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Angiogenesis in Melanoma

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Summary

The process of angiogenesis is crucial for progression and metastasis of the majority of solid tumors including melanomas. The purpose of this review is to summarize existing knowledge of the mechanisms of angiogenesis in melanoma as well as current anti-angiogenic therapeutic strategies and their targets. Here, we have focused primarily on the role of key growth factors which are secreted by melanoma cells and known to trigger angiogenic responses, and their receptors as expressed on both endothelial and melanoma cells. Many of these growth factors function in synergy with receptors for extracellular matrix, integrins and matrix metalloproteinases. All these systems of molecules are activated during major stages of angiogenesis such as endothelial migration, proliferation and reorganization of surrounding extracellular matrix. The blockade of these molecules and their downstream pathways leads to inhibition of melanoma vascularization. Thus, these classes of molecules are essential for melanoma angiogenesis and, therefore, might serve as promising targets for therapeutic intervention. Many recently developed compounds targeting key pathways in angiogenesis are in their final stages of clinical trials.

Keywords

Melanoma; Angiogenesis; Extracellular matrix; Growth factors; Integrins; Matrix metalloproteinase

Introduction

Angiogenesis, the process of formation of neovasculature from pre-existing blood vessels, is widely considered as an essential process to ensure the supply of nutrients and oxygen to rapidly growing tumors as well as to provide a route for tumor cell metastasis¹. Angiogenesis is a distinct feature of several human cutaneous melanomas and was first described by Warren and Shubik following transplantation of human melanoma tumor tissue into a hamster cheek pouch². These studies were later confirmed by Hubler and Wolf ³. Rapid angiogenesis of cutaneous melanomas dramatically enhances the risk of lethality and contributes to the progression of the most common type of cancer in young adults.

Like any other cutaneous neoplasia, melanomas follow discrete sequential transformation processes in which the nevus attains dysplastic radial growth phase (Figure.1). This is followed by a vertical growth phase $^{4-6}$, The vertical growth phase requires high angiogenic activity, which, in turn, contributes to melanoma cell metastasis. However, the most critical step in

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metastasis is the spreading of melanomas to lymphatic vessels surrounding the tumor tissue. Following lymph node invasion, melanoma cells metastasize to the lung, liver, cerebrum, and other sites. Exposure to ultraviolet radiation is known to cause genetic changes in skin, which modulate the cutaneous immune response and increase production of several growth factors 7^{-9} . This causes uncontrolled proliferation of melanocytes, which, in turn, dramatically

increases the consumption of oxygen and nutrients eventually leading to cell starvation and hypoxia^{10,11}. To fulfill this increasing demand additional vasculature needs to be developed. Thus, to increase the blood supply, the tissue begins to produce a spectrum of growth factors which trigger the process of angiogenesis.

Studies using human melanoma xenograft models in nude mice indicated that melanoma tumor cells serve as a source of several growth factors including but not limited to vascular endothelial growth factor (VEGF) and basic fibroblast growth factor (bFGF), two powerful stimulators of rapid angiogenesis¹². Despite recent advances in our understanding of the molecular mechanisms of tumor growth/metastasis and the availability of modern technology to detect and treat various human cancers, our knowledge about the mechanisms underlying cutaneous tumor angiogenesis and metastasis remains rudimentary.

In this review we have focused on the role of cytokines and growth factors secreted by melanoma cells in tumor angiogenesis. These growth factors are involved in both autocrine and paracrine regulation of melanoma progression. Importantly, growth factors and their receptors function in intimate relationship with extracellular matrix proteins, their receptors, integrins, and matrix metalloproteinases. These classes of molecules are essential for melanoma angiogenesis and, therefore, serve as promising targets for therapeutic intervention. Most of the relatively recently developed compounds are at the final stages of clinical trials. Thus, this review is intended to summarize the basic biological knowledge regarding melanoma tumor angiogenesis as well as current clinical approaches to treat late stages of malignant melanomas.

Growth factor dependent melanoma tumor growth and angiogenesis

Angiogenesis in melanoma is stimulated by a variety of growth factors such as VEGF, basic FGF, acidic FGF, platelet derived growth factor (PDGF), and transforming growth factors α and β (TGF- α and β) are among the others (Figure 2). Angiogenesis serves as a turning point in melanoma tumor growth and metastasis.

