangiogenesis is the one based on VEGFRs and Neuropilins (NRPs) signal transduction. There are, to date, five growth factors in the VEGF family: VEGF-A, B, C, D and the Placenta growth factor (PIGF) (Koch and Welsh, 2012). Moreover, these growth factors can suffer alternative splicing (VEGF-A, B and PIGF) and further processing (VEGF-A, C and D). Which are associated with differences in binding to receptors (more or less affinity towards VEGFRs and/or NRPs) and to the very own extracellular matrix (Koch and Welsh, 2012; Ferrara, 2010).

As mentioned before, the angiogenic process is most regulated by VEGF-A, VEGFR-1 and VEGFR-2. While the best described lymphangiogenic regulators are VEGF-C and VEGFR-3 (Tammela et al., 2005). There are, however, overlaps between growth factors and receptors involved in both processes (Scavelli et al., 2004).

Another well established participant in angiogenesis is the Delta-like-4 (Dll4)-Notch signaling axis. Notch is the receptor for the ligand Dll4 (among others), involved in proliferating signals in different cellular types (Artavanis-Tsakonas and Lake, 1999). Papers show that, during angiogenesis, the branching of pre-existing vessels occurs concomitantly with variations in Dll4 and Notch expression in endothelial cells (Adams and Alitalo, 2007). Where the tip cells (the ones leading the branching process) have an up-regulated expression of Dll4, and the stalk cells (that form the tube following tip formation) have an up-regulated expression of Notch (Dufraine et al., 2008). It has been shown that tumor cells can have an increased expression of another Notch ligand, Jagged-1, associated with pro-angiogenic effects (Zeng et al., 2005). And that a soluble form of the Notch 1 receptor (Notch 1 decoy) could have anti-angiogenic properties, thus, being a promising therapeutic approach towards cancer (Funahashi et al., 2008).

2. The interactions between the extracellular matrix (ECM) and the vasculature (blood and lymphatic vessels)

2.1. Tumor angiogenesis and the extracellular matrix

Angiogenesis is regulated by the homeostatic balance between pro- and anti-angiogenic factors (Carmeliet and Jain, 2000). The "angiogenic switch" occurs when an "imbalance" between those factors is established, pending to the side of greater concentrations of positive angiogenic regulators, usually such imbalance occurs at the transition between potentially malignant lesions and the cancer (Hanahan and Folkman, 1996). The literature points to at least 27 endogenous inhibitors of angiogenesis, such as proteins and small molecules (Nyberg et al., 2005). The matrix-derived angiogenesis inhibitors seem to be arresten, canstatin, collagen fragments, endostatin-like fragment from type XV collagen (EFC-XV),

endorepellin, endostatin, fibronectin fragments, fibulin, thrombospondin-1 and -2 and tumstatin (Nyberg et al., 2005).

It has been shown that the cancer cells may escape from ECM endogenous inhibitors by upregulating pro-angiogenic factors (Fernando et al., 2008). *In vivo* studies of cancer cells over expressing thrombospondin-1, endostatin and tumstatin showed that all cancers presented an ability to escape from these inhibitory situations by production of pro-angiogenic molecules such as VEGF, platelet-derived growth factor-A and -B (PDGF-A and – B, fibroblast) growth factor (FGF) and angiopoetin-2 (Fernando et al., 2008).

The ECM can play not just an anti-angiogenic role, but also, a pro-angiogenic influence in tumorigenesis (Campbell et al., 2010; Han et al., 2008). There were discovered different pro-angiogenic functions played by already known matrix-derived factors such as collagen IV, laminin and fibronectin (Campbell et al., 2010). Han et al. (2008) showed that the extracellular matrix protein-1 (ECM1) had a direct correlation with tumor angiogenesis and metastatic potential in laryngeal carcinoma. The very own pro-angiogenic growth factors and cytokines such as VEGF, FGF-2, interleukin-8 (IL-8 or CXCL8), tumor growth factor-β (TGF-β), PDGF and angiopoietins are constitutively present in the ECM and may be mobilized from it to exert their functions in favor of the tumor (Crivellato et al., 2008). It is fundamental to keep in mind that tumorigenesis is a dynamic process involving a network of molecular events related to each other. Therefore, the ECM can play different roles depending on several microenvironmental factors of the tumor, such as many different cell types (epithelial cells, endothelial cells and stromal cells), its proteic and glycoproteic components. The natural tissue homeostasis regulators can, in fact, prevent the processes involved in tumor progression. However, when these physiological regulators reduce homeostatic effectiveness, the tumor microenvironment may start to act in an opposite manner, promoting tumor growth and development (Bissell and Hines, 2011).

