

Modes of resistance to anti-angiogenic therapy

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Abstract | Angiogenesis inhibitors targeting the vascular endothelial growth factor (VEGF) signalling pathways are affording demonstrable therapeutic efficacy in mouse models of cancer and in an increasing number of human cancers. However, in both preclinical and clinical settings, the benefits are at best transitory and are followed by a restoration of tumour growth and progression. Emerging data support a proposition that two modes of unconventional resistance underlie such results: evasive resistance, an adaptation to circumvent the specific angiogenic blockade; and intrinsic or pre-existing indifference. Multiple mechanisms can be invoked in different tumour contexts to manifest both evasive and intrinsic resistance, motivating assessment of their prevalence and importance and in turn the design of pharmacological strategies that confer enduring anti-angiogenic therapies.

Hypertension

A medical condition in which the blood pressure is chronically increased.

The long-standing proposition that induction of chronic angiogenesis is a hallmark of cancer is now solidly grounded in a substantial body of research involving genetic and pharmacological perturbation of elements in the vascular regulatory circuitry. The 'angiogenic switch'¹ is increasingly recognized as a rate-limiting secondary event in multistage carcinogenesis², as documented in animal models of cancer and correlated in advanced pre-malignant stages, as well as their malignant derivatives, in a growing list of human cancer types. That this acquired capability is functionally important for manifestation of the disease has been further validated by the approval of angiogenesis inhibitors as cancer therapeutics, most notably ones targeting the vascular endothelial growth factor (VEGF) pro-angiogenic signalling pathways³. The pioneers of the clinical proof-of-concept for angiogenesis inhibitors are [bevacizumab](#) (Avastin, Genentech/Roche), a ligand-trapping monoclonal antibody, and two kinase inhibitors ([sorafenib](#) (Nexavar, Bayer) and [sunitinib](#) (Sutent, Pfizer)) targeting the VEGF receptor (VEGFR) tyrosine kinases, principally VEGFR2 (also known as [KDR](#)). Since March 2008, bevacizumab has been approved for treating patients with late-stage [colon cancer](#), [non-small-cell lung cancer](#) and [breast cancer](#), all in combination with chemotherapy. Sorafenib and sunitinib have both been approved for treating renal carcinoma, a highly vascularized (and angiogenic) tumour type. In addition, sunitinib has been approved for treating gastrointestinal stromal tumours, and sorafenib for hepatocellular carcinomas³⁻⁶. Numerous ongoing clinical trials seek to expand the applications of each of these VEGF

pathway inhibitors, and dozens of other angiogenesis inhibitors (many also targeting VEGF signalling) are being clinically evaluated (see [Angiogenesis Inhibitors Therapy](#) URL and clinical trials URLs in Further information). Moreover, two VEGF pathway inhibitors (the RNA aptamer pegaptanib and a Fab derivative of bevacizumab) have been approved for treating the angiogenic (wet) form of macular degeneration⁷⁻⁹.

Many of the demonstrable clinical benefits and side effects (such as hypertension) of the kinase inhibitors targeting the VEGF signalling pathway (sorafenib, sunitinib and dozens more in various stages of preclinical and clinical testing) can be attributed to inhibition of the activity of the VEGFRs, but it must be emphasized that all are intrinsically (owing to their chemistry) selective but not specific. Thus, virtually every VEGFR kinase inhibitor has an attendant variety of additional moderate-to-high affinity kinase targets in the 'kinome'¹⁰, of which some may convey added therapeutic benefit (such as inhibition of platelet-derived growth factor receptor (PDGFR), in the case of sunitinib), and others may evoke new toxicities. The components of the VEGF signalling pathway, the constellation of drugs developed to inhibit VEGF signalling, the specific mechanistic effects of VEGF signalling and VEGF inhibition, and the clinical trials evaluating their effects are described in depth and discussed in a Review by L. Ellis and D. J. Hicklin also in this Focus issue¹¹. Additional reviews, in this Focus on targeting angiogenesis¹²⁻¹⁴ and elsewhere¹⁵⁻²¹, describe and discuss other targets and strategies for inhibiting tumour angiogenesis.

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At a glance

- Angiogenesis inhibitors targeting the vascular endothelial growth factor (VEGF)-mediated pro-angiogenic signalling pathways are producing demonstrable clinical benefit for an increasing number of cancer types. However, in some cases (both in humans and in mouse models of cancer) anti-angiogenic therapies produce initial responses followed almost inevitably by progression, thereby affording appreciable but limited survival advantage. In other cases there is no objective benefit. Increasing evidence supports the proposition that progression and mortality following a period of benefit reflects an adaptive response by tumours, manifesting 'evasive resistance' to angiogenesis inhibitors. By contrast, patients for whom there is no tangible benefit indicate that an intrinsic resistance to angiogenesis inhibitors exists.
- Evasive resistance to VEGF pathway inhibitors (and arguably other angiogenesis inhibitors) involves a number of distinct and interrelated mechanisms that may be variably important. The emergent mechanisms of evasive resistance include revascularization consequent to upregulation of alternative pro-angiogenic signals; protection of the tumour vasculature either by recruiting pro-angiogenic inflammatory cells or by increasing protective pericyte coverage; accentuated invasiveness of tumour cells into local tissue to co-opt normal vasculature; and increased metastatic seeding and tumour cell growth in lymph nodes and distant organs.
- Intrinsic resistance is likely to involve similar molecular and cellular mechanisms to those that mediate evasive resistance. Whereas rapid adaptive responses (fast evasion) may underlie some cases of apparent intrinsic resistance, there is evidence to suggest that certain tumours, owing to their stage of progression, treatment history, genomic constitution and/or host genotype, may have a pre-existing tumour microenvironment that conveys such indifference.
- If the postulate of evasive and intrinsic resistance to angiogenesis inhibitors is further validated in preclinical and clinical investigations, we foresee a future of cancer therapeutics in which combinatorial strategies involving angiogenesis inhibition are integrated with drugs targeting resistance mechanisms to afford enduring efficacy.