VEGF and FGF

Transformed melanocytes are known to produce high amounts of bFGF as well as VEGF during conversion of the dysplastic naevus to the vertical growth phase. This extensive growth factor production continues until new blood vessel formation is complete. Recent reports indicate that basal expression of VEGF by transformed melanocytes plays a more important role in promoting angiogenesis in melanomas than hypoxia-induced VEGF up-regulation^{13–15}. At the same time, bFGF regulates endothelial cell proliferation and angiogenesis by both autocrine and paracrine mechanisms. Since bFGF is devoid of the classic signal peptide for secretion, tumor cells release this factor by exocytosis from endoplasmic reticulum. Significant amounts of bFGF were found to be associated with extracellular matrix as well as with basement membrane of the newly formed blood vessels in human melanomas¹⁶. Digestion of extracellular matrix by matrix metalloproteinases of melanoma or endothelial origin promotes release of matrix bound bFGF, which, in turn, stimulates endothelial cell proliferation and vascular tube formation in melanomas^{17,18}. Therefore drugs targeting these molecules are currently under various stages of clinical trial (Table 1).

PIGF

Another important stimulator of melanoma angiogenesis is placental growth factor (PIGF). PIGF exists in several isoforms that are generated by alternative splicing of the same gene product and it exhibits distinct heparin-binding properties. PIGF-1 and -2 are expressed by melanoma cells and are known to bind to neuropilin-1 and -2 receptors expressed on endothelial cells¹⁹. In addition, PIGF acts through binding to VEGF receptor-1. Also, PIGF forms heterodimers with VEGF and, as a result, is able to bind to VEGF receptor-2 on endothelial cells (Figure 2). PIGF enhances melanoma tumor angiogenesis not only by acting on pre-existing endothelial cells, but also by mobilization and recruitment of VEGFR-1 positive hematopoietic precursors from bone marrow²⁰. Further, PIGF is known to contribute to angiogenesis by acting on VEGFR-1-expressing smooth muscle cells/pericytes, thereby enhancing blood vessel maturation. PIGF secreted by both primary and metastatic melanomas induces tumor cell proliferation, indicating PIGF might contribute to melanoma tumor growth through an autocrine mechanism. Several studies also indicated that PIGF mediated neovascularization resulted in rapid vessel branching and formation of large and stable blood vessels in various in vivo models²¹.

IL-8 and TGF-1

Several growth factors and their receptors are implicated in the pathogenesis of a variety of human melanomas since their upregulation is associated with melanoma progression. Among these is interleukin-8 (IL-8). Its expression was found to be very minimal in normal epidermis and benign melanocytic lesions. However, it is dramatically increased in a majority of cutaneous melanomas examined. In a clinical study analysis IL-8 serum levels in melanoma patients are significantly elevated compared to healthy individuals²². Further, serum IL-8 levels correlate with advanced disease stage as well as with overall survival of melanoma patients. A recent study by Liu et al²³ indicated that transforming growth factor-1 (TGF-1) is able to enhance expression of IL-8 in human melanoma cells and promote angiogenesis in several mouse xenograft models. Several studies conclusively demonstrated that while tumor cell-derived IL-8 directly supports the growth of melanomas, IL-8 of endothelial origin further promotes melanoma cell migration. Additionally, tumor-derived IL-8 is able to induce endothelial cell migration (Figure 2).

IL-8 is also known to modulate vascular permeability. It functions through activation of Gprotein coupled receptors on endothelial cells, leading to enhanced actin stress fiber formation, which, in turn, results in cell retraction and gap formation between endothelial cells Experimental evidence from several laboratories indicated that over-expression of IL-8 in poorly vascularized and nonmetastatic melanoma cells resulted in enhanced angiogenesis, rapid tumor growth and increased metastatic potential of melanomas^{24,25}.

Plasminogen system

Besides a number of growth factors discussed above, there are other important players in melanoma which might eventually serve as therapeutic targets. Urokinase plasminogen activator and its receptor (uPA/uPAR) have been demonstrated to play a crucial role in several stages of melanoma tumor progression including melanoma cell migration, invasion and metastasis²⁶. At the same time, uPA secreted from melanoma tumor cells is able to regulate endothelial cell functions including migration and the organization of endothelial cells into tube-like structures. Analysis of biopsy specimens of skin lesions indicated that uPA expression highly correlated with disease progression in a majority of patients. Delbaldo et al²⁷ demonstrated that uPA and plasminogen activator inhibitor type 1 mRNAs accumulate in atypical naevocytes and in melanoma cells, but not in benign naevocytes. Further, these observations suggest that up-regulation of the uPA gene is an early event of melanocyte transformation and that unbalanced enzyme activity is associated with the malignant