2.2. Tumor Lymphangiogenesis and the extracellular matrix

The interactions between lymphatic endothelial cells (LECs) and the ECM, are quite similar to those observed between the vascular endothelial cells and ECM. Lymphangiogenic tissue homeostasis is achieved through balance of pro- and anti-lymphangiogenic factors and one of the endogenous lymphangiogenic inhibitors that is shared with vascular endothelial cells is endostatin (Ji, 2006). It has been shown that the use of angiogenic inhibitors such as endostatin, platelet factor-4 (PF-4) and interferon- α (INF- α) had significant inhibitory effects on proliferative and migratory activities of lymphatic endothelial cells (Shao et al., 2008; Shao and Xie, 2005). LECs present an incomplete or absent basement membrane, with the presence of anchoring filaments serving as binding structures to the surrounding tissues (Leak and Burke, 1968). There is still much research to be conducted in the field of the crosstalk among LECs, the surrounding ECM, and associated stromal or cancer cells. Focusing on ECM signaling, basically the most important molecules involved in lymphangiogenesis are hyaluronan and integrins (Ji, 2006; Paupert et al., 2011).

Hyaluronan is a mucopolysaccharide, abundantly present in the extracellular matrix, which provides a favorable microenvironment for cell proliferation and migration (Laurent and Fraser, 1992). Lymphatic vessel endothelial hyaluronan receptor-1 (YVE-1) is the receptor for hyaluronan constituvely present in lymphatic endothelium, which is homologue to CD44 (Banerji et al., 1999). Itano *et al.* (2008) showed that tumor-derived hyaluronan serves as a scaffold for interactions between cancer and host cells, favoring promotion of both tumor angiogenesis and lymphangiogenesis.

Other important molecules related to ECM-lymphatic endothelial interactions in lymphangiogenesis are the integrins, which are a large family of heterodimerictrans

membrane glycoproteins (Paupert et al., 2011). Mishima s. (2007) have shown that prospero homeobox protein-1(PROX-1), through induction of α 9 integrin expression on lymphatic endothelial cells, stimulated migration of LECs towards VEGF-C expressing cells. α 4 β 1 integrin interaction with matrix fibronectin may be involved in tumor associated lymphangiogenesis and metastasis (Garmy-Susini et al., 2010).

3. Crosstalk between tumor and endothelial cells

Tumor cells establish a specific stromal microenvironment fostering tumor growth, in which blood/lymph-angiogenesis are involved. A complicated cytokine networks induces the tumor-associated blood/lymph-angiogenesis continually, providing several interactions among tumor cells and stromal cells, in which endothelial cells (ECs) are included (Onimaru et al., 2011).

3.1. Crosstalk between tumor and vascular endothelial cells

It is well known that the tumor cells can secrete growth factors and stimulating molecules such as VEGFs, PDGF, FGF-2, epidermal growth factor (EGF), angiopoietins, etc, leading to a proliferative activity of vascular endothelial cells. Research has been performed recently in order to comprehend the crosstalk mechanisms between tumor and vascular endothelial cells that can occur, in the tumor microenvironment. Recently, Nör et al. (2001) using an in vivo model of human tumors surrounded by human microvessels demonstrated that tumors with vascular endothelial cells (HDMECs) stably expressing Bcl-2, had a 3-fold increase in growth as well as an increase in microvascular density and survival of endothelial cells compared with control HDMECs and tumor cells. Bcl-2 can stimulate, in an autocrine manner, proangiogenic activities in vascular endothelial cells mediated by nuclear factor kappaB (NF-kB) and further production of angiogenic CXC chemokines (Karl et al., 2005). Tumor cells can also present the ability to up-regulate the expression of Bcl-2, as an antiapoptotic gene (Hanahan and Weinberg, 2011). Moreover, the crosstalk between tumor and vascular endothelial cells is mediated by Bcl-2, as Kaneko et al. (2007) showed in their study. They demonstrated that vascular endothelial cells expressing Bcl-2, via the transcription factor signal transducer and activator of transcription-3 (STAT3), could secrete VEGF, which induces the up-regulation of Bcl-2 and angiogenic chemokines such as chemokine ligand 1(CXCL1)and CXCL8 in head and neck squamous cell carcinoma (HNSCC) cells, increasing survival and proliferative activities in both cell types (Kaneko et al., 2007), and also enhancing the invasiveness potential of tumor cells (Warner et al., 2008).