The clinical achievements with bevacizumab, sunitinib and sorafenib constitute a milestone event for the field of angiogenesis research, eliciting survival benefits in many aggressive tumours, but there is a sobering addendum: these VEGF pathway inhibitors are failing to produce enduring clinical responses in most patients^{5,22–24}. Rather, the introduction of anti-angiogenic therapy results in transitory improvements, in the form of tumour stasis or shrinkage and in some cases increased survival. Inevitably, however, the tumours begin to grow again and the disease progresses, after a fleeting period of clinical benefit that is typically measured in months²⁵. This seemingly preordained return to growth and progression in the face of ostensibly potent angiogenesis inhibitors conflicts with the widely accepted proposition that angiogenesis is an essential capability for the manifestation of lethal cancer^{2,26}. There is, therefore, a clear need to understand the mechanistic basis of this apparent conundrum for the therapeutic targeting of tumour angiogenesis.

In this Review, we elaborate a hypothesis for the transitory efficacy of the current generation of pathway-specific angiogenesis inhibitors, one we predict will prove general to potent angiogenesis inhibitors. Our proposition is based both on emerging data from clinical and preclinical investigations, and on mechanistic insights from studying the biological regulatory mechanisms operative in the tumour microenvironment that govern

tumour phenotypes, most notably angiogenesis and invasion. We envision two general modes of resistance to angiogenesis inhibitors, in particular those targeting the VEGF pathways: first, adaptive (evasive) resistance; and second, intrinsic (pre-existing) non-responsiveness (FIG. 1). Multiple mechanisms are suspected to underlie both modes of resistance, as outlined below.

Mode I: evasion of anti-angiogenic therapy

Over the past few years a provocative 'contrarian' concept has emerged from the hopeful but simplistic view that angiogenesis inhibition would prove insurmountable^{27,28}; the new postulates arose from considering the results of both preclinical and clinical research into angiogenic mechanisms and the largely modest effects that angiogenesis inhibitors have had on tumours and cancer patients. The evolving hypothesis is that angiogenic tumours can adapt to the presence of angiogenesis inhibitors, variously acquiring the means to functionally evade the therapeutic blockade of angiogenesis^{25,29–36}. In contradistinction to traditional concepts of drug resistance being acquired by mutational alteration of the gene encoding a drug target or by alterations in drug uptake and efflux^{37,38}, evasive resistance is largely indirect: alternative ways to sustain tumour growth are activated but the specific therapeutic target of the anti-angiogenic drug remains inhibited^{30,31,33,36,39–41}. The current experimental evidence, which is not yet definitive, suggests that there are at least four distinct adaptive mechanisms that manifest evasive resistance to anti-angiogenic therapies: first, activation and/or upregulation of alternative pro-angiogenic signalling pathways within the tumour (FIG. 2); second, recruitment of bone marrow-derived pro-angiogenic cells, both of which can obviate the necessity of VEGF signalling, thereby effecting reinitiation and continuance of tumour angiogenesis (FIGS 2,3); third, increased pericyte coverage of the tumour vasculature, serving to support its integrity and attenuate the necessity for VEGF-mediated survival signalling (FIG. 4); and fourth, activation and enhancement of invasion and metastasis to provide access to normal tissue vasculature without obligate neovascularization (FIG. 5).

The logical bases for these four adaptive mechanisms and the emerging experimental evidence that supports their existence are presented in the subsections below.

Upregulation of alternative pro-angiogenic signalling circuits.

Evidence for the existence of evasive resistance manifested by alternative pro-angiogenic signalling was revealed during preclinical trials in a genetically engineered mouse model of pancreatic neuroendocrine (islet cell) cancer, *Rip1-Tag2* (REF. 30). When *Rip1-Tag2* mice were treated with a monoclonal antibody (DC101) that specifically blocked VEGFR signalling (in particular VEGFR2), there was an initial response denoted by tumour stasis and reductions in tumour vascularity. However, the response phase was transitory (10–14 days) and was followed by tumour regrowth, during which the typically dense tumour vasculature was restored, indicative of re-initiation and persistence of tumour angiogenesis. The relapsing tumours were found to express higher

