**Review Article**

**Integrin-Mediated Cell-Matrix Interaction in Physiological and Pathological Blood Vessel Formation**

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Physiological as well as pathological blood vessel formation are fundamentally dependent on cell-matrix interaction. Integrins, a family of major cell adhesion receptors, play a pivotal role in development, maintenance, and remodeling of the vasculature. Cell migration, invasion, and remodeling of the extracellular matrix (ECM) are integrin-regulated processes, and the expression of certain integrins also correlates with tumor progression. Recent advances in the understanding of how integrins are involved in the regulation of blood vessel formation and remodeling during tumor progression are highlighted. The increasing knowledge of integrin function at the molecular level, together with the growing repertoire of integrin inhibitors which allow their selective pharmacological manipulation, makes integrins suited as potential diagnostic markers and therapeutic targets.

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**1. Introduction**

Invasive cancer is among the leading causes of death worldwide, and rates are still increasing, due to ageing and changes in lifestyle [1]. Cancer is a collective term for many diseases, rather than a single disease, with the common characteristic that tissue growth goes haywire [2]. Patients who have undergone cancer treatment show an increased risk of developing a second tumor, mainly due to the same risk factors that were responsible for the first tumor but also in part due to the treatment of the first tumor with mutagenic chemotherapeutics or radiation [3]. Therefore, new strategies for cancer treatment with as little as possible adverse side effects are needed that effectively eradicate the primary tumor and also do not increase the risk of recurrence.

A tumor initially grows without any connection to the vasculature until it reaches a critical size of about two mm in diameter. Then it remains in a dormant state, in which proliferation and apoptosis due to lack of oxygen, are in a dynamic equilibrium unless it develops in a well-vascularized region or is able to recruit its own vasculature. Hanahan and Weinberg have proposed six hallmarks of cancer, one of them being the induction of angiogenesis [4, 5]. For further growth, the tumor needs to hook up to the vascular system by forming neovessels.

During tumor progression, an angiogenic switch is activated causing a continuous neovessel formation emanating from the normally quiescent vasculature, which sustains tumor growth [6]. This process called tumor angiogenesis is a collective term that is generally used for all types of tumor neovascularization. In addition to vessel co-option and to endothelial cell (EC) sprouting, tumor vessels can also develop by intussusceptive or glomerular angiogenesis, or, in a way of vascular mimicry, even tumor cells themselves can form vessel-like hollow structures. These types of vessel formation can occur in parallel, and also gradual transitions are possible. Vessel formation by the latter types requires less energy than sprouting angiogenesis, is thus carried out faster, and usually can be observed in, for example, gliosarcoma multiforme, melanoma, and breast and colon cancer [7].

For neovessel formation, ECs need to migrate into a previously avascular region and to extensively remodel the extracellular matrix (ECM). In this process, integrins, which are cell adhesion receptors for various ECM proteins and immunoglobulin superfamily molecules, are the most important matrix receptors [8, 9]. Therefore, integrins are
appealing targets for cancer therapy using a variety of integrin-specific antagonists, ranging from endogenous antagonists over humanized or chimeric antibodies to peptides and small nonpeptidic compounds [10–12].

In this paper, based on the general assembly of blood vessels, the specific organization of tumor vasculature will be described, as well as the dynamic sequence of events by which a tumor gains access to the body's vasculature. In this context, the role of integrins and possibilities of their pharmacological manipulation are explored.

2. The Static Picture: The Extracellular Matrix of Blood Vessels

The tissue's ECM is a structure-shaping molecular scaffold and also a repository for cytokines and other growth factors [13]. Cells embedded in this matrix need to be supplied with oxygen and nutrients, signaling molecules need to be received and emitted, and metabolic waste products need to be disposed of. These tasks are optimally fulfilled by the cardiovascular system with its intricate and dynamic network of blood vessels. Depending on their functions, different types of blood vessels show special histological and molecular adaptations. The heart, as a double-acting pump, drives the blood circulation within the vasculature via the aorta through arteries and arterioles into capillaries, from where the blood flows back through venules and veins. Due to the prevailing pressure conditions, the body fluid is forced through the vessel wall to form the lymph, which then is drained by lymph vessels back to the blood circulation. Additionally, the vasculature serves as "highway" system for leukocytes to patrol the body during immunological surveillance and to quickly reach sites of inflammation. The vascular wall is capable of self-sealing upon smaller injuries, and leukocytes are able to penetrate the blood vessel wall in a complex interplay without any obvious vessel leakage. Pathologically, tumor cells capitalize the blood vessel system to disseminate from a primary tumor and to colonize distant organs where they develop metastases.

2.1. General Organization of the Vessel Wall. Histologically, the walls of blood vessels comprise three concentric layers, that is, tunica intima, tunica media, and tunica adventitia [14], which are separated by two sheet-like structures of ECM proteins. The membrana limitans interna and externa establish a border between tunica media and tunica interna and adventitia, respectively. These ECM sheaths tightly connect the cell layers of the vessel wall to form a functional unit, which becomes evident when too weak cell-matrix interactions lead to life-threatening aneurysms.

The tunica intima comprises a single layer of squamous ECs and lines the inner surface of all blood vessels. The tunica media, which is usually the thickest layer in arteries, is composed of mural cells, which are smooth muscle cells in larger blood vessels and pericytes in capillaries. The tunica adventitia finally interconnects the blood vessel with the surrounding connective tissue, and it is usually most prominent in veins.

In different vessel types, that is, arteries, arterioles, capillaries, venules, and veins, this general blueprint is modified corresponding to the respective functional requirements. For example, endothelia, which are continuous in most instances, can become fenestrated, as in exocrine or endocrine gland tissues, or even discontinuous, as in liver, spleen, or bone marrow, in order to facilitate the exchange of hormones or metabolites. Elastic and muscular arteries illustrate other examples for a modification of this general blueprint. In order to even the pulsatile blood flow coming from the heart, the proteins elastin and fibrillin are abundant in the tunica media ECM of elastic arteries, which is the direct cause for the vessel wall's elastic properties. Muscular arteries possess numerous concentric sheaths of smooth muscle cells. By means of vasoconstriction and vasodilation, they can distribute and direct the blood to different organs.

2.2. Extracellular Matrix in the Vessel Wall. The ECM of blood vessels together with their resident cells contributes to essentially all physiological functions of blood vessels and has been reviewed recently [15].

The subendothelial basement membrane (BM) compartmentalizes the vessel's single-layered endothelium from the vascular connective tissue. The molecular architecture of BMs has recently been reviewed [16–18]. Fibronectin, incorporated between endothelial and perivascular cells, is essential for blood vessel morphogenesis [19]. The presence of von Willebrand factor (vWF) is characteristic for the subendothelial BM, where also other BM proteins, such as the network-forming collagens IV and XVIII can be found, together with laminins, nidogens, and perlecan. Thirteen different collagens are present in the vascular wall [20, 21]. The network-forming collagen IV [22] plays a key role for the mechanical stability of the BM [23], which, especially in arterial regions of the circulatory system, has to withstand a considerable blood pressure.

In the tunica media of elastic and muscular arteries, covalently crosslinked supramolecular aggregates of elastin form concentric lamellae and fibers in a proportion of up to 50% of the vessel's dry weight and confer resilience to pulsatile blood flow [24–26]. Regions of the ECM that consist mostly of elastin are confined by EMILINs, that is, homotrimeric elastin microfibril interphase-located proteins [27]. Anchored to microfibrillar bridges of fibrillin-1 and fibrillin-5 between these concentric elastin lamellae, vascular smooth muscle cells (VSMCs) are sandwiched in a fishbone-like pattern and thus can effectively regulate the vessel's caliber [25, 28–31]. Dependent on the vessel type, distinct fibulins are involved in the assembly of the ECM. While fibulin-1 is widespread and occurs in the BMs of all blood vessels, heart valves and septa, fibrulin-3, and fibrulin-4 occur in the walls of capillaries and larger blood vessels [32]. The innermost and outermost elastic lamellae are referred to as membrana limitans interna and membrana limitans externa, respectively. Between the elastic lamellae, type I and III collagens are deposited that bear tensile forces exerted on the vessels and limit their elastic dilatability. In contrast, in the interstitial connective tissue between the subendothelial membrane and the membrana limitans interna, type VI and
type VIII collagens are found [21, 33]. The connection of the membrane limitans interna to the subendothelial BM by type XVIII collagen is assumed [34]. Also type XVI collagen, which is produced by VSMCs and found close to both elastic microfibrils and fibrillar type I and type III collagens, may contribute to the connection between the elastic and collagenous phases of the ECM [35, 36], especially, as type XVI collagen contains a binding site for the major collagen receptor on VSMCs, integrin α1β1 [37, 38].