Another study from the same laboratory established that vascular endothelial cells could secrete CXCL8, interleukin-6 (IL-6) and EGF into the extracellular medium (Neiva et al., 2009). These authors determined that these molecules could induce intra-cellular fosforilation of STAT3, protein kinase B (PKB or Akt) and extracellular signal-(ERK) in HNSCC cells, leading to increased survival and migration potential in the tumor cells (Neiva et al., 2009).

Zeng *et al.* (2005) identified a different crosstalk mechanism involving Notch activation in endothelial cells via the Notch ligand Jagged-1 expression in HNSCC cells. The over expression of Jagged-1 was due to the activation of the mitogen-activated protein kinase(MAPK) signaling pathway in tumor cells after the addition of growth factors such as hepatocyte growth factor (HGF), tumor growth factor-a (TGF-a and EGF. This suggests a possible crosstalk between tumor and vascular endothelial cells mediated by such secreted molecules (Zeng et al., 2005). Liu *et al.* (2010) demonstrated that tyrosine kinase receptor tie-2/TEK positive glioma and brain tumor stem cells had an increased adhesion potential to endothelial cells, which was observed in *in vitro* and *in vivo* studies. Such receptor had already been identified in normal and tumor associated endothelial cells and it is, positively

related to an increase in tumor malignancy and angiogenesis (Peters et al., 1998). It has been shown, however, that this receptor is also present in brain tumor cells and that, stimulated by angiopoietin-1, tie-2 positive tumor cells increase the adhesion on vascular endothelial cells *in vitro* and *in vivo*, mediated by up-regulation of integrin β 1 and N-cadherin (Liu et al., 2010).

3.2. Crosstalk between tumor and lymphatic endothelial cells

It is known that tumor cells can produce and secrete VEGF-C and VEGF-D. VEGF-C is the most important growth factor in mediating tumor-associated lymphangiogenesis, which can be followed by migration of cancer cells through these newly formed lymphatic vessels, leading to the invasion and metastasis (Achen et al., 2005). As mentioned above, proangiogenic molecules can also participate in promoting tumor lymphangiogenesis (Scavelli et al., 2004). Issa et al. 2009 have brilliantly shown that tumor-secreted VEGF-C could not only induce, in an autocrine manner, an increased matrix-degradation and migration in a three dimensional model (mainly by consequent production of matrix metalloproteinase 9 [MMP-9]), but also, VEGF-C was capable of up-regulating chemokine ligand 21(CCL21) secretion in the lymphatic endothelium. This up regulation could also enhance the chemoatraction of C-C chemokine receptor type-7 (CCR7)-expressing tumoral cells, which could move toward the surrounding lymphatics (Issa et al., 2009). Moreover, Tarquinio et al. (2011) observed that VEGF-C was responsible for the up-regulation of Bcl-2 on lymphatic endothelial cells through VEGFR-3 and consequent PI3k/Aktphosphorylation. And that there's an increased expression of Bcl-2 on endothelial cells from human oral tumors with lymph node metastasis as compared to stage-matched oral squamous cell carcinomas without metastatic spread (Tarquinio et al., 2011).

4. Interactions between tumor angiogenesis and lymphangiogenesis with associated non-endothelial stromal cells

It is increasingly important to study how stromal cells can affect tumor angiogenesis and lymphangiogenesis. The literature highlights very interesting data concerning the proangiogenic and lymphangiogenic molecules secreted by such cells, some of it further described in this review. It is also important to acknowledge that, during the co-evolution between tumor cells and their microenvironment, those two interconnected mechanisms are frequently involved. Tumor cells can alter the surrounding stroma and stromal cells can strongly influence tumorigenesis and tumor progression (Polyak et al., 2009). Therefore, crosstalk between all cellular types such as dendritic cells, fibroblasts, macrophages, pericytes, lymphocytes and tumor cells is crucial for cancer establishment and development.

4.1. Role of non-endothelial stromal cells in tumor angiogenesis and lymphangiogenesis

4.1.1. Dendritic cells—Dendritic cells (DCs) are observed in peripheral tissues, being the sentinel of the innate immune system. Once in contact with an invading pathogen and being stimulated through inflammatory cytokines, those cells can ingest antigens, become mature, and after they circulate in the afferent lymphatics reaching the lymph nodes with consequent antigen presentation to lymphocytes, what characterizes DCs as a pivotal cell in the regulation of adaptive responses (Sozzani et al., 2007). Within the tumor microenvironment, DCs can produce different pro-angiogenic factors, such as tumor necrosis factor- α (TNF- α), TGF- β , and granulocyte macrophage colony-stimulating factor (GM-CSF), as well as they can response to different mediators related to angiogenesis (Sozzani et al., 2007). Tumor-secreted factors, suppressor myeloid cells, and tumor-associated macrophages have been demonstrated to inhibit DCs maturation in cancer, mainly by activation of the transcriptional factor STAT3, leading to a decreased effectiveness in immune responses, such as tolerance and immune suppression, facilitating the development of some malignancies (Fainaru et al.,

2010). Fainaru *et al.* (2010) have demonstrated *in vivo* that xenografted tumors with an increased level of angiogenesis were infiltrated by an immature population of DCs, whereas dormant a vascular tumors had an infiltration of mature DC population. Moreover, such as some tumoral cells, DCs can also "transform" into endothelial-like cells (Gottfried et al., 2007).