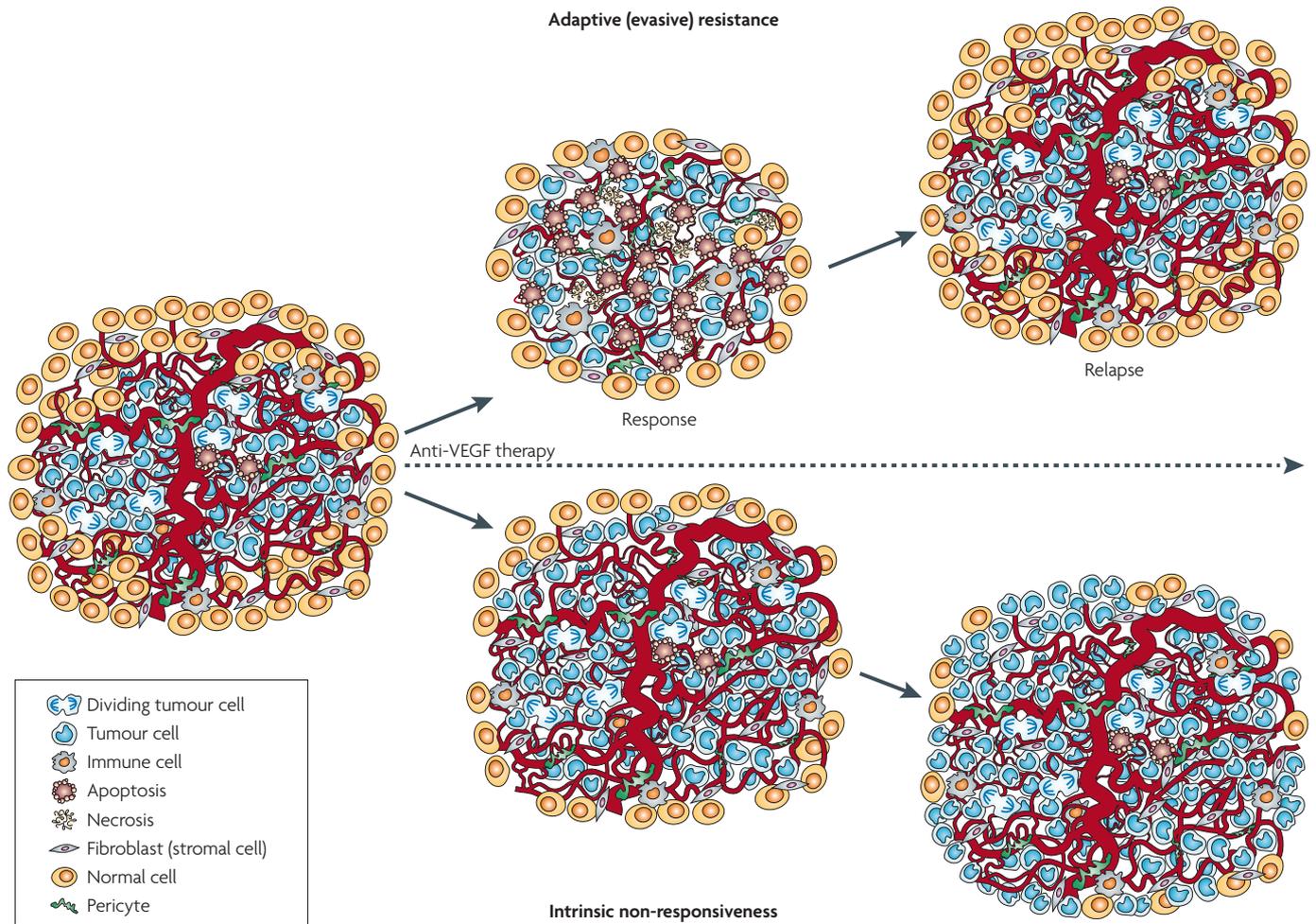


Figure 1 | Two modes of resistance in response to anti-angiogenic therapy imply adaptive evasion and intrinsic non-responsiveness of tumours. Adaptive or evasive resistance refers to the ability of a tumour, after an initial response phase, to adapt so as to evade the therapeutic blockade by inducing or accentuating mechanisms that enable neovascularization despite the therapeutic blockade, or reduce dependence on such growth of new blood vessels by other means, leading to renewed tumour growth and progression. By contrast, intrinsic non-responsiveness is a pre-existing condition defined by the absence of any (even transitory) beneficial effect of an anti-angiogenic therapy, ranging from the inability to shrink or stabilize tumours to the lack of improvement in quality of life. Consequently tumours grow and progress unabated during the course of anti-angiogenic therapy. VEGF, vascular endothelial growth factor.

levels of the mRNAs for the pro-angiogenic factors fibroblast growth factor 1 (*Fgf1*) and *Fgf2*, ephrin A1 (*Efna1*) and *Efna2*, and angiopoietin 1 (*Angpt1*) than did unperturbed tumours. Notably, the tumours had regions of acute hypoxia at the peak of the response phase, and tumour-derived cells subjected to hypoxic conditions similarly upregulated most of these genes. In order to begin assessing the functional significance of these upregulated genes in the observed revascularization, mice were first treated with the VEGFR inhibitor alone, and then at the peak of the response phase they were also treated with an FGF-trap (FGFR-Fc fusion protein) to suppress signalling through the FGF ligands. The combination attenuated the revascularization and slowed tumour growth, indicating that FGF signalling was involved in regulating the restored angiogenesis. A concurrent study showed that induction of the pro-angiogenic cytokine *IL8* served to maintain an

angiogenic capability in tumours that were otherwise impaired owing to absence of the transcription factor hypoxia-inducible factor 1 α (HIF1 α), an inducer of VEGF expression³⁶. Further corroboration comes from a recent study that implicated evasive resistance (through upregulation of pro-angiogenic factors such as VEGF, PDGFA and FGF2) in tumours whose growth is impaired by ectopic expression of the endogenous angiogenesis inhibitors *thrombospondin 1*, tumstatin and endostatin (also known as *COL4A3* and *COL18A1* fragments, respectively)⁴⁰.