The ECM of the tunica media is synthesized by VSMCs, which are all encapsulated by an (incomplete) BM containing the usual BM proteins, type IV collagen and laminins [33, 39]. Depending on microenvironmental cues, VSMCs can reversibly acquire distinct phenotypes, which can be characterized as either (i) contractile and differentiated or (ii) secretory, migratory, and less differentiated [37, 39]. Under physiological conditions, the contractile phenotype prevails, at which the VSMCs transduce forces on the pericellular matrix especially by the collagen-binding integrin α1β1, by the laminin-binding integrin α7β1 and by dystroglycan [37]. In contrast, in the secretary, proliferatory, and migratory phenotype, the integrin equipment of the VSMCs predominantly consists of the fibronectin receptor, α5β1, and the integrins α4β1 and α9β1. Consistently, in the proximity of secretory VSMCs, the fibronectin splice variants V (IIICS) and EIIIA with binding sites for the integrins α4β1, α5β1, and α9β1 are abundant [39]. In capillaries, scattered pericytes, each encapsulated by an own BM, stabilize the endothelium and its subendothelial BM [40–42].

The fibroelastic connective tissue of the tunica adventitia connects the blood vessel with the perivascular connective tissue. It is rich in versican, a glycoprotein, which can interact with fibrillin-1 [43], fibulin-1 [44], and fibulin-2 [45], as well as with other ECM molecules.

2.3. Receptors for ECM Molecules. To interact with their microenvironment and to spatiotemporally regulate their differentiation state, morphology, metabolism, and survival, cells are equipped with a variety of receptors for all the ECM molecules [13]. Integrins are the largest family of these receptors, and they mediate adhesion to collagens, laminins, and fibronectin. In addition, there are other receptors and coreceptors, such as the syndecans [46].

Binding to a wide variety of different ECM molecules and transmitting signals bi-directionally in an outside-in and inside-out manner, integrins constitute functional hubs, which, according to an interesting concept in network theory and systems biology, integrate networks of angiogenic signaling cues that orchestrate the behavior of ECs and VSMCs during angiogenesis [47, 48]. Thus, therapeutically targeting integrins as the operationally important circuit-integrating hubs rather than single pathways of the complex system may result in a more pronounced inhibition of angiogenesis [47].

ECs express the vitronectin receptors αvβ3 and αvβ5; moreover, on ECs and pericytes the following integrins are expressed: the collagen receptors α1β1 and α2β1, the laminin receptors α3β1, α3β6, and α6β4, the osteopontin receptor α9β1, and the fibronectin receptors α4β1 and α5β1 [49]. Pericytes additionally express the laminin receptor α7β1, and the osteopontin receptor α8β1, and integrin αvβ3 is also expressed on glial cells [49].

As EC-derived tumors, angiosarcomas express the integrins α1β1, α2β1, α3β1, α5β1, and α6β1, and in benign and malignant mesenchymal tumors as well as in the desmoplastic stroma of carcinomas, integrins α1β1 and α5β1 are widely distributed [50]. Integrins α1β1 and α2β1 bind to the same ligand in the ECM and are VEGF-dependently upregulated on migrating ECs, and antagonists against both integrins inhibit VEGF-mediated angiogenesis without affecting the existing vasculature [51, 52]. Therefore, and against the background of gene ablation studies, they are believed to differentially regulate angiogenesis [49]. Important coreceptors for integrin α2β1 are the syndecans-1 and -4, which weaken the invasiveness of tumor cells into a collagenous matrix [53].

Cells bind to fibronectin and vitronectin preferentially via the RGD-dependent integrins αvβ3 and α5β1 [54]. Fibronectin can also be bound by the leukocyte-specific integrins α4β1 and α4β7 [55]. Cell-fibronectin interactions are modulated by proteoglycans, glycoproteins of the ECM, and the coreceptors syndecans [56].

Integrin αvβ3 was identified as a marker for angiogenic vascular tissue [57]. In contrast to quiescent ECs, integrin αvβ3 is highly expressed on activated ECs during tumor angiogenesis, as well as on some tumor cells [58, 59]. In the tumor microenvironment, angiogenic ECs can interact due to their increased levels of the integrins αvβ3 and αvβ5 with provisional matrix proteins, such as vitronectin, fibrinogen, vWF, osteopontin, and fibronectin. Also, partially proteolyzed collagen in the tumor exposes RGD sites and is a further ligand for integrin αvβ3 [60]. Thus, the ECM of the tumor microenvironment both provides survival signals and facilitates invasion. Integrin-αvβ3-mediated adhesion to platelets protects malignant cells from clearance through the immune system, and moreover, αvβ3 integrin also helps tumor cells to adhere to the vessel endothelium and to spread into adjacent tissues [61].

The pharmacological inhibition of integrin-αvβ3-mediated cell-matrix interaction impedes tumor angiogenesis and growth [62], as does a replacement of the β3 subunit with a mutated nonphosphorylatable subunit in a murine model [63], which provides evidence for a proangiogenic role of integrin αvβ3, in contrast to integrin αvβ5, which does not seem to play an essential role in angiogenesis [64]. Interestingly, the analysis of αv-knock-out mice revealed that, despite being embryonic or perinatally lethal, the vascular endothelium was not impaired in the absence of the αv subunit, whereas the primary cause of death was brain hemorrhage [65–67]. Also endothelial Tie-2-specific knockout of the αv subunit did not result in any vascular or angiogenesis defect [67]. Moreover, in an integrin subunit β3- and also β5-deficient mouse model, pathologic angiogenesis and tumor growth are increased [68]. A possible cause for these seemingly contradictory phenomena could be a relief of a transdominant inhibition by αvβ3 on other integrins or other molecules, which would enhance their proangiogenic function [69, 70]. Likewise, there could be a compensatory role of other integrins with overlapping function [49]. Moreover, inhibition could also stabilize the
integrin αvβ3 in its unligated conformation and thus induce apoptosis by triggering an integrin-mediated death program [71].

Integrin αvβ8 is important for vascular development in the embryonic brain and in the yolk sac [72]. It is expressed on astrocytes but not on ECs or pericytes, nevertheless plays an important role in angiogenesis, as it binds in addition to several ECM proteins also to the latency-associated peptide (LAP) of TGFβ1, which in cooperation with the membrane-type metalloproteinase MT1-MMP/MMP14 results in activation of TGFβ and triggering of its downstream signal cascades [73–75].

Collagen IV, an essential component of BMs, is bound by integrin α1β1, which is expressed on mesenchymal cells and can also bind to other collagens [76, 77]. Further collagen-binding integrins are α2β1, the main receptor for fibrillar collagens, which is expressed on epithelial and some mesenchymal cells as well as on thymocytes [78], α10β1 in cartilage [79], and α1β1, a key receptor for fibrillar collagen on fibroblasts [80]. The integrins α1β1 and α2β1 are involved in the regulation of collagen and MMP synthesis and thus of special importance for ECM turnover [81–83]. Discoidin domain receptors DDR1 on epithelial cells and DDR2 on mesenchymal cells are further collagen receptors with tyrosine kinase function and are relevant for cancer [84]. Other collagen receptors are glycoprotein GPIV on platelets [85], the leukocyte-associated immunoglobulin-like receptor LAIR-1/CD305 [86], and the urokinase-type plasminogen activator receptor-associated protein uPARAP/Endo180, which is involved in matrix turnover during malignancy [87].