phenotype. Studies from Hearing et al²⁸ indicated that mouse melanoma cells express the single chain form of uPA on their cell surface and that these cells are capable of plasminogendependent extracellular matrix degradation. Further, melanoma cells that had been treated with anti-uPA antibody showed significantly decreased metastatic activity compared to controls. Studies from Min et al²⁹ demonstrated that molecules which prevent uPA binding to its receptor are able to inhibit bFGF-induced neovascularization in vivo and mouse melanoma tumor growth in syngeneic mice. Additional studies also demonstrated inhibition of melanoma tumor cell metastasis by uPA inhibitors in similar models. Taken together, these studies suggest that uPA/uPAR system is required for melanoma tumor angiogenesis and growth, as well as metastasis (Figure 2). Therefore, antagonists of the uPA/uPAR system might be utilized as potential inhibitors of tumor progression through several mechanisms^{30,31}.

Integrin signaling in melanoma tumor progression and angiogenesis

Integrins are heterodimeric transmembrane glycoproteins consisting of α and β subunits. Their large ectodomain is required for ligand binding, while a short cytoplasmic domain transfers signals from the extracellular environment to the cytoplasm via a short transmembrane domain (Figure 3). Integrins mediate cellular processes such as migration, invasion, proliferation and metastasis in a variety of human cancers including melanomas³². Even though integrins are classically described as cell adhesion molecules, recent advances in integrin biology indicate that they play a very significant role in signal transduction, gene expression, cellular proliferation and regulation of apoptosis or anoikis of normal as well as transformed cells (Figure 3). Beyond this, integrins also play very crucial roles in embryogenesis, inflammation, immunity, tissue differentiation, regulation of cell growth, hemostasis and angiogenesis³³.

Integrins $\alpha_{\nu}\beta_3$ and $\alpha_{\nu}\beta_5$ are known to upregulate on endothelial cells during tumor neovascularization³⁴. Several $\alpha_{\nu}\beta_3$ integrin ligands including vitronectin, osteopontin and bone sialoprotein are known to modulate both VEGF as well as FGF-2-induced tumor angiogenesis in mouse models³⁵. Studies from several groups indicate that $\alpha_{\nu}\beta_3$ integrin plays a very crucial role in the progression of cutaneous melanomas from the benign radial growth phase to the metastatic vertical growth phase³⁶. Studies utilizing $\alpha_{\nu}\beta_3$ integrin inhibitory peptides, antibodies or small molecular compounds indicate that $\alpha_{\nu}\beta_3$ integrin indeed is required for tumor endothelial cell survival, therefore its inhibition leads to the regression of tumor vasculature³⁷. These studies further indicate that the expression of α_{ν} , α_2 , α_3 and α_4 integrin subunits are elevated in malignant melanoma tumor cells compared to normal melanocytes. Similarly, laminin receptor $\alpha_6\beta_4$ integrin expression is significantly reduced in melanoma cells compared to benign cells³⁸. In contrast, clinical samples of human malignant tissue show a significant increase in $\alpha_6\beta_4$ expression³⁹. Therefore, the role of $\alpha_6\beta_4$ integrin in melanoma tumor growth and metastasis is controversial. Among the beta subunits, expression of β_1 and β_3 are more frequently elevated in several tumor types, including melanomas (Table 2).

Other major functions assigned to integrins expressed by melanoma cells are to support ECM anchorage-dependent tumor cell proliferation and anchorage-independent invasion. Integrins are also crucial in tumor angiogenesis and tumor cell metastasis⁴⁰. Expression analysis at different stages of melanoma progression have indicated that the levels of β_3 and β_1 integrins promote melanoma transition from the radial growth phase to the vertical growth phase⁴¹. This phenomenon is further connected with rapid tumor neovascularization as well as tumor cell metastasis. Another interesting phenomenon observed in melanoma patients is the enhanced expression of integrin ligands. One such $\alpha_v\beta_3$ integrin ligand is osteopontin⁴². Constitutive expression of osteopontin in dermal cells leads to sustained anti-apoptotic signaling in melanocytes and a high rate of proliferation. A recent study utilizing numerous clinical samples of melanomas in various clinical grades indicated that the expression of osteopontin is

positively correlated with an invasive phenotype⁴³. Further clinical studies have indicated that serum osteopontin levels are significantly higher in patients with benign or metastatic tumors compared to healthy individuals⁴⁴. Therefore, serum osteopontin levels could potentially be a prognostic marker for various human cancers.