4.1.2. Fibroblasts—Cancer-associated fibroblasts (CAFs) have been shown to exert proangiogenic and developmental activities in melanoma (Li et al., 2003) and prostate cancer (Niu and Xia, 2009). Quiescent fibroblasts can be activated in a paracrine manner when present in the tumor microenvironment (Lorusso and Ruegg, 2008). Like some immune cells, stromal fibroblasts can also inhibit cancer in its initial stages (Rasanen and Vaheri, 2010).

However, tumor progression involves CAFs, which can promote tumor development (Orimo and Weinberg, 2006). In a 4T1 murine breast cancer model, CAFs where shown to promote tumor growth and metastasis by modulating the tumor immune microenvironment (Liao et al., 2009). Through DNA vaccine targeting the activated-CAFs proteins, these cells were eliminated resulting, in a shift of tumor immune polarization, followed by a suppressed recruitment of cancer associated macrophages and reduced tumor angiogenesis and lymphangiogenesis (Liao et al., 2009). Interestingly, fibroblasts may initiate their pro-angiogenic activity through VEGF secretion by tumor cells. The release of VEGF into the stromal microenvironment induces microvascular permeability, leading to extra vasation of plasma proteins such as fibrin, which in turn leads to a generated influx of fibroblasts, inflammatory cells and endothelial cells (Kalluri and Zeisberg, 2006). Fibroblasts can produce fibronectin and type I collagen rich extracellular matrices, VEGF and matrix metalloproteinasis (MMPs which are capable of conducting and initiating tumor angiogenesis and progression (Kalluri and Zeisberg, 2006).

In relation to lymphangiogenesis, Itano's research group has extensively worked with the pivotal role that ECM rich in hyaluronan may play in inducing tumor initiation and progression as well as intratumoral lymphangiogenesis (Itano et al., 2008; Koyama et al., 2008). Using an *in vivo* model of hyaluronan synthase2 (Has2) transgenic mice, it was observed that an overproduction of hyaluronan facilitated intratumoral lymphangiogenesis, suggesting that cancer cells could maintain favorable conditions for lymphangiogenesis through hyaluronan-induced CAFs recruitment with consequent VEGF-D secretion (Koyama et al., 2008).

4.1.3. Macrophages—These cells are phagocitary antigen-presenting cells, which originated from the bone-marrow, circulating briefly in the blood as monocytes and later differentiating into macrophages in the tissues or in inflammatory foci (Guruvayoorappan, 2008). Tumor-associated macrophages (TAMs) have been shown to affect not only tumor angiogenesis, but also tumor cell invasion onto surrounding stroma with degradation of the basement membrane by MMP production; immuno supression by expression of prostaglandin E2 (PGE2), interleukin-10 (IL-10) and TGF-β and metastasis by secretion of factors such as EGF that can guide tumor cells toward blood vessels (Lewis and Pollard, 2006). TAMs can secrete a number of potent pro-angiogenic growth factors and cytokines such as VEGF, TNF-α, bFGF and other angiogenesis modulating enzymes, including matrix metalloproteinase 2 (MMP-2), matrix metalloproteinase 7(MMP-7), matrix metalloproteinase 9(MMP-9), matrix metalloproteinase 12 (MMP-12) and cyclo-oxygenase-2 (COX-2) (Lewis and Pollard, 2006). Lamagna *et al.* (2006) stated that TAMs can have dual roles within the tumor microenvironment. They can act together with mature dendritic and NK cells, fighting against tumor cells (Lamagna et al., 2006). However, when

As reviewed in angiogenic processes linked to cancer, tumor-associated macrophages can promote tumor lymphangiogenesis as well (Schoppmann et al., 2002; Skobe et al., 2001). Schoppmann et al. (2002) have discovered populations of TAMs in samples of cervix squamous intraepithelial lesions, that expressed VEGF-C and D, as well as their receptor VEGFR-3. They have also isolated VEGFR-3 expressing monocytes in the circulation, suggesting that peritumoral lymphangiogenesis may occur, in part, due to the infiltration of such sub fraction of monocytes and the consequent differentiation into VEGF-C and D expressing tumor-associated macrophages (Schoppmann et al., 2002). Also, it has been demonstrated that VEGF-C could have a macrophage-recruitment activity, despite its lymphangiogenic and angiogenic activities (Skobe et al., 2001). Infiltration of TAMs has been correlated with increase of peritumoral lymphatics and poor prognosis in lung adenocarcinoma (Zhang et al., 2010). Also, in cutaneous squamous cell carcinoma, it was described that the microenvironment consisted of increased lymphatic density with high levels of macrophage-derived VEGF-C (Moussai et al., 2011).