A suggestion of analogous evasive resistance by FGF-dependent revascularization has come from a clinical investigation involving a study of *glioblastoma* patients being treated with the VEGFR inhibitor *cediranib* (Recentin, Astra Zeneca)⁴². There was a demonstrable response phase, and then a relapse or progression phase. Provocatively, levels of FGF2, one of the factors associated

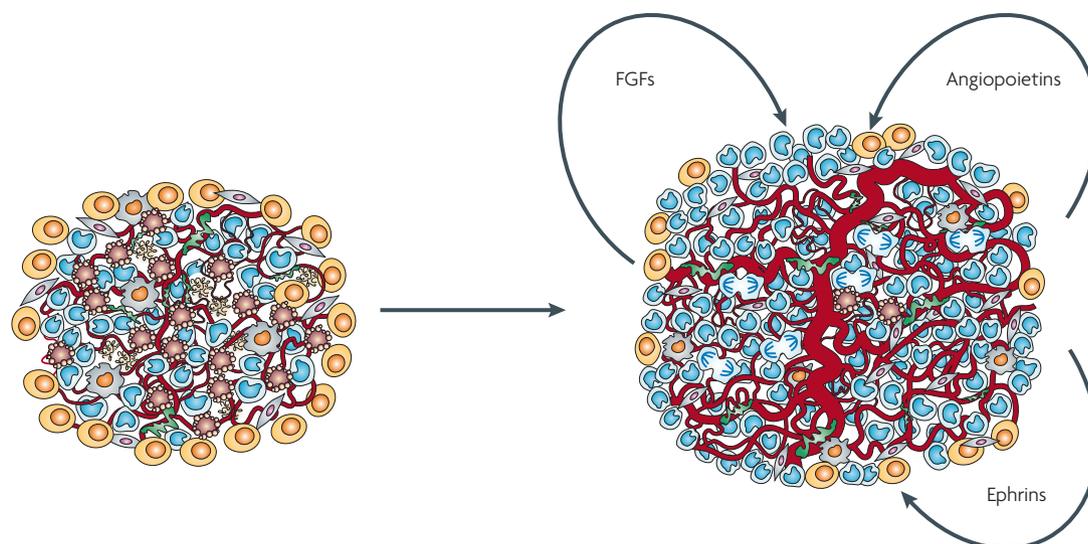


Figure 2 | Induced pro-angiogenic factor substitution re-establishes tumour neovasculature. Activation and/or upregulation of other pro-angiogenic signalling pathways, including those involving members of the fibroblast growth factor (FGF), ephrin and angiopoietin families, can circumvent the anti-angiogenic therapy and, in due course, provoke neovasculature and subsequent tumour relapse.

with evasive resistance in the preclinical models, were found to be higher in the blood of relapsing patients than in that of the same patients during the response phase, indicative of an analogous adaptive mechanism involving upregulation of FGF2. The fact that receptor tyrosine kinase inhibitors can transiently increase plasma levels of pro-angiogenic factors such as VEGF and placental growth factor (PlGF, also known as PlGF) has been previously recognized in the clinic, and even proposed as a predictive biomarker for tumour response^{43–45}. Such propositions, however, may require refinement or reconsideration in light of recent experiments that revealed increased levels of pro-angiogenic factors in the circulation of sunitinib-treated control (non-tumour-bearing) mice, implicating a systemic endocrine response to inhibition of VEGF and PDGF signalling in normal tissues⁴⁶. Increased VEGF expression was detected in a number of tissues in the sunitinib-treated mice, as were dose-dependent increases in plasma levels of granulocyte colony-stimulating factor (G-CSF, also known as CSF3), stromal cell-derived factor 1 α (SDF1 α , also known as CXCL12), stem cell factor (SCF, also known as KIT ligand) and osteopontin. Thus, the presence of pro-angiogenic factors in the blood may not be strictly indicative of tumour revascularization nor of their involvement in it. On the other hand, vascular imaging of the glioma patients treated with cediranib indicated that relapse was accompanied by re-initiation of tumour angiogenesis and loss of vascular normalization⁴², implicating a mechanism to evade the VEGF blockade. This could well involve upregulated FGF2, consistent with its increased level in the bloodstream.

The resulting uncertainties about the instructive value of increased levels of pro-angiogenic ligands in the circulation underscore the importance of analysing treated human tumour tissue to assess the angiogenic and histopathological phenotypes in responding and

relapsing tumours in the face of anti-VEGF therapy. In mouse models, the capability to do this has been instrumental in revealing the existence and importance of alternative pro-angiogenic signalling for evasive resistance. Critical evaluation of the existence (and prevalence) of this mechanism in human tumours would be markedly facilitated by increased use of trial designs involving surgical resection following a regimen of anti-angiogenic therapy, more frequent imaging (such as dynamic contrast enhanced magnetic resonance imaging (DCE-MRI) and possibly Doppler ultrasound) of tumours to assess vascular status during the course of treatment, and development of refined pre- and post-treatment biopsy strategies amenable to more routine use.

Recruitment of vascular progenitor cells and pro-angiogenic monocytes from the bone marrow. Low oxygen conditions (hypoxia) caused by vessel regression during the course of anti-angiogenic therapy can not only lead to upregulation of pro-angiogenic factors within the tumours but also to the recruitment of various bone marrow-derived cells (BMDCs) that have the capacity to fuel tumours by eliciting new blood vessels (FIG. 3). Pro-angiogenic BMDCs consist of vascular progenitors and vascular modulatory cells. Endothelial and pericyte progenitors differentiate into endothelial cells, which form the inner lining of blood vessel walls, or pericytes, the cells that envelop blood vessels, respectively^{47,72} (BOX 1). By contrast, pro-angiogenic monocytes, such as tumour-associated macrophages⁴⁸, and immature monocyte cells — including TIE2⁺ (also known as TEK⁺) monocytes⁴⁹, VEGFR1⁺ hemangiocytes^{50,51} and CD11b⁺ (also known as ITGAM⁺) myeloid cells^{52,53} — exert their functions as vascular modulators by expressing a variety of cytokines, growth factors and proteases without being physically part of the vasculature^{54,55}. Evidence that low oxygen tension triggers the recruitment of BMDCs

Doppler ultrasound
A test that uses reflected sound waves to evaluate blood flow.

Hemangiocytes
CXCR4⁺ VEGFR1⁺
haematopoietic progenitors.