Studies from various laboratories indicate that expression levels of $\alpha_2\beta_1$ and $\alpha_5\beta_1$ are higher during the vertical growth phase of melanoma compared to the radial growth phase or malignant transformation^{45–47}. Melanoma cells during the vertical growth phase bind to collagen type-1 through $\alpha_2\beta_1$ and $\alpha_5\beta_1$ integrins (Table 2). This stimulates the expression of MMP-1 and -2, which are crucial for collagen fibril degradation in the dermis which in turn facilitates vertical spreading of melanomas. The accumulated, denatured collagen acts as the $\alpha_v\beta_3$ integrin ligand as the denaturation process exposes RGD sequences⁴⁸. The binding of denatured collagen to $\alpha_v\beta_3$ integrin further stimulates MMP-2 expression in these cells, increasing their invasive potential. One of the major photolytic degradation products due to over-expression of MMPs in dermis is fibronectin. The fibronectin receptor $\alpha_5\beta_1$ is expressed abundantly in most melanomas studied. Over- expression of $\alpha_5\beta_1$ in mouse melanoma cells leads to enhanced expression of MMP-2 and MMP-7⁴⁹. To summarize, activation of integrins leads to enhanced expression of MMPs, which in turn degrade ECM components thereby generating a series of integrin ligands. These integrin ligands generated by photolytic degradation bind to the invasive melanomas and activate the signaling cascades required for malignant transformation.

As integrins are known to positively regulate angiogenesis, tumor growth and metastasis, several inhibitors of integrins are currently under clinical trials⁵⁰. Most clinical trials are focused on inhibitors of the $\alpha_{v}\beta_{3}$ integrin complex or α_{v} integrin alone³⁷. These include cilengitide, ATN-161, CNTO-95 and vitaxin (the last two are humanized monoclonal antibodies). These compounds specifically mask ligand binding sites and promote the internalization of targeted integrins. Peptide integrin inhibitors currently under clinical trials include cilengitide and ATN-161. Cilengitide is a cyclic RGD peptide that specifically inhibits $\alpha_{\nu}\beta_{3}$ and $\alpha_{\nu}\beta_{5}$ integrin function. In preclinical studies, cilengitide substantially reduced tumor growth in a mouse melanoma xenograft model. Even though cilengitide has been in clinical trials for some time, the outcome of this trial has yet be published 34. Similarly, the peptide integrin inhibitor ATN-161 is still under phase 1 clinical trials and data indicates that it exhibits anti-angiogenic and anti-metastatic activities⁵¹. Results of phase 1 clinical trials of vitaxin indicated that it stabilized disease and reduced the risk of metastases. However, results of phase 2 clinical trials indicated a very modest response and were not very encouraging. Therefore, in recent phase 2 clinical trials, vitaxin was administrated in combination with a standard chemotherapeutic compound (dacarbazine). Beyond this, several integrin inhibitors including small molecule compounds are currently in the preclinical phase of development. E7820, an aromatic sulfonamide derivative known to inhibit α_2 integrin, entered its phase 1 clinical trial in early 2004⁵⁰. Similarly, volociximab, a humanized monoclonal antibody specifically targeting $\alpha_5\beta_1$ integrin is also currently in phase 2 clinical trial. Phase 1 clinical studies aimed to determine the optimal concentration did not find any dose limiting toxicity of this antibody. Overall, integrin inhibitor clinical trials are encouraging and several small molecule compounds have successfully completed preclinical studies.

Role of matrix metalloproteinases in melanoma angiogenesis

Matrix metalloproteinases (MMPs) are the major class of proteases that play crucial roles in tissue remodeling during embryonic development, bone resorption, wound healing and angiogenesis. Their catalytic activity is regulated partially by tissue inhibitors of matrix metalloproteinases and it has been demonstrated by several research groups that MMPs produced by actively proliferating tumor cells facilitate angiogenesis, tumor growth and metastasis⁵². As MMPs are actively involved in efficient matrix degradation, MMP expression

and catalytic activity are tightly regulated, at the stages of transcription, post-translational extracellular activation, and by suppression by its inhibitors⁵³. Several studies indicated that the basal level of MMP production in benign or normal melanocytes is typically low and expression of MMPs is highly correlated with disease progression. MMP activation is accomplished by removal of the N-terminal propeptide domain through exogenous or autocatalytic cleavage⁵⁴. Previous studies demonstrated that serine proteases such as plasmin activate most of the MMPs through this mechanism. MMP-2, which is abundantly expressed in early stages of malignant transformation, is known to achieve activation in a membraneassociated manner in endothelial cells and melanoma tumor cells. Further, cell surface associated membrane-type matrix metalloproteinase (MT1-MMP) is also known to activate MMP-2 through this mechanism⁵⁵. To date the most extensively studied MMPs in melanomas are MMP-2 and MMP-9. It has been demonstrated by several groups that the expression and activation of these enzymes has been correlated to the invasive and metastatic phenotypes of melanomas⁵⁶. Previous reports indicated that MMP-2 and MMP-9 are constitutively expressed in malignant melanomas and their expression is highly associated with melanoma atypia and dedifferentiation in melanocytic lesions⁵⁷.