4.1.4. Pericytes—Pericytes are cells that surround endothelial cells, in the formation of blood vessels. Reinmuth et al. (2001) have demonstrated that both pericytes and tumor cells could stimulate, in a paracrine manner, endothelial cells through VEGF secretion. Tumor and endothelial cells could produce PDGF towards pericytes, enabling them to secrete VEGF (Reinmuth et al., 2001).

4.1.5. T cells—Very interesting work has been done concerning VEGF interactions with T cells, although little is known about such associations. Shin et al. (2009) have demonstrated that T lymphocytes activated by monoclonal antibodies anti-CD3, anti-CD28 or by some antigens, transcribed mRNA for VEGFR-1 and 2, where only VEGFR-1 was present in the cell surface. Furthermore, the addition of VEGF to either non-activated or activated T lymphocytes led to up-regulation of IL-10 and slight down-regulation of interferon- γ (INF- γ) and anti-angiogenic cytokines. This suggest that high concentrations of VEGF may cause T cells to migrate towards a tumor microenvironment, and that this interaction could play a role in IL-10 mediated immune evasion by tumor cells (Shin et al., 2009).

4.1. 6. Mast cells—Mast cells (MCs) are immune cells that mediate the transition between innate and adaptive responses, orchestrating inflammatory responses against bacteria, parasites and viruses (Metz and Maurer, 2007). MCs present dual roles in tumorigenesis. They can display a suppressor activity by secreting different inhibitory interleukines and also the TNF- α or they can help tumor progression by promoting angiogenesis, ECM degradation and immuno suppression (Theoharides and Conti, 2004). Concerning the proangiogenic capacity, MCs can secrete molecules such as VEGF, FGF-2, IL-8 and TGF- β (Crivellato et al., 2008). The recruitment of these cells toward the tumor microenvironment is mainly due to the stem cell factor (SCF) produced by cancer cells (Crivellato et al., 2008).

MCs may play a very important role not just in angiogenesis, but also in tumor lymphangiogenesis. It was shown, *in vivo*, that endostatin was capable of decreasing the infiltration levels of VEGF-C expressing MCs in induced skin squamous cell carcinoma. The capacity of endostatin in inhibiting adhesion and migration of MCs on fibronectin has also been shown *in vitro*, suggesting that the reduction in aggressiveness caused by endostatin could be, in part, due to the decrease in levels of VEGF-C, which is consequence of MC migration and adhesion inhibition (Brideau et al., 2007).

5. Tumor-stroma interactions and therapeutic applications

The increasing knowledge of the signaling mechanisms between cancer cells and the surrounding stroma has opened room for therapeutic approaches targeting the tumor microenvironment. It is true, for example, that the studies about the VEGF signaling have allowed the development of the so called therapeutic agent Bevacizumab (a monoclonal antibody against VEGF-A). The effects of this drug will be also targeted to stroma-derived VEGF (Hofmeister et al., 2008). Still concerning anti-angiogenic therapy, the soluble Notch1 decoy (Funahashi et al., 2008) seems to be an elegant approach towards tumor/ endothelial crosstalk. The most studied cell component of the tumor microenvironment is the associated endothelial cell. In this respect, very interesting work have also targeted an endothelial specific membrane molecule, CD105 (endoglin). Moreover, in vivo studies are being carried out in order to suppress the anti-apoptotic effects of this surface protein using these antibodies (Hofmaister et al., 2008). Maybe the future in stroma-directed therapies is to develop new strategies targeting molecular surface markers of the cells composing the tumor microenvironment.

6. Concluding remarks

The purpose of this review was to present, in a dynamic topic format, a concise and updated review about the work that has been done concerning microenvironmental mechanisms and intercellular interactions in tumor angiogenesis and lymphangiogenesis. Such processes involving neo vasculature have been well described as important keys in tumor growth and metastasis and, for that reason, it is increasingly necessary to enlarge the knowledge of these interactions and signaling pathways, with the interest of pursuing new and more complete therapeutic approaches to treat cancer.

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Figure 1.