Laminin, as a further integral component of BMs, is bound by the integrins α3β1, α6β1, α6β4, and α7β1 [88–91] and also by α-dystroglycan [92, 93] and by the 67 kDa laminin receptor 67LR [94]. 67LR is increased in various tumors and correlates with their metastatic potential [95, 96]. The different laminin receptors may also act cooperatively in laminin binding, for example, laminin-binding β1 integrins and 67LR [97] or integrin α6β4 and syndecan 1 [98].

Integrin α3β1, which in the vascular wall binds to laminins-411 (laminin 8) and-511 (laminin 10), thrombospondin (TSP), TIMP2, tetraspanin CD151, and to the C-terminal domain of the collagen IV α3 chain, is controversially ascribed either a positive or a negative role in angiogenesis (cf. [99]).

There is controversy whether the hemidesmosomal integrin α6β4, which is expressed on a subset of ECs [100] and on tumor ECs [101], aggravates pathological angiogenesis [101] or whether it is a negative regulator of angiogenesis that is downregulated at its onset [102].

Thus, many molecules of the ECM scaffold, for example, laminins, collagens, fibronectin, and vitronectin, are ligands for integrins that link the cell’s cytoskeleton to the ECM. Loss of this matrix-integrin contact triggers apoptotic cell death [103]. Picking up signals from the cell’s microenvironment, integrins functionally sense, interpret, and distribute information, which allows the cell to modulate its proliferation, differentiation, migration, and shape [104]. The modulatory and regulating function of integrins is emphasized by direct interaction with a multitude of proteins, such as MMPs, uPA/uPAR, tissue inhibitor of matrixmetalloproteinase-2 (TIMP-2), vWF, TSP-1, osteopontin, syndecan-1, insulin-receptor substrate-1 (IRS-1), cytohesin-1, integrin cytoplasmic domain-associated protein-1 (ICAP-1), integrin-linked kinase (ILK), calcium- and integrin-binding protein (CIB), β3-endotoxin, talin, actinin, tensin, nischarin, and the Ras-related protein Rab 25 [9].

The subendothelial BM of the tunica intima serves as a mechanical support to which ECs are anchored by various adhesion molecules, especially integrins [46, 105–108]. Additionally, the subendothelial BM provides microenvironmental information that regulate the metabolic activity of attached ECs, such as their production of leukocyte adhesion molecules [107] or antithrombotic prostacyclins [109], as well as other properties, for example, the tightness of intercellular contacts [108]. Therefore, angiogenesis is regulated not least by integrins which are adhesion receptors for matricellular proteins, ECM proteins, and immunoglobulin superfamily molecules, on nearly all cells including ECs [8, 58].

In addition to their mechanical function [110], integrins also assist growth factor receptors and play important roles in signaling processes, in particular as soluble growth factors, and other signaling molecules are bound by integrins as well [111]. For example, the proangiogenic VEGF-A165 is bound by integrins αvβ3 and α3β1 [112] and also by the tenascin-C- and osteopontin-receptor integrin α9β1 [113]. The latter integrin, furthermore, binds the lymphangiogenic growth factors VEGF-C and VEGF-D [114]. Angiopoietins-1 and -2 are bound by integrin α5β1 [115]. Integrin α6β1 is a receptor for the proangiogenic CCN-family member CYR61, and is involved in in vivo tube formation [116, 117]. The fibronectin receptor integrin αvβ3, which is the best-studied integrin in relation to angiogenesis and is upregulated during wound healing and retinal vascularization and especially on tumor blood vessels, also binds to fibroblast growth factor FGF-1 [118]. Semaphorin 7A binding is also reported for the collagen receptor integrin α1β1 [119].

Stimulated by PDGF, vascular smooth muscle cells express the laminin receptor integrin α7β1, which plays an important role in recruitment and differentiation of VSMCs [120, 121].

Integrin α9β1 is not only involved in lymphangiogenesis [114] but also plays a role in EC adhesion [122]. While binding of TSP-1 to integrin α9β1 promotes angiogenesis [123], VEGF-A is another ligand of integrin α9β1 [115].

2.4. Vascular-Relevant Integrin-Deficient Mouse Models. The crucial involvement of integrins in EC biology has been elucidated substantially by the examination of genetic knock-out studies [124]. By ablation of the respective genes, the EC integrins α1β1, α2β1, α4β1, α5β1, α6β1, α6β4, α9β1, αvβ3, and αvβ5 as well as the VSMC integrin α7β1 and the glial cell integrin αvβ8 have been implicated in regulation of cell growth, survival, and migration during angiogenesis (for recent reviews of the findings from knock-out mice cf. [8, 10]). However, due to redundancy and compensatory
mechanisms, the interpretation of knock-out results is often difficult.

Itgb1−/− mice die at E5.5 before they start to develop their vasculature [125, 126]. Mice with a conditional knock-out in Tie-2-positive ECs survive until E9.5–E10.5, and they are capable of vasculogenesis, but their angiogenesis is disturbed showing defects in sprouting and branching [127–129]. Another endothelial-specific knockout of the integrin β1 subunit is mediated via VE-cadherin-Cre recombinase and becomes manifest later in embryogenesis resulting in lethality between E13.5 and E17.5 [130]. In this mouse model, loss of β1 integrin leads to a decreased expression of the cell polarity gene PAR3 and thus to disruption of EC polarity and lumen formation [130].

Itga1−/− mice, deficient for the collagen-binding integrin α1β1, show a normal vascular development and a reduced tumor angiogenesis in adulthood, which has been attributed to increased MMP activity [131], while α2β1-deficient Itga2−/− mice show an enhanced tumor angiogenesis in adulthood, but an otherwise normal vascular development [131, 132], and integrin α2β1 is involved in the PlGF-dependent regulation of VEGFR-1 [132]. Although integrin α1β1 and α2β1 bind to the same ligand in the ECM, their differential knockout results in opposing effects on angiogenesis, suggesting a regulatory role for this pair of integrins.

Da Silva and coworkers generated EC-specific conditional α3 integrin knock-out mice and showed that these mice, in contrast to a global ablation, are viable and fertile but display enhanced tumor growth, elevated hypoxia-induced retinal angiogenesis and tumor angiogenesis, and increased VEGF-mediated neovascularization [99]. The authors also could show that α3β1 is a positive regulator of EC-derived VEGF, which again represses VEGFR2 expression. Their data demonstrated that endothelial α3β1 negatively regulates pathological angiogenesis and implicated an unexpected role for low levels of EC-derived VEGF as an activator of neovascularization.

Itga4−/− mice, deficient for fibronectin- and VCAM1-binding integrin α4β1, are embryonic lethal with 50% dying at E9.5–10.5 due to failure of chorion-allantois fusion and 50% dying at E11.5 due to cardiovascular defects [55].

Mice, which by ablation of Itgα5 are deficient for the fibronectin receptor integrin α5β1, show normal vasculogenesis but no angiogenesis, which results in embryonic lethality at E10-11 due to defects in posterior somites, yolk sac, and embryonic vessels [133, 134]. This demonstrates the requirement of the integrin α5 subunit during embryonic development of early blood vessels and other tissues. Accordingly, integrin α5β1, which is poorly expressed on normal quiescent ECs, is markedly upregulated during tumor angiogenesis [135].

Among the laminin-binding integrins, integrin α6 is not essentially required for vascular development, although α6-deficiency is lethal with skin blistering defects resembling epidermolysis bullosa [136]. In line with the α6 knock-out mice, Itgb4−/− mice, lacking a functional laminin-binding integrin α6β4 by deletion of its signaling domain, show normal vascular development, although with reduced angiogenesis [101], but die of severe skin defects [100]. In neovascularization, the endothelial expression of integrin α6β1 is downregulated [102]. While it is not required for EC proliferation and survival, it promotes tumor angiogenesis [101]. In contrast, genetic ablation of α7β1, which is expressed on VSMCs but not on ECs, leads to incomplete cerebral vascularization and hemorrhage and also to placental vascular defects, which results in partial embryonic lethality and demonstrates that integrin α7β1 is important for recruitment and survival of VSMCs [121, 137].