Currently, cell surface associations of secreted MMPs through post translational modification have developed wide interest in the scientific community. It has been previously demonstrated that MMP-2/TIMP-2 associated with the cell surface in melanomas exhibits enhanced catalytic activity against its substrates compared to MMPs in secreted phase. Malignant melanoma cells are known to express a number of MMPs, including MMP-1, -2, -9, -13, and -14, as well as inhibitors of MMPs such as TIMP-1, -2 and -3^{56} . A recently published study from Kerkela et al clearly demonstrates a specific distribution of MMPs within cutaneous squamous cell carcinomas⁵⁷. Another recent clinical study also indicated that increased MMP-2 expression in melanomas was highly correlated with metastasis. Further, increases in expression of MMPs were shown to highly correlate with low survival rates in patients with malignant melanoma tumors⁵⁸. It is also very important to note that not only expression of MMPs, but also their functional activity, is required for malignant tumor progression. Genetic overexpression of MT1-MMP in melanoma cells induced activation of MMP-2 and this activation is crucial for extracellular matrix degradation when localized on the leading edge of invasive carcinomas. In a clinical study of human melanoma lesions consisting of different stages of tumor progression it was found that MMP-2 and MT1-MMP positive tumor cells were often restricted to the interface between the tumor stroma and the invasive part of the tumor⁵⁷. Surprisingly, expression of MMPs is not restricted to tumor cells but is also found abundantly in stromal cells. This clearly indicates a major contribution of host-derived proteases to melanoma tumor progression.

One of the major MMPs found to be expressed in human melanoma is MMP-1⁵⁹. A series of studies has also indicated that MMP-1 expression is highly associated with malignant melanoma progression. In vitro studies indicated that degradation of collagen types I and IV and tumor cell invasion through Matrigel required MMP-1 expression (Table 3). Other than MMP-1 and -2, the major MMP expressed in melanoma tumor cells is MMP-9 which is also referred to as gelatinase B^{60} . MMP-9 expression in melanoma tumor cells was found exclusively during the horizontal growth phase but not during the vertical growth phase. This clearly suggests that expression of MMP-9 expression was only detected in advanced stages of disease, not in early melanocyte lesions⁶¹. Further, melanomas expressing constitutively high levels of MMP-9 exhibited increased lung colonization in experimental lung metastasis models. These advancements in understanding of MMP-9 biology indicate that MMPs expressed either by neoplastic or stromal cells are important in the metastasis of melanomas⁶².

Several studies using either model cell lines or animals have demonstrated that the balance between MMPs and their inhibitors finally determines melanoma tumor progression⁶³⁻⁶⁸. To date, tissue inhibitors of matrix metalloproteinases (TIMPs) are extensively studied as they are natural inhibitors of MMPs and hence could be potential therapeutic targets. Numerous conclusive studies demonstrate that overexpression of (TIMP) -1,-2 and -3 significantly reduces melanoma tumor cell invasion, migration, tumor growth and metastasis⁶⁹. Further, several studies have indicated that TIMPs significantly reduce tumor neovascularization in the several tumor models studied. Even though TIMPs are known to inhibit tumor cell metastasis in several experimental animal models, in human melanoma cells TIMP expression significantly enhances tumor cell proliferation⁷⁰. Therefore, the role of TIMPs in melanoma tumor growth remains controversial.