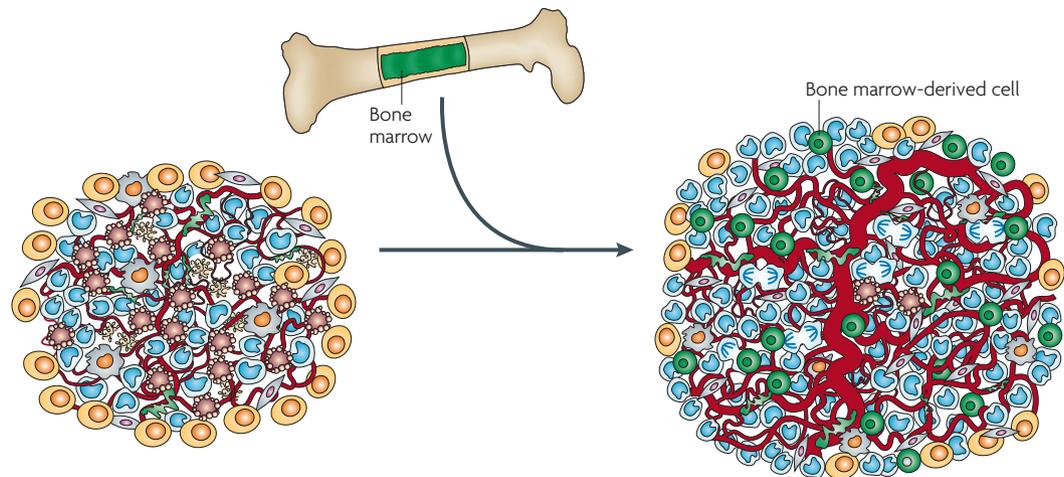


Figure 3 | Recruitment of bone marrow-derived cells can endorse restored neovascularization. Low oxygen conditions in tumours (hypoxia), acting in part through induced increases in hypoxia-inducible factor 1α and its targets stromal cell-derived factor 1α and vascular endothelial growth factor, can attract a heterogeneous population of bone marrow-derived cells consisting of vascular progenitors and pro-angiogenic monocytic cells. Endothelial and pericyte progenitors are incorporated as components of new vessels to directly build new blood vessels, and pro-angiogenic monocytes fuel the tumours with pro-angiogenic cytokines, growth factors and proteases, all of which facilitate neovascularization.

stems from observations in experimentally induced ischaemic tissue, wherein endothelial progenitors and other $CXCR4^+$ BMDCs are recruited, in part through increases in HIF 1α and its downstream effectors SDF 1α (a CXCR4 ligand) and VEGF itself^{18,56,57}. In a study of neovascularization in glioblastoma multiforme (GBM), a tumour type characterized by extensive hypoxia and necrosis, HIF 1α was found to promote angiogenesis and tumour growth by inducing recruitment of various pro-angiogenic bone marrow-derived CD45⁺ (also known as *PTPRC*⁺) myeloid cells, including subpopulations defined by expression of TIE2, VEGFR1 and/or CD11b, as well as mature F4/80⁺ tumour-associated macrophages; endothelial and pericyte progenitor cells were also prevalent^{55,58}. GBM tumours lacking HIF 1α had few of these abundant BMDCs, and were severely impaired in their angiogenic and tumour growth phenotypes⁵⁵.

These results collectively suggest that anti-angiogenesis-induced hypoxia can in some contexts elicit BMDC recruitment and thereby foster an adaptive mechanism that enables tumours to overcome hypoxia. Given that many tumours contain hypoxic areas, influx of BMDCs may depend on a threshold of or correlate with the degree of low oxygen tension⁵⁹. Evidence supporting the link between therapy-induced hypoxia and BMDC recruitment has come from a study demonstrating that vasculature-disrupting agents, which ablate blood vessels within a tumour and thereby cause acute hypoxia and necrosis, can trigger a transient accumulation of endothelial progenitor cells at the tumour margins in sufficient numbers to facilitate revascularization⁶⁰. Notably, the untreated transplant tumours did not contain appreciable levels of BMDCs, consistent with the interpretation that this was an adaptive response enabling evasive resistance to potent anti-angiogenesis therapy.

There is intriguing evidence from clinical investigations that this evasion mechanism operates in patients with GBM who are undergoing anti-VEGF therapy. One recent study suggests that hypoxia determines survival outcome in bevacizumab-treated GBM patients⁶¹, and a second reports that SDF 1α , a hypoxia-regulated retention factor for CXCR4⁺ BMDCs, was detectable in the blood of patients with GBM at the time of tumour progression and relapse (discussed above and in REF.42). Therefore, SDF 1α could be an effector and biomarker of response for relapsing tumours, raising the possibility that both forms of evasive resistance (upregulation of pro-angiogenic factors such as FGF2 and recruitment of BMDCs) could be operative and collaborative in GBM. However, there remains the challenge to develop means to distinguish systemic paracrine reactions of drug-treated normal tissue from those of tumour-specific adaptive responses. We predict that such mechanisms will prove to be involved in other tumour types, and anticipate that both these clinical investigations and the aforementioned preclinical studies will motivate further investigation of the possibility.

Increased and tight pericyte coverage protects tumour blood vessels. A growing body of evidence indicates that pericytes, the periendothelial support cells of the microvasculature^{62–64}, are also important cell constituents of the aberrant tumour vasculature. When therapies impair neovascularization and/or elicit vascular regression, some tumours evidently rely on pericytes to help keep a core of pre-existing blood vessels alive and functional (FIG. 3). This concept has evolved from the observation by several groups that, although inhibition of VEGF signalling can lead to substantial reduction in tumour vascularity, distinctive functional vessels remain that are slim and tightly covered with pericytes^{19,65–68}.