Deletion of Itga8 resulting in lack of integrin α8β1, a receptor for fibronectin and tenasin, results in partial embryonic lethality, but no defects in vascular development (Müller and Reichardt, cited in [138]).

Itga9−/− mice lacking integrin α9β1, which is the receptor for tenasin-C, osteopontin, VCAM-1, and also for VEGF-A, -C, and -D [113, 114], have defects in large lymphatic vessels and die postnatally at P8-12 from a bilateral chylothorax [139].

Ablation of Itgav, resulting in simultaneous loss of the two integrins αvβ5, a receptor for vitronectin, osteopontin, and Del-1 (developmental locus 1), and αvβ3, a receptor for a variety of ECM proteins, such as fibronectin, vitronectin, laminins, fibrinogen, fibrin, TSP, tenasin-C, vWF, denatured collagen, osteopontin, MMP-2, Del-1, bone sialoprotein, FGF-2, thrombin, and CCN1 (cystein-rich protein 61), leads to 80% embryonic lethality at E9.5, and the other 20% die at P0 with brain hemorrhage [65]. On the other hand, Itgb3−/− mice, which are just integrin-αvβ3 deficient, show 50% embryonic and early postnatal lethality and an enhanced angiogenesis in surviving adult animals, indicating that this integrin is not strictly required for vascular development [140]. Surprisingly, animals with an intact but nonfunctional β3 integrin subunit develop normally but show defects in angiogenesis in adulthood [63]. In contrast, Itgb5−/− animals lacking integrin αvβ5 develop normally and angiogenesis is not significantly affected, indicating that this integrin is not mandatory for vascular development [64]. Integrins β3 and β5 doubly deficient mice show enhanced tumor growth and angiogenesis. This strongly suggests that these integrins are not required for vascular development or for pathological angiogenesis, pointing out that the mode of action of αvβ3 antagonists and antiangiogenic therapeutics is still insufficiently understood [68]. Ablation of Itgb8 leads to the loss of integrin αvβ8 on glial cells and thus to disrupted blood vessel formation in the brain, thereby demonstrating that this integrin is mandatory for brain’s blood vessel development [72]. Moreover, the phenotype of β8-deficient mice resembles that of αv-deficient mice, which provides evidence that most defects in αv-deficient mice are due to the loss of integrin αvβ8 [72].

2.5. Integrin Structure. The family of integrins contains 24 structurally related N-glycosylated heterodimeric proteins assembled noncovalently from 18 α-subunits and eight β-subunits. Each subunit comprises a large extracellular domain, a single transmembrane domain, and with the exception of the β4 integrin subunit, a short noncatalytic cytoplasmic tail [141]. Integrins are of special importance as
they mediate cell matrix crosstalk via both outside-in and inside-out signaling [54, 142]. Moreover, the 24 different integrins possess promiscuous and redundant ligand specificities, which is of importance when distinct signals are to be transduced or when in a particular context a defined cellular response is elicited, as is discussed by Rüegg and Alghisi [11].

Integrin structure and function have been studied in detail at the molecular level [143, 144]. The extracellular headpiece is formed by a disk-like propeller domain of the α subunit and globular domains of the β subunit [145, 146]. The joint globular head harbors the ligand-binding site [146, 147]. The crystal structure of the integrin-αvβ3-binding site with an inserted RGD ligand [148] helped to map functional amino acid residues on other integrins [149]. Recently, the binding pocket of integrin α5β1 has been mapped by swapping regions of zebrafish and human α5 subunit in a gain-of-function approach [150].

2.6. Integrin Signaling. Depending on their activity, integrins adopt distinct conformations (Figure 1). In the inactive resting conformation, the headpiece of the heterodimer bends towards the plasma membrane, and the transmembrane domains of the α and β subunits are associated [146]. Upon ligand binding, the previously bent integrin ectodomain adopts an activated upright conformation [106, 151]. This conformational change is conveyed through the transmembrane domains towards the cytoplasmic tails [54, 105, 152], where cytoskeletal proteins and signaling molecules relay the incoming signal intracellularly [153]. In inside-out signaling, the binding of intracellular molecules, such as talin or kindlins [154, 155], to the cytoplasmic integrin tails leads via a separation of the transmembrane domains [156] to a switch blade-like erection of the extracellular domains [147, 157, 158]. Likewise, in outside-in signaling, ECM ligand binding to the integrin headpiece also induces a conformational change in the hybrid domain and thereby a separation of the integrin subunits' legs [144]. This parting of the legs separates the cytosolic tails and allows binding of cytosolic proteins and thus clustering of integrins and formation of focal adhesion sites (Figure 1).

By clustering into focal adhesions, integrins recruit talin, paxillin, α-actinin, tensin, and vinculin and thereby mechanically couple the ECM scaffold to the actin cytoskeleton. Additionally, integrins bind scaffolding molecules, such as p130 CRK/BCAR1, and recruit and activate kinases, such as focal adhesion kinases (FAKs), Src family kinases (SFKs), and integrin-linked kinase (ILK), the latter forming a complex with the adapter molecules parvin and PINCH/LIMS1 [159].

Figure 1: Integrin activation. Integrins are a family of heterodimeric transmembrane adhesion receptors that bidirectionally relay signals with the extracellular matrix (ECM) and also with other cells. When activated, a conformational change increases the affinity, and clustering increases the avidity towards the ligand. (1) By inside-out signaling, integrins can reversibly undergo a conformational change from a bent inactive to an upright activated conformation with intermediate ligand affinity, at which the cytoplasmic domains are still close together. (2) Upon ligand binding, the integrin adopts a high-affinity conformation with a concomitant parting of the legs and a separation of the cytosolic α- and β-tails that unlocks docking sites for cytosolic molecules. (3) Clustering of ligand-occupied and activated integrins establishes a mechanical link between ECM and cytoskeleton and leads to the recruitment of scaffolding molecules and kinases. (4) The assembly of focal adhesions triggers intracellular signaling cascades. Details can be found in the text.
In addition, tetraspanins can recruit integrins to membrane microdomains, thus regulating integrin function [160]. Thereby, the rather unstable nascent adhesions are transformed into focal complexes, focal adhesions, fibrillar adhesions, or podosomes. This clustering of integrins leads to a reorganization of the plasma membrane around the focal adhesion into caveolin-containing lipid rafts, to which also growth factor receptors often localize, and to the assembly of adhesion signaling complexes [161–163]. This allows a regulation of growth factor signals by integrin-mediated caveolae trafficking [164, 165]. In the assembly of such integrin adhesions, up to 156 distinct molecules, amongst other adaptors, proteins, kinases, and phosphatases, are involved [48, 163]. Membrane lipid-protein interactions that modulate the homo- or heterotypic association of receptor molecules in the cell surface, or between adjacent cells, have been reviewed recently [166]. From the focal adhesion sites signal pathways diverge that regulate diverse cellular programs, such as adhesion, migration, proliferation, and survival. To provide an overview, integrins generally relay their signals via the FAK, ERK, and NF-κB pathways [153].

In most cases, in mechanosensory signaling FAK, Src, and SH2, domains containing protein tyrosine phosphatase 2 (SHP2) are involved [167]. Upon integrin binding, FAK autophosphorylates and binds to Src, which further phosphorylates FAK and several downstream binding partners, amongst others, JNK and Rho [168–170].

Activated FAK also recruits PI3K, which mediates the activation of AKT and procures integrin-mediated cell survival, and likewise the antiapoptotic AKT can be activated via Ang-1 [171]. Moreover, signals relayed via integrins and Src can be integrated by FAK with growth factor receptor-relayed signals via Ras, MEK, and MAPK [172]. Growth factors can activate Ras signaling independently from integrin-related adhesion signals. Nevertheless, MEK1 and Raf1 are important interfaces between integrin-relayed and growth-factor-relayed signaling, because both MEK1 and Raf1 need to be activated via adhesion-mediated activation of Src and FAK in order to activate MAPK [173, 174].