As MMPs are known to play very crucial roles during the processes of tumor progression, several inhibitors specifically targeting MMPs are currently undergoing clinical trials. In the early '90s MMP inhibitors generated great enthusiasm among several research groups wishing to take them to clinical trials. Preclinical trials of MMP inhibitors were very promising, showing minimum side effects compared to other drugs available at the time⁷¹. Several current inhibitors, which have been tested in preclinical and clinical trials, are broad category MMP inhibitors. Pharmacological inhibitors such as batimastat and its analog marimastat, which interfere with the catalytic site of the MMPs, were the first inhibitors studied in detail. A recent review by Coussens et al discusses the status of several MMP inhibitors in clinical trials 72 . The first clinical trial of MMP inhibitors started in 1997 with marimastat and prinomastat. Phase 1 and 2 clinical trials were mostly focused on the optimal biological dose of MMP inhibitor rather than clinical outcome. Phase 2 and 3 clinical trials include three major strategies: (1) the clinical outcomes of MMP inhibitors were directly compared with standard chemotherapy. (2) MMP inhibitors were used in combination with standard chemotherapy. (3) Effects of MMP inhibitors were directly compared with placebo, which was administered in those patients who had no clinical evidence of disease after standard chemotherapy 71. Therefore, the results of the MMP inhibitor clinical trials were very conclusive. Most of the MMP inhibitors tested in clinical trials were not very promising due to the lack of positive outcomes and the appearance of substantial drug side effects, which were not observed in preclinical studies. Therefore, most of the MMP inhibitor clinical trials were terminated following phase 3 clinical trials⁷¹. This failure of early MMP inhibitor clinical trials substantially suppressed the initiation of new clinical trials of MMP inhibitors targeting cancer.

Anti-angiogenic therapies in melanoma

Tumor vasculature is essential for a tumor's growth, progression and metastasis to distant sites. According to the original theory of Judah Folkman, the destruction of tumor vasculature must initiate the process of tumor suppression and death⁷³. Based on the results of the most recently published studies, inhibition of angiogenesis might make tumors highly susceptible to radiation and chemotherapy⁷⁴. Most importantly, the process of angiogenesis generally is not required for the normal physiology of normal adult organisms, with the exception of the female reproductive cycle or recovery from an injury. Therefore, minimal side effects are anticipated from the inhibition of this process. Based on preclinical data, one might predict the side effects will be restricted to impaired wound healing, a process known to be dependent on angiogenesis. Taken together, the predicted specificity and effectiveness of the approach initiated an interest in angiogenesis as a therapeutic target.

It is not surprising that numerous anti-angiogenic agents are currently in clinical trials or in development^{75–78}. Table 4 shows several classes of anti-angiogenic compounds divided into groups based on their molecular specificity. The list of molecular targets for anti-angiogenesis therapy includes: 1) Components or fragments of extracellular matrix and metalloproteinase

inhibitors^{79,80}; 2) Angiogenic growth factors and their receptors (expressed both on tumor as well as on endothelial cells). This category includes monoclonal antibodies and soluble forms of receptors designed to bind and neutralize growth factors, small molecules designed to inhibit growth factor-receptor interaction or tyrosine kinase activity of receptors^{81–83}. In addition, we included oligosaccharide-based inhibitors of growth factor release from extracellular matrix into this category. 3) A class of compounds which targets cellular receptors for extracellular matrix, integrins expressed on both melanoma and endothelial cells⁸⁴. There are monoclonal antibodies, peptides and small molecule inhibitors in this category. All of the drugs presented in Table 4 have demonstrated significant inhibition of tumor angiogenesis and tumor growth in preclinical studies. Many experimental models included melanoma⁸⁵.

At present, it is quite intriguing to follow the results of clinical studies. Not all of the trials have been reported and discussed in detail in the literature. As a result, the overall picture of the efficacy of various groups of anti-angiogenic compounds is constantly changing, making any sort of generalization challenging. An interest in anti-angiogenic therapy dramatically increased after the US Food and Drug Administration (FDA) approved bevacizumab (Avastin) for several forms of cancers⁸⁶. Inhibition of angiogenesis by bevacizumab appears to be a promising and effective strategy for solid tumors. It has been successfully used alone as well as in combination with chemotherapy treatments in several types of cancers⁸⁶. Bevacizumab is relatively well tolerated; however, some observed side effects were surprising and not predicted based on the known mechanism of action for this antibody. Thus, it appears that additional preclinical studies will be required to fully address the effects of bevacizumab and its underlying mechanism on normal functional characteristics of vascular and hemostatic systems. A PUBMED search using the combination of key words "bevacizumab and melanoma" in the clinical trials category resulted in "0" hits. However, based on information from the NIH website "www.ClinicalTrials.gov" it appears that at least seven studies of bevacizumab as monotherapy or in combination with other compounds (carboplatin, paclitaxel, sorafenib, dacarbazine, interferon-alpha, imatinib were listed) in melanoma are currently recruiting patients. Results of bevacizumab clinical trials in melanoma are listed as "to be published" on

http://www.centerwatch.com/patient/trialresults/druglst_Avastin_bevacizumab.html Since bevacizumab was the first FDA approved anti-angiogenic compound, it will most likely serve as a gold standard for other therapies which are currently in development. However, it is clear that drugs designed to target the same pathway, or even the same molecule, are not anticipated to display similar efficacies or similar profiles of side effects in melanoma or other cancer patients.