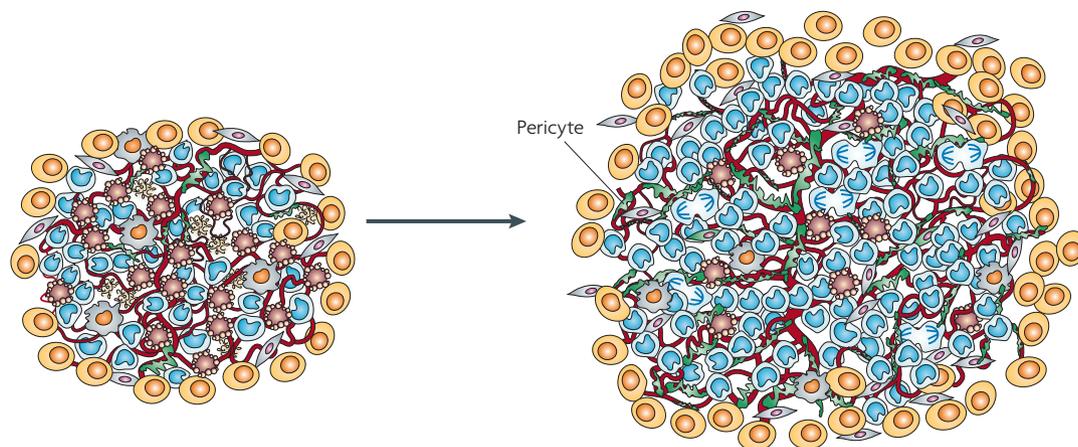


Figure 4 | Increased pericyte coverage protects tumour blood vessels. Although inhibition of vascular endothelial growth factor signalling pathways can lead to vessel regression, a few ‘normal-appearing’ slim and functional vessels remain; these vessels are densely and tightly covered with pericytes, and are markedly distinct from the vessels that are seen in tumours of untreated animals, which are typically dilated, tortuous and irregularly shaped, and variably covered with less closely associated pericytes. Such coating by pericytes arguably helps the tumour endothelium to survive and function, and thereby enables tumours to grow (perhaps more slowly) during the course of an anti-angiogenic therapeutic regimen.

Notably, the morphology of these surviving tumour vessels is readily distinguishable from the typically dilated tumour vessels of untreated animals, which are, by contrast, variably covered with less closely associated pericytes^{19,66,67,69}. These observations suggest that endothelial cells can induce pericyte recruitment to protect themselves from death consequent to the lack of the crucial tumour-derived survival signals conveyed by VEGF. This hypothesis is supported by the findings that tumour vessels lacking adequate pericyte coverage are more vulnerable to VEGF inhibition^{19,70} and that tumour pericytes, which are juxtaposed to endothelial cells, express appreciable levels of VEGF and potentially other factors that support endothelial cell survival^{71,72}. In addition, pericytes are capable of attenuating the proliferation rate of endothelial cells^{73,74}, a necessity for proper maturation and stabilization of newly formed vascular structures. One can therefore speculate that in treated tumours, blood vessels heavily covered with pericytes survive because pericytes mediate endothelial cell quiescence and survival (this is also the case for normal tissue), therefore rendering tumour endothelial cells less responsive to anti-angiogenic agents.

Based on the stabilizing effects of pericytes and their functional capability to ameliorate the effects of VEGFA depletion on angiogenic endothelium, there is a rationale for targeting both cell types. Indeed, improved efficacy has been observed in a variety of mouse tumour models, beginning with the *Rip1-Tag2* transgenic mouse model of pancreatic neuroendocrine cancer in studies combining inhibitors of VEGF and PDGF signalling to target endothelial cells and pericytes, respectively^{19,20,75,76}. Clinical trials are currently ongoing or in development that aim to target endothelial cells and pericytes simultaneously and assess the potential benefits for anti-tumoural efficacy. A related agenda will be to clarify

the clinical situations in which singular VEGF pathway inhibition (with its attendant transitory normalization) is preferable, and those in which combining VEGF inhibitors with agents that disturb vascular ‘normalization’ by disrupting pericyte support (such as PDGFR inhibitors) has greater benefit.

The strategy of dissociating pericytes from the tumour vasculature (with inhibitors of PDGFR signalling, for example) to render the tumour endothelium more sensitive to VEGF inhibitors is attractive, but emerging clues suggest an additional and undesirable effect: severe reduction or lack of pericyte coverage may disrupt the integrity of the vasculature, enabling tumour cells to transit into the circulatory system, thereby facilitating metastasis. In one study, genetic disruptions of pericyte coverage elicited increased metastasis in the *Rip1-Tag2* pancreatic islet tumour model⁷⁷. In several experimental therapeutic studies, both anti-VEGF strategies, as well as treatment modalities involving dual targeting of endothelial cells and pericytes, increased the incidence of metastasis concomitant with severe impairment of the vasculature and growth of the primary tumour (O. Casanovas, unpublished observations; R. Kerbel, unpublished observations). If substantiated by further studies in mouse models and in human patients, the result would solidify another clinically significant mechanism of evasive resistance: when angiogenesis is severely inhibited and the integrity of the remaining tumour vasculature disrupted by anti-angiogenic therapeutic strategies, increased intravasation might provide an alternative means for tumour cells to survive and grow, through blood-borne metastasis. However, resistance mechanisms involving increased intravasation and metastatic seeding will probably remain dependent either on evading angiogenesis inhibition at the new sites, or on co-opting normal tissue vasculature to support tumour expansion, as discussed below.

Intravasation

Part of the metastasis process in which cancer cells invade through the basal membrane into blood vessels.

Co-option

Blood vessel co-option: tumour cells grow around existing blood vessels in the tissue.