An endothelial-specific ablation of FAK results in impaired blood vessel development and embryonic lethality [175]. Downstream of FAK, Src couples integrin-mediated and VEGF-receptor-mediated proangiogenic signaling in ECs [176–178]. However, endostatin can also activate Src via integrin α5β1 and thereby disassemble actin stress fibres and focal adhesions and thus inhibit cell migration, which is regulated by integrins via the Ras/ERK pathway [179–181]. Important for adhesion and migration of endothelial and VSMCs are also p130Cas and PLC-γ, which can interact with FAK [182–185].

PI3K is of pivotal importance for angiogenesis, because its deletion results in embryonic lethality E9.5 to E10.5, when angiogenesis is important for vascular development. PI3K deletion also causes decreased Tie-2 expression and thus creates a phenotype resembling Tie-2 deficiency [186, 187]. Moreover, EC-specific deletion of the PI3K isofrom p110α impairs angiogenesis [188]. In ECs, adhesion via integrins elicits a survival signal via FAK/PI3K/mTOR/4E-BP1 and Cap-dependent translation [189]. Furthermore, the activation of PI3K by Ras is important for lymphangiogenesis [190].

In addition to a direct activation of ERK, integrins can also activate a Raf/MEK/ERK signaling cascade [189, 191, 192]. Raf-deficient and MEK-deficient mice have severe vascular defects [193, 194]. Growth-factor-mediated ERK signaling is linked with integrin-mediated signaling via FAK [195]. Integrin-mediated ERK signaling is important for cell proliferation and migration of ECs [191, 196]. Integrin α1β1 is unique among the collagen-binding integrins because it promotes cell proliferation by activating the Ras-Shc-MAPK pathway, and cell cycle progression is regulated via FAK, Rac, and cyclin D by integrin-mediated adhesion and matrix stiffness [197–199].

Integrins can also activate the NF-κB pathway in ECs and protect them from apoptosis [200–202]. Additionally, NF-κB signaling regulates the expression of cyclooxygenase-2 (COX-2), which again is involved in EC spreading and migration and in the induction of VEGF and FGF-2 [177, 203, 204]. However, inhibition of the NF-κB pathway increases angiogenesis pathologically [205].

Integrins alone are not oncogenic, but some oncogenes may depend on integrin signaling for tumor growth and invasion. For example, integrin-triggered FAK signaling is essential for Ras- and PI3K-mediated oncogenesis [206, 207]. Also the expression of the cancer stem cell marker CD44 is integrin-regulated, and it can be speculated that integrin-related signals are needed to maintain a cancer stem cell population [12, 208]. On the other hand, there is evidence that the collagen receptor integrin α2β1 has a tumor-suppressing function [209, 210].

Ligated integrins promote survival, whereas unligated integrins recruit caspase-8 to the plasma membrane and promote apoptosis in a process termed integrin-mediated death [71, 211], which differs from anoikis induced by loss of cell adhesion to the ECM [103, 212]. Loss of caspase-8 confers resistance to integrin-mediated death of tumor cells, and unligated integrin αvβ3 promotes the malignancy of such tumors [213, 214]. Cell survival is promoted by integrin ligation-dependent upregulation of BCL2 and FLIP/CFLAR, activation of the PI3K-AKT pathway, NF-κB signaling, and p53 inactivation [176, 202, 215–217]. Survival is also promoted by crosstalk between integrins and growth factor receptors, for example, αvβ3 and FGFR or αvβ5 and VEGFR2 [195, 218].

In various steps of angiogenesis and tumor progression, crosstalk between integrins and growth factor receptors on tumor cells and also on host cells is important. This crosstalk can consist in either an activation of a latent growth factor, a regulation of common pathways for signaling or internalization and recycling, a collaborative or a direct activation, or also a negative regulation [111]. The outcome of a growth factor signal in a particular context is often determined by a synergistic and reciprocal interaction of integrins with growth factor receptors, such as tyrosine kinase receptors like VEGFRs and Tie-2, Met, and FGFR, and semaphorins regulate integrin function as well [111, 219–221]. A complex of VEGF with the fibronectin heparin II domain increases, upon cell binding via integrin α5β1 and
the signaling via VEGFR2 synergistically [222]. Expression of integrin α1β1 on tumor-associated fibroblasts has a tumor-promoting effect, because it upregulates the expression of insulin-like growth factor 2 (IGF2), which is another example of integrin-regulated growth factor signaling [223].

Beside binding ECM proteins and thus regulating adhesion and migration, integrins can also directly interact with pro- and antiangiogenic factors [221]. Integrin α5β1 can bind to matrix-bound VEGFR-1 [224]. In addition, integrin α9β1 can directly interact with VEGF-A, -C, and -D and also with hepatocyte growth factor (HGF) [113, 114, 225]. Moreover, integrin α3β1 and αvβ3 bind VEGF-A165 and VEGF-A189 [112]. FGF is directly bound by integrin αvβ3 [226]. Angiopoietins also can directly interact with many integrins [115, 221, 227, 228].

In the context of a hypoxic tumor microenvironment, it is especially interesting that the expression of integrins α1β1 and α2β1 is upregulated by VEGF [51].

3. The Dynamic Process: Connection of a Tumor to the Host Vasculature

Angiogenesis is an important step in the metastatic cascade, which not only provides the tumor with nutrients but also is a route for dissemination. An important trigger for this is hypoxia [229].

3.1. An Angiogenic Switch Triggers the Angiogenic Cascade.

In avascular tissue regions, an oxygen diffusion limit of about 150 μm restricts tumor growth to just a few millimeters in diameter. Thus, in this prevascular phase of tumor dormancy, there is a dynamic equilibrium between proliferation and hypoxia-induced apoptosis [230]. The dormant phase ceases when a tumor recruits its own vasculature by the secretion of angiogenic factors into its environment [231], a process denoted as angiogenic switch [2, 6]. After this angiogenic switch is thrown, the tumor hooks up to the body’s vascular system and thus resumes its growth.

In tumor development, the establishment of an angiogenic phenotype is a crucial and general step [232–234]. Depending on tumor type and environment, this induction of new vessel sprouting can occur at different stages of the tumor progression pathway, and it leads to exponential macroscopic tumor growth [2, 4, 6]. In addition, recent data indicate that angiogenesis also contributes to the microscopic pre-malignant phase of neoplastic progression [5].

Infiltration of bone-marrow-derived monocytes that differentiate into macrophages can trigger this angiogenic switch in spontaneous tumors by releasing both numerous proangiogenic cytokines, for example, VEGF, TNFa, IL-8, and bFGF [235, 236] and MMPs (e.g., MMP-2, -7, and -9) together with elastase and uPA [236]. These matrix-degrading enzymes loosen the avascular ECM for the angiogenic ingrowth of neovessels.

From the multitude of proangiogenic molecules, such as FGF-1 and -2, G-CSF, HGF, IL-8, PD-ECGF, PGE-1 and -2, PlGF-1, and -2, TGF-α and -β, TNF-α, and VEGF-A through E, only the VEGFs and PlGFs are specific for ECs [230]. VEGF-A, which exists in five splice variants, is the most intensively studied one [237]. Mediated by HIF-1, VEGF-A synergizes with FGF-2. VEGF is upregulated under hypoxic and hypoglycemic conditions prevailing within tumor tissue [230].

The role of chemokines in tumor angiogenesis and neovascularization has been reviewed recently [238]. Tumor cells express CCL2/MCP-1 (C-C-motif ligand 2/monocyte chemotactic protein-1), and thus, tumor-associated macrophages (TAMs) are recruited, resulting in an inflammatory response. These TAMs are again a source for angiogenic growth factors, such as, VEGF and FGF-2 [239, 240]. MCP-1 also mediates the recruitment of mural cells in an Ang-1–dependent manner in an ex vivo model [241].