In sum, the process of angiogenesis in melanoma is crucial for tumor development and metastasis and undoubtedly requires further detailed investigation of its underlying mechanisms. Since this process involves a synergistic action of several classes of molecules and signaling pathways, there are numerous possibilities for the development of non-overlapping therapeutic strategies. Several angiogenic compounds are already in clinical trials for melanoma and available preliminary results are encouraging.

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Figure 1. Malignant progression of melanoma

Aberrant proliferation of normal melanocytes leads to formation of benign naevus. The rapidly proliferating transformed melanocytes form dysplastic naevi, which are asymmetric with irregular borders. This is followed by a rapid radial growth phase, which is characterized by intraepidermal growth. The last step in benign melanoma growth is the vertical growth phase. It is also called the dermal invasion phase. Malignant melanomas develop vasculature very rapidly providing a route for further metastasis.



Figure 2. Cell surface receptor mimicry between malignant melanoma cells and endothelial cells Malignant melanoma cells express VEGFR-2, VEGFR-1 and co-receptor neuropilin-1/2 which are commonly not expressed on most of the cancer cells. Therefore, VEGF is able to induce similar intracellular signaling responses in both endothelial and melanoma cells. Placental growth factor can independently bind to VEGFR-1 and neuropilin-1/2 to induce intracellular signaling. PIGF also form heterodimeric complexes with VEGF and interact with VEGFR-2. Other receptors expressed on both melanoma and endothelial cells include uPA receptor, CXCR-1/2 and FGFR-1. Therefore, there is a spectrum of growth factors/chemokines which are able to induce similar signaling responses in both cell types.



Figure 3. Regulation of integrin signaling by ECM proteins and growth factor receptors

Integrins and their ligand expression are up-regulated in most of the human melanomas. Ligand binding to the extracellular domain of specific integrin leads to initiation of signaling events required for a broad spectrum of processes such as cellular proliferation, survival, and actin cytoskeleton regulation. Integrin interactions with its ligands are the prerequisites for cellular adhesion, invasion and migration. Growth factors, via receptors expressed on endothelial cells or melanoma cells, are able to regulate integrin function activity, i.e. its affinity and avidity of interactions with ECM proteins. The process of functional crosstalk between growth factor receptors and integrins is regulated by several intracellular signaling molecules which are able to shuffle between growth factor receptors and integrin cytoplasmic domains.

Table 1

Major growth factors expressed in malignant melanoma and clinical significance

Molecule	Functions	Clinical Trial	Reference
VEGF/VEGFR	Enhance EC proliferation, migration and angiogenesis	Phase 2/3	[¹⁴],[¹⁵]
FGF/FGFR	Enhance expression of VEGF and uPA and prevent EC apoptosis	Phase 1	[¹⁷], [¹⁸]
PIGF	Recruit bone marrow-derived hematopoietic precursors site of angiogenesis	None	[19], [20]
IL-8	Induce tumor neovascularization and tumor cell proliferation	Phase 2	$[^{24}], [^{25}]$
uPA	Increase tumor vasculature and migration and metastasis of tumor cells	Phase 1	$[^{30}], [^{31}]$
FGF/FGFR PIGF IL-8 uPA	Enhance expression of VEGF and uPA and prevent EC apoptosis Recruit bone marrow-derived hematopoietic precursors site of angiogenesis Induce tumor neovascularization and tumor cell proliferation Increase tumor vasculature and migration and metastasis of tumor cells	Phase 1 None Phase 2 Phase 1	$ \begin{bmatrix} 17\\ 19\\ 19\\ 14\\ 124\\ 16 1 1 1 1 1 $

Table 2

Major integrin receptors expressed on endothelial/malignant melanoma cells and clinical significance