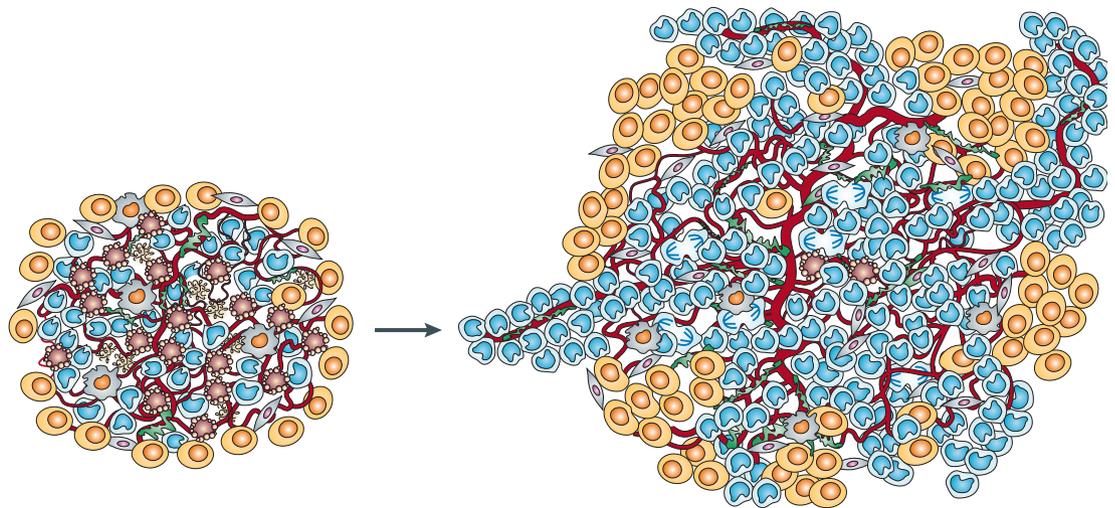


Figure 5 | Increased tumour cell invasiveness to escape oxygen and nutrient deprivation. When tumours are not able to satisfactorily reinitiate angiogenesis, tumour cells may invade adjacent normal tissue to achieve vascular sufficiency in a dispersed fashion. Tumour cells can migrate along the outside of blood vessels (perivascular invasion), using them as conduits into normal tissue, or infiltrate through the extracellular matrix. In either case they depend on normal vasculature for sustenance.

Increased capabilities for invasion without angiogenesis.

There are increasing clues for another insidious form of adaptation: change to a condition of increased invasiveness. It is increasingly evident that in some tumours in which angiogenesis is thwarted genetically or pharmacologically, cancer cells adapt by migrating more aggressively into normal tissue. The evasive phenotype of increased invasiveness was first described in mouse models of orthotopic GBM in which neovascularization was blocked by genetically deleting angiogenic factors such as VEGF, HIF1 α and matrix metalloproteinase 9 (MMP9), or inhibited pharmacologically with the VEGF inhibitor SU5416 (*semaxanib*). In this setting the tumours became more invasive and continued to grow, albeit more slowly^{31,39,55}. The glioblastoma cells were seen to co-opt normal blood vessels, evidently using them as highways or conduits to journey deep into the brain, thereby achieving vascular sufficiency in a dispersed fashion, a phenotype referred to as perivascular tumour invasion. Congruent with the results obtained in orthotopic mouse models of GBM, three recent studies implicate pro-invasive adaptation in humans, as observed by MRI in a subset of GBM patients who had developed multifocal recurrence of tumours during the course of anti-VEGF therapy with bevacizumab^{78,79} (M. Prados and N. Butowski, unpublished observations).

A pro-invasive adaptation mechanism has also been observed in the *Rip1-Tag2* pancreatic islet tumour model in the course of assessing the effects of inhibitors targeting VEGFR³⁰ or multiple receptor tyrosine kinases implicated in pro-angiogenic signalling (D.H., unpublished observations, and O. Casanovas, unpublished observations). The reduction in tumour vascularity evoked by the angiogenesis inhibitors was accompanied by clear signs of increased tumour cell invasiveness. How might such tumours become more invasive? We envision multiple adaptive mechanisms. First, tumours may increase

the activity of a pre-existing invasion programme that was not previously the driving force of expansive tumour growth, given the capability for angiogenesis. Alternatively, some tumours may switch on a distinctive invasive growth programme to that arising spontaneously during progression. For example, untreated GBMs often invade normal brain tissue by infiltrating as single cells into the brain parenchyma, migrating along basement membranes of ventricles, leptomeninges and blood vessels⁸⁰. By contrast, when GBMs were genetically or pharmacologically impaired in their angiogenic capability, tumour cells were observed to predominantly migrate as multicellular layers along normal blood vessels^{31,39,55,81}. Additionally, VEGF signalling may in some contexts serve to directly restrict perivascular tumour cell invasion: addition of VEGF to cultured GBM cells, or the generation of GBM cells that ectopically express high levels of VEGF, reduced the propensity of those cells to invade both *in vitro* and *in vivo*. Notably, these tumour cells express VEGFR1 and VEGFR2 (REF. 55). Adaptive mechanisms in the face of angiogenesis inhibition involving increased invasiveness are also predicted to culminate in increased metastasis in some tumour types, and indeed current studies with VEGF pathway inhibitors are uncovering both increased invasiveness and metastasis, as discussed above. Importantly, the existence and the prevalence of such adaptive mechanisms across the spectrum of human cancers remain to be established, but there are sufficient clues and rationale to raise this possibility here in order to provoke future investigations.