Multiple sequential steps are required for angiogenesis to be successful and in all steps of this angiogenic cascade integrins, which mediate interactions of cells with surrounding insoluble ECM proteins, in addition to soluble growth factors, play an important role [15]. In a first step, the BM of an existing vessel is degraded by MMPs that are expressed by ECs, such as MMP-1, MMP-2, MMP-9, and MT1-MMP/MMP14 [242–244], at which MMP-9 is required for tumor vasculogenesis rather than angiogenesis [245]. Subsequently, cell-matrix contact influences the outgrowth of tip cells and the proliferation of stalk cells that thereupon form endothelial tubes [246]. A new BM is assembled by newly synthesized BM proteins. Finally, the newly generated capillaries undergo maturation, pruning, and expansion.

3.2. Tumor Vessels Can Arise by Different Types of Vessel Formation.

During embryonic morphogenesis, endothelial precursor cells called angioblasts initiate the body’s vasculature by forming tubes in a process called vasculogenesis. This is subsequently accompanied by sprouting (angiogenesis) of new vessels from already existing ones. Once morphogenesis is completed, the adult vasculature is largely quiescent, except for transient events, such as wound healing or menstruation [247]. However, angiogenesis takes place under many pathological conditions, such as atherosclerosis, endometriosis, osteomyelitis, diabetic retinopathy, rheumatoid arthritis, psoriasis, and tumor growth [230]. During tumor progression, the quiescent vasculature becomes permanently activated to sprout new vessels that enable blood supply and thus help sustain tumor growth [5, 6]. Due to its increased metabolic rate, tumor tissue requires blood supply for expansive growth, which is circumstated by the observation that tumor cells, which are p53 deficient and thus show a reduced apoptosis rate, die beyond an oxygen diffusion limit in the range of 150 μm [248]. Tumor cells proliferate around the continuously formed neovessels which markedly differ from normal vessels in morphology and molecular composition [219, 249]. Tumor vasculature generally appears highly tortuous, chaotic, and disorganized. The vessels themselves are leaky due to a discontinuous endothelium, a poorly formed BM, and a lack of mural cells. In addition, tumor cells sometimes mimic ECs. This poor quality of tumor-associated blood vessels compromises blood flow, impairs drug delivery, and facilitates tumor cell intravasation leading to hematogenous or lymphatic metastasis. In addition to histological vessel malformations,
tumor vessels show an anomalous composition of their ECM, for example, tenascin-C and -W, and the oncofetal fibronectin ED-B splice variants are associated with tumor vessels [250, 251]. ED-B fibronectin is synthesized by neo-fibronectin ED-B splice variants are associated with tumor vessels [250]. ED-B fibronectin mediates survival of tumor stem cells, enhances proliferation, invasiveness, and metastasis in tumor cells, and blocks immunosurveillance [250, 253]. Tenascin-C stimulates angiogenesis in ECs, mediates survival of tumor stem cells, enhances proliferation, invasiveness, and metastasis in tumor cells, and blocks immunosurveillance [250, 253]. Tenascin-W is more strictly associated with tumorigenesis and can be used as a tumor biomarker for breast and colon cancer, because it is undetectable in healthy stroma but overexpressed in the tumor stroma [254, 255].

Vascularization mechanisms in cancer have been reviewed recently [256, 257]. New tumor blood vessels can either arise by vessel co-option or be formed by tumor angiogenesis, but there is also evidence for vasculogenesis or recruitment of circulating bone-marrow-derived endothelial progenitor cells that differentiate into ECs [230, 258–260] (Figure 2(A)). Depending on the tumor type, tumor blood vessels build different and characteristic vascular beds, and, according to the function of the vascular bed and the osmotic pressure of the surrounding tissue, endothelia represent highly heterogeneous “vascular addresses” [230]. Tumor vessels constantly change their shape due to persistent growth, and about 30% of the vasculature comprise arteriovenous shunts bypassing capillaries. The concomitant poor perfusion leads to hypoxia of ECs, which consequently synthesize more proangiogenic molecules and thus crank tumor angiogenesis [230].

3.2.1. Endothelial Sprouting. Endothelial sprouting can be triggered by hypoxia, hypoglycemia, and inflammatory or mechanical stimuli, such as blood pressure, and is regulated by many angiogenic growth factors, such as VEGF, and matrix proteases. When neovessels sprout from capillaries, pericytes are selectively lost, and upon receiving an angiogenic stimulus, select ECs differentiate into tip cells that invade the avascular ECM (Figure 2(B)). These tip cells migrate into the ECM following the stimulatory gradient. Behind the tip cells, other ECs begin to proliferate and, as stalk cells, form cord-like structures. These develop into endothelial tubes [130, 261, 262] that subsequently anastomose and thus allow blood flow. Finally, pericytes and smooth muscle cells are recruited, a new BM is synthesized, and the ECs become quiescent again.

The molecular background of capillary sprouting and the key role of VEGF have been reviewed by Carmeliet [231]. Upon a hypoxic stimulus, VEGF is produced, and as a consequence the endothelium’s permeability is increased and the BM loosened by the activity of MMPs [243, 263] and the urokinase plasminogen activator system [264]. The MMP inducer EMMPRIN/CD147 also upregulates soluble VEGF isoforms 121 and 165 and VEGFR-2 on ECs and thus promotes sprouting angiogenesis [265]. Integrin αvβ3 mediates migration into the fibrin-rich cancer stroma and furthermore can associate with MMP-2, thus enabling ECs to maintain the BM in the sol state and to promote tumor cell invasion [266]. In addition to VEGF, FGF, PDGF, and PIDGF are involved, and Ang-2/Tie-2 signaling regulates the detachment of pericytes. Later, PDGF-BB recruits pericytes and smooth muscle cells to the newly formed EC tube, and TGF-β1 and Ang-1/Tie-2 stabilize the EC-mural cell interaction [231].

3.2.2. Intussusceptive Angiogenesis. Another way of tumor neovascularization is intussusceptive angiogenesis, which represents a nonproliferative and noninvasive mechanism for the enlargement of a capillary plexus by intussusceptive growth, arborization, and remodeling [267] (Figure 2(C)). As this mode of vascularization is mostly independent from EC proliferation and migration, as well as BM degradation, this process is more economical and, occurring within hours or even minutes, is noticeably faster than sprouting
angiogenesis [268]. It begins with the formation of translu-
minal pillars from the EC walls. Their subsequent expansion
splits the preexisting vessel into two, thereby enhancing the
vascular surface. In a subsequent process of arborization,
the disorganized capillary network is remodeled into a
functional tree-like structure by serial pillar formation. In
a final remodeling step, the branching angles are modified,
and the capillary network is pruned. The formation of new
capillaries is initiated by sprouting angiogenesis that is later
accompanied or followed by intussusceptive angiogenesis,
which increases the EC surface [269]. Intussusceptive angi-
ogenesis is synergistically regulated by VEGF and Ang-1,
and it seems to be induced by laminar shear stress on the
vessel walls, whereas oscillating shear favors sprouting
angiogenesis [269].

3.2.3. Glomeruloid Angiogenesis. In many aggressive tumors,
glomeruloid angiogenesis gives rise to complex vascular
structures termed glomeruloid bodies, in which several
microvessels together are ensheathed by a BM of varying
thickness containing sparse pericytes [270] (Figure 2(D)).
The frequency of occurrence of such glomeruloid bodies
is an indication for the tumor’s aggressiveness and the
patient’s survival [271]. The formation of such glomeruloid
bodies is rather a remodeling than true angiogenesis, because
proliferating and migrating tumor cells can actively pull
capillaries of the surrounding host vasculature and adjacent
capillary branching points into the tumor node. Thereby,
formed coiled vascular structures develop subsequently into
glomeruloid bodies that are connected to the surrounding
vasculature via numerous narrowed capillaries [256].

3.2.4. Vessel Co-Option. Malignant cells can initially grow
in the vicinity and along pre-existing microvessels and thus
use the host vasculature for their own benefit (Figure 2(E)).
This co-option of the host vasculature was originally believed
to be limited to the initial phase of tumorigenesis [272].
Meanwhile, however, there is evidence that vessel co-option
might persist during all stages of primary and metastatic
growth of various tumors [256], for example, cutaneous
melanoma, which appears to grow by co-opting the vascular
plexus in its surrounding connective tissue, while there is no
sign of directed vessel ingrowth [273].