<u> </u>			<u> </u>	
Integrin Receptor	Functions	Expression	Clinical Trial	Reference
$\alpha_v \beta_3$	Enhance tumor vasculature and tumor growth	EC, MM	Phase 2	[³⁶]
$\alpha_v \beta_5$	Similar to $\alpha_v \beta_3$ integrin induce tumor growth and angiogenesis	EC	Phase 2	[³⁴]
$\alpha_5\beta_1$	Essential for metastasis of melanoma cells	EC, MM	Phase 2	[⁴¹]
$\alpha_6\beta_4$	Required for vascular maturation, associated with aggressive behavior of tumor cells	EC, MM	None	[³⁸]
$\alpha_2\beta_1$	Essential for endothelial and tumor cell interaction with basal lamina	EC, MM	Phase 1	[⁴⁵]
$\alpha_3\beta_1$	Required for malignant progression of melanoma cells	MM	None	[⁴⁶]
$\alpha_4\beta_1$	Required for melanoma cell metastasis	EC, MM	Phase 1/2	[⁴⁷]

					Table 3	
Maj	or MMPs ex	pressed in n	nalignant	melanoma	and clinical	significance

MMP	Functions	Clinical Trial	Reference
MMP-1	Collagen type I and IV degradation and melanoma cell invasion	Phase 2/3	⁵⁹
MMP-2	Melanoma cell invasion and angiogenesis	Phase 2/3	[⁶³]
MMP-3	Promotes tumor growth and angiogenesis	Phase 2/3	[⁶⁴]
MMP-7	Growth, invasion, metastasis of tumor cells and angiogenesis	Phase 1/2	[⁶⁵]
MMP-9	Tumor cell invasion, metastasis and angiogenesis	Phase 1 to 3	$[^{62}]$
MMP-10	Malignant melanoma tumor cell invasion and metastasis	None	[⁶⁶]
MMP-13	Expression correlates with tumor progression and aggressive tumor behavior	None	[⁶⁷]
MMP-14	Melanoma tumor cell invasion and angiogenesis	None	[⁶⁸]

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Maior angiogeni	c molecules targeted for	cancer therapy. clinical a	Table 4 purpaches and outcome.				
Category	Molecular Target	Class	Name	Preclinical studies	Clinical trials status in melanoma or other cancers	Some observed side effects	Refs
Angiogenic growth factors and its receptors	VEGF	Monoclonal antibody	Bevacizumab	Very potent inhibitor of melanomas in several in vivo	Phase II (response observed)	Minimal toxicity, delayed wound healing	75
	Tyrosine kinase receptors	Small molecule	Sorafenib	Inducts Inhibitor of various tumors in vivo and in vitro	Phase II (minimal effect as a	Well tolerated	76
G	VEGF	Soluble VEGF receptor	VEGF-trap	Potent inhibitor in various in vivo	sıngle agent) Phase I/II	Same as above	77
	VEGF receptor-1	Small molecule	Semaxinib (SU-5416)	Effective in vivo	Phase II (partial response	Deep vein thrombosis	78
	VEGF receptors	Small molecule	PTK/ZK	Effective in vivo (including	observed) Phase I/II	Well tolerated	81
	Heparanase inhibitor (VEGF and FGF)	Mixture of oligosaccharides	PI-88	Inclanomas) Inhibitor of VEGF/ FGF release and tumor growth and	Phase I (partial response, recommended	Well tolerated, thrombocytopenia	82
., ,	VEGF receptor-2	Monoclonal antibody	DC101	Very potent in vivo	IOT pnase II) 	I	83
for extracellular	Integrin $\alpha v \beta_3$	Monoclonal antibody	Vitaxin (MEDI-522)	Potent in vivo inhibitor of tumor growth and	Phase I (recommended for phase II)	Limited toxicity (MI)	37
integrins,	αν integrins alphavbeta3/alphav beta5	Monoclonal antibody Small molecule	CNTO95 Cilengitide (EMD 121974)	Same	Phase I/II Phase I/II	Well tolerated Well tolerated	84 34
Components	MMPs	Small molecule	Batimastat	Inhibits melanoma growth and lung	Phase I/II, limited activity	Poor solubility	79
extracentuar matrix and profeases	MMPs Extracellular matrix	Small molecule Fragment of matrix	Marimastat Endostatin	Inclusions in Ince Inhibits melanoma growth in vivo Potent angiogenesis inhibitor in vitro and in vivo	Phase <i>VI</i> Phase I	Muscle pain and stiffness Minimal toxicity	80
Complex	Angiogenesis inhibitors	Small molecule	Thalidomide and lenalidomide	Effective in vivo	Phase II	Neuropathy, skin rash	78
of action	and minimionroutators Cyclooxygenase-2	Small molecule	Celecoxib	Effective in vivo	Phase I/II (shows response)	Well tolerated	85

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