Vessel co-option is regulated dependent on the tumor
type and the host environment, but the key regulators are
again VEGF and angiopoietins [272, 274]. Ang-1 binds to
Tie-2 and thus triggers signaling cascades, assuring survival
and quiescence of ECs, and thus causing tumor vessel
maintenance, whereas the nonsignaling Tie-2 ligand Ang-2
acts as a negative regulator and destabilizes the capillary walls
by detachment of pericytes [272, 274]. Subsequently, VEGF
via its receptor VEGFR-2 promotes both survival of ECs and
growth of new vessels [237, 275].

3.2.5. Vascular Mimicry. Aggressive melanomas can form
fluid-filled vessel-like channels without any EC lining in
a nonangiogenic process termed vascular mimicry [276]
(Figure 2(F)). These channels allow perfusion independent
of angiogenesis, and they can arise by two types of vascu-
logenic mimicry, designated the tubular and the patterned
matrix type [277]. These tubular vessel-like networks resemble
the pattern of embryonic vascular networks, and, in their
gene expression pattern, aggressive tumors that form
such channels resemble endothelial, pericytes, and other
precursor stem cells, suggesting that tumor cells might
disguise as embryonic stem-cell-like or other cell types
[256]. Vascular mimicry of the patterned matrix type
looks completely different and is characterized by a fluid-
conducting meshwork of extravascular patterned depositions
of matrix proteins such as lamins, collagens IV and VI, and
heparin sulfate proteoglycans that anastomose with blood
vessels [277–279]. Although it is not yet elucidated how such
channels are connected to the vasculature, the latter type
of vascular mimicry has been reported for many cancers,
such as breast, ovarian, and prostate carcinoma, melanoma,
soft tissue sarcomas, osteosarcoma, and phaeochromocy-
toma [277, 280]. In aggressive melanoma, the expression
of tissue factor pathway-associated genes, such as tissue
factor (TF), TF pathway inhibitor-1 (TFPI-1), and TFPI-
2, is upregulated, suggesting an anticoagulation mechanism
in the channel-forming tumor cells [281]. Fluid propelled
through these channels by a pressure gradient might facilitate
the supply with nutrients and oxygen, and, additionally, this
fluid-conducting network could substitute for a lymphatic
vascular system and drain extravasated interstitial fluid
in tumors that lack lymphatic vessels, for example, uveal
melanoma [279, 280].

4. Manipulation of Cell Matrix Interaction in
Tumor Angiogenesis

Cell-matrix interactions regulate signaling pathways that are
intricately interconnected with cytokine-regulated pathways,
which complicates the analysis of their contribution to a
particular step in angiogenesis [153]. ECM receptors can be
manipulated with a wide variety of different compounds
ranging from endogenous compounds, such as matrikines,
over their synthetic analogues and peptides mimicking only
integrin-binding sites to function-blocking antibodies and
small molecules with integrin inhibitory function. Other
starting points for an antiangiogenic therapy are the inhibi-
tion of signaling cascades downstream of the ECM receptors
or cytokine receptors and as a new avenue the blocking of
microRNAs with antisense RNAs in ECs [282, 283]. An
efficient antivascular cancer therapy can target either the
angiogenic signaling pathways or the vascularization mech-
anism [256]. A combination of conventional chemotherapy
with angiopressive or vascular disrupting therapy is often
problematic and needs careful design [256].

4.1. Pharmacological Intervention of Integrin-ECM Interac-
tion. In addition to soluble growth factors, such as VEGF,
there are several endogenous angiogenesis inhibitors, for
example, endostatin, endorepellin, and tumstatin, which
share the common feature that they all are proteolytic frag-
ments of ECM molecules [284, 285]. In tumor angiogenesis
within a primary tumor, such ECM fragments are generated by the release of MMPs, in order to degrade the BM. This results not only in labile and leaky tumor vessels but at the same time keeps metastases from growing, as these endogenous angiogenesis inhibitors are distributed via the blood stream [230]. Therefore, they are of pharmacological interest with regard to their use as angiogenesis inhibitors. Intensive efforts have been directed towards the development of integrin antagonists for the treatment of cancer and many other diseases, ranging from autoimmune diseases over inflammatory to thrombotic diseases, and their applications seem promising [11, 286]. Integrin-mediated interactions of cells with their surrounding ECM can be manipulated by antibodies, peptides, small nonpeptidic compounds, and endogenous inhibitors (Figure 3). Integrin antagonists with antiangiogenic activities have been reviewed recently with special emphasis on drugs that are in clinical trials [11].

Spurred by the success in pharmacologically targeting RGD-dependent integrins, there are also attempts to pharmacologically manipulate RGD-independent integrins, such as the collagen- and laminin-binding integrins, as reviewed recently [287]. The collagen-binding subgroup of integrins with their common A domain comprises interesting targets in the development of drugs against thrombosis, inflammatory diseases, and cancer. TSPs-1 and -2 are naturally occurring potent angiogenesis inhibitors, and their antiangiogenic effects can be imitated by short-peptide mimetics that among other targets bind to β1 integrins [288, 289].

An endogenous inhibitor, which blocks the interaction of integrin α1β1 with collagen I and also binds to heparan sulfate proteoglycans, is arresten, the C-terminal fragment of the collagen IV α1 chain [290, 291]. Endorepellin, a C-terminal fragment of perlecan specifically blocks the function of integrin α2β1 [292] and interestingly also binds to endostatin, thus counteracting its antiangiogenic effect [293]. Additionally, integrin α1β1 can be specifically inhibited with obtustatin from the snake venom of Vipera lebetina obtusa [294, 295]. The interaction of integrin α2β1 with collagen can be specifically inhibited with the C-type lectin rhodocetin from the snake venom of Callolase-lasma rhodostoma [296, 297]. In addition, it can also be selectively antagonized by the protein angiocidin, which was first detected in lung carcinoma cells [298, 299]. The aromatic tetracyclic polyketides maggiemyccin and anhydro-maggiemyccin from Streptomyces, which have been described as potential antitumor agents [300], inhibit collagen binding by blocking the A domain of the integrin subunits α1, α2, α11, and to a lesser extent α10 while cell adhesion to fibronectin, mediated by integrins α5β1, αvβ3, αvβ5, is unaffected [301]. Recently, the sulfonamide derivative BTT-3016 has been described as a potent antithrombotic small–molecule inhibitor of integrin α2β1 with only slight effect on other collagen–binding integrins and no effect on fibronectin- or vitronectin-binding integrins [302]. Another sulfonamide derivative, E7820, which does not interfere with integrin–ligand interaction, reduces integrin α2 expression on the mRNA level [303]. Angiogenesis can be inhibited with antibodies against the α subunits of the integrins α1β1 and α2β1, whereas quiescent vessels are not affected [230].

In a phase I clinical trial, endostatin, the C-terminal fragment of collagen XVIII, blocks the function of integrin α5β1 [179, 304, 305] and also binds to heparin and with lower affinities to other heparan sulfate proteoglycans that are involved in growth factor signaling [306, 307]. Endostatin’s antiangiogenic activity can also be mimicked with derived short non-RGD but arginine-rich peptides [308]. Integrin α5β1 can also be blocked by the synthetic non-RGD peptides PH SCN, named ATN-161, [309] and cyclic CRRETAW [310], as well as by the peptide mimetics SJ749 [311] and JSM6427 [312], and it can be inhibited by the affinity-matured humanized chimeric monoclonal antibody M200/volociximab [313].

Angiostatin is a proteolytic fragment of plasminogen that effectively inhibits integrin αvβ3 [314], and its antiangiogenic effect can also be achieved by its isolated kringle-5 domain [315]. Kringle-1 to 3 show the same antiproliferative effect as the whole angiostatin, but hardly inhibit migration, whereas kringle-4 inhibits EC migration but shows only a marginal antiproliferative effect [316]. Other endogenous integrin αvβ3 inhibitors are the collagen XVIII fragment endostatin [304], and the C-terminal fragment of the collagen IV α3-chain termed tumstatin [317], which also binds to integrin α6β1 [318]. Tumstatin has two binding sites for integrin αvβ3. The N-terminal site mediates an antiangiogenic signal, whereas the C-terminal binding site is associated with the antitumor cell activity [318, 319].

Canstatin, the NC1 domain of the collagen IV α2 chain, inhibits both integrins αvβ3 and αvβ5 [320] and seems to interact with integrin α3β1 too [321]. A hemopexin-like domain comprising C-terminal fragment of MMP-2, termed PEX, also antagonizes integrin αvβ3 by preventing its binding to MMP-2 and thus inhibiting proteolytic activity on the cell surface, especially during vessel maturation [322, 323]. Fastatin and other FAS1 domains, which are present in the four human proteins peristin, FEEL1, FEEL2, and βHig-h3, also function via integrin αvβ3 as endogenous regulators of pathogenic angiogenesis [324]. Next to these natural antagonists there is a variety of synthetic RGD—containing peptide inhibitors that mimic a motif that occurs on many ECM molecules, such as fibronectin, vitronectin, fibrinogen, osteopontin, TSP, vWF, and partially degraded collagen. Most integrins of the αv subfamily and the integrins α5β1 and αIIbβ3 bind to this motif. Therefore, adhesion and spreading of ECs to the ECM can be competitively inhibited by RGD peptides, whereby anchorage-dependent ECs undergo apoptosis [230]. To this group belong compounds, such as cilengitide/EMD121974 [325], S137 and S247 [326, 327], the TSP-derived peptide TP508/chrysalin [328], and several integrin αvβ3- and αvβ5-specific peptidomimetics, such as BCH-14661, which preferentially inhibits αvβ3 and BCH-15046, which blocks αvβ3, αvβ5, and α5β1 [329], SCH221153 [330], and ST1646 [331]. Another inhibitor is the non-peptide antibiotic thiolutin, which intracellularly blocks paxillin and thus, indirectly, integrin αvβ3-mediated adhesion to vitronectin [332]. Antibodies against the β3 subunit inhibit contact of ECs to vitronectin and concomitantly VEGF-induced tyrosine phosphorylation of VEGFR-2 in cell culture studies [333]. Moreover, integrin
**Figure 3:** Options to manipulate integrin function. Essential cellular functions, such as adhesion, migration, proliferation, and survival, which are all regulated by integrins, can pharmacologically be manipulated with a panoply of matrikines, antibodies, peptides, and small molecule inhibitors, many of which are used as therapeutic tools in combination with conventional chemo- or radiotherapy to attack tumor cells and vasculature. Details are described in the text.

αvβ3 can be effectively antagonized with the monoclonal antibody LM609/MEDI-552 and its humanized derivative abegrin/etaracizumab/vitaxin [57, 334–337]. In contrast, the humanized anti-αv antibody CNTO95 targets both integrins αvβ3 and αvβ5 [338]. The humanized Fab fragment 17E6/abciximab/ReoPro of the monoclonal antibody c7E3 inhibits the integrins αvβ3 and also αMβ2/Mac-1 [339, 340], whereas the human-specific monoclonal antibody 17E6 targets all αv integrins [341]. Currently, humanized or chimeric integrin antibody antagonists of αvβ3, αvβ5, and α5β1, and peptide inhibitors of these integrins are in clinical trials as antiangiogenic agents [180].

### 5. Applications and Outlook

Integrins and their binding partners are of special interest as potential therapeutic targets, and several are already in clinical trials. However, the results fall short of the initial expectations, pointing out that monotherapy with a single angiogenesis inhibitor is not sufficient to counteract the numerous angiogenic factors involved in tumor progression [231]. Moreover, there are some caveats in aiming at integrins as therapeutic targets. Obviously, integrins are expressed on virtually all cells under physiological as well as pathological conditions, and it is a major challenge to target exclusively integrins on tumor or tumor-associated cells. Another problem is that low concentrations of antagonists alter the signaling of integrins and other receptors. When administered in nanomolar concentrations, the RGD-containing inhibitors cilengitide and S 36578 alter the trafficking of integrins and VEGFR2 in tumor ECs, thus stimulating angiogenesis and tumor growth [342].

Current tumor therapy aims at vessel eradication in order to disrupt the connection of the tumor to the vascular system and thus cut off the supply of nutrients and oxygen. This can be done with compounds that preferentially affect tumor endothelia rather than normal cells, that is, (i) specific angiogenesis inhibitors, (ii) tumor vessel toxins that attack inherent weaknesses in static tumor vessel endothelia and associated vascular structures, and (iii) dual-action compounds [343]. However, within the last years, a paradigm shift has taken place [344, 345]. Vessel normalization by pruning immature vessels and increasing pericytes and BM coverage of the remaining vessels comes to the fore, rather than vessel eradication, because mere antiangiogenic treatment can worsen malignancy [346]. A malformed tumor vasculature creates and aggravates a hypoxic and acidic milieu which hampers drug delivery and perfusion [347–349], and, due to its leaky endothelium, it promotes tumor cell dissemination [346]. Therefore, chemotherapeutic efficacy can be ameliorated by a concomitant vessel normalization therapy which improves delivery and efficacy of cytotoxic drugs and also sensitizes the tumor cells to radiation [345, 350].

In vessel normalization, the interaction of cells with their surrounding ECM via integrins is of special importance. However, many antiangiogenic compounds, for example, ATN-161, endostatin, and integrin inhibitors, show...
The text discusses the role of integrins in tumor therapy, highlighting the paradoxical effects of RGD mimetics and the importance of targeting compounds that can act either anti- or proangiogenic. A promising approach may be a combination therapy that targets several integrins and receptors.

Additionally, integrins can be used as biomarkers to non-invasively assess the efficacy of chemotherapeutic and radiotherapeutic drugs. Integrin-targeted probes can be used to visualize tumor angiogenesis and the response to chemo- and radiotherapy by various imaging methods, such as magnetic resonance imaging (MRI), positron emission tomography (PET), and ultrasonography.

In summary, the above data illustrate the importance of integrins and integrin-binding and signaling proteins in both physiological and pathological blood vessel formation. Thus, they may be potential targets for antiangiogenic tumor therapy. Although our knowledge concerning this matter has increased remarkably within the last years, the understanding is far from complete.

**Abbreviations**

- Ang: Angiopoietin
- BM: Basement membrane
- CRDGK, CRGDKGPD, CRRETAWAC, PHSCN, RGD:
- ECM: Extracellular matrix
- EC: Endothelial cell
- FAK: Focal adhesion kinase
- FGF: Fibroblast growth factor
- G-CSF: Granulocyte colony-stimulating factor
- HGF: Hepatocyte growth factor
- HIF: Hypoxia-inducible factor
- IL: Interleukin
- MAPK: Mitogen-activated protein kinase
- MEK: MAPK/ERK kinase
- MMP: Matrix metalloproteinase
- NC: Noncollagenous
- NF-κB: Nuclear factor κ-light-chain enhancer of activated B cells
- NRP-1: Neuropilin-1
- PDGF: Platelet-derived growth factor
- PD-ECGF: Platelet-derived endothelial cell growth factor
- PGE: Prostaglandin E
- PI3K: Phosphatidylinositol-3 kinase
- PLC: Phospholipase C
- PIGF: Placenta-derived growth factor
- Ras: Rat sarcoma protein
- Src: Sarcoma oncogene
- TGF: Transforming growth factor
- Tie: Tyrosine kinase with immunoglobulin-like and EGF-like domain
- TIMP: Tissue inhibitor of metalloproteinases
TNF: Tumor necrosis factor
uPA(R): Urokinase-type plasminogen activator (receptor)
VCAM: Vascular cell adhesion molecule
VEGF(R): Vascular endothelial growth factor (receptor)
VSMC: Vascular smooth muscle cell
vWF: Von Willebrand factor
TSP: Thrombospondin.

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