We have presented a set of non-traditional (evasive) therapeutic resistance mechanisms that stand in contrast to the well-established mechanisms operative in cancer cells, where genes are mutationally altered, deleted and amplified so as to afford genetic resistance to drugs targeting the genetically unstable tumour cells. We cannot

Leptomeninges

The two innermost layers, comprised of the arachnoid mater and pia mater, that envelop the brain and spinal cord.

exclude the possibility that such genetic resistance mechanisms will be discovered in tumour endothelial cells in response to chronic therapy with anti-angiogenic drugs, perhaps in particular metronomic chemotherapy. Indeed, some tumour endothelial cells have been reported to be aneuploid⁸², suggesting the capability to develop mutations that convey drug resistance. Additionally, it is increasingly evident that angiogenic signalling pathways are not unilaterally specific, and indeed are found to be functioning in certain tumour types, as reviewed by L. Ellis and D. J. Hicklin in this Focus issue¹¹ and in REF. 83, which raises the possibility of classical resistance circumventing the direct impairment of tumour cells treated with ‘angiogenesis inhibitors’. We suspect, however, that the aforementioned evasive resistance mechanisms will prove to be the primary basis for the transitory efficacy of anti-angiogenic drugs, in particular those targeting VEGF signalling.

Mode 2: indifference to anti-angiogenic therapy

We envision a second mode of resistance to anti-angiogenic therapy, one for which the experimental clues are more diffuse but the rationale is nevertheless persuasive: intrinsic, pre-existing non-responsiveness of a tumour (FIG. 1). One such clue involves the clinical trials of the currently approved VEGF pathway inhibitors. A substantial minority of the individuals tested in clinical trials for bevacizumab, sorafenib and sunitinib failed to show even transitory clinical benefit^{24,42}. Although the trial designs did not typically involve frequent serial monitoring that could document intrinsic resistance, it seems likely that some of these non-responding patients fall into this category of no discernible response, with their tumours evidently ‘growing through’ the therapy. Although this resistance could reflect rapid adaptation and the onset of evasive resistance, we expect that pre-existing resistance will prove to be in force in some cases. Its basis could be in tumours that, by virtue of their particular developmental ontogeny, have already activated one or more of the aforementioned evasive resistance mechanisms, not in response to therapy but rather in response to the selective pressures of their microenvironment during premalignant development and malignant progression.

The operative definition of this mode of pre-existing resistance is the absence of a discernable (even transitory) beneficial effect of an angiogenesis inhibitor, even when the subject’s tumour(s) is serially analysed. There is no tumour shrinkage (that is, no partial or complete responses in the clinical parlance), no cessation of tumour growth (stasis), nor even retardation in growth rate; neither is there quality of life benefit, nor increased survival. In short, the patient (or the tumour-bearing animal) is intrinsically refractory to the anti-angiogenic therapy, such that disease progression continues unabated. Rigorous characterization and discrimination of this mode of non-responsiveness as distinctive to the potentially rapid adaptive mechanisms discussed above is experimentally challenging because the major difference may be in the timing of activating the two modes, as variations of the same basic mechanistic themes. Nonetheless, there is good reason to suspect that intrinsic resistance exists in some individuals afflicted with cancer, appreciation of which may prove important for clinical management of anti-angiogenic therapies. The clues and rationales for these prospective mechanisms of pre-existing resistance are discussed briefly below.

Pre-existing multiplicity of redundant pro-angiogenic signals.

There are intriguing clues of pre-existing resistance to angiogenesis inhibition consequent to treatment with bevacizumab, which captures the VEGF ligands. Bevacizumab is currently approved by the US Food and Drugs Administration (FDA) for treatment of late-stage metastatic colon, breast and lung cancer, but only in combination with conventional chemotherapy for all three indications (see [FDA approval of bevacizumab](#) URL in Further information). In the case of breast cancer, an analysis of human breast cancer biopsies covering a spectrum from low- to high-grade malignancies revealed that late-stage breast cancers expressed a plethora of pro-angiogenic factors, including FGF2, in contrast to earlier stage lesions, which preferentially expressed VEGF⁸⁴. Thus one can imagine that the pre-existence of FGF2 and the other pro-angiogenic factors in late-stage metastatic tumours could enable continuing angiogenesis in the face of bevacizumab therapy, such that inhibition of VEGF signalling does not affect angiogenesis.

Box 1 | Pericytes

Pericytes (mural or vascular smooth muscle cells) are perivascular cells with a prominent nucleus, a small content of cytoplasm and long processes that wrap around blood capillaries (peri- meaning around; -cyte meaning cell). They communicate with endothelial cells by direct physical contact through a jointly synthesized basement membrane and reciprocal paracrine signalling. Pericytes and endothelial cells are thereby interdependent and, as such, defects in either endothelial cells or pericytes can affect the vascular system. Pericytes have various demonstrable functions in different physiological contexts, including stabilization and homeostatic regulation of mature blood vessels; facilitation of vessel maturation in the context of neovascularization; provision by their intimate association of endothelial cell survival signals; and limitation of cell transit across the vascular wall. The functional significance of pericytes in development is underscored by genetic depletion or disruption of pericyte association with the developing vasculature in mice, which results in blood vessel dilation, widespread microvascular leakage and microaneurysms, and subsequent lethality during late gestation. In tumours, pericytes are typically less abundant and more loosely attached to blood vessels than in normal tissues, but their association is still important, as shown in a growing body of experimental evidence which indicates that pericytes help to maintain the integrity and functionality of the tumour vasculature.

Metronomic chemotherapy
Administration of chemotherapeutic drugs at comparatively low doses at frequent or continuous schedule, with no extended interruptions.

Notably, bevacizumab trials in late-stage metastatic breast cancer have been equivocal, as reviewed by L. Ellis and D. J. Hicklin in this Focus issue¹¹ and in REF. 85. The exception involved a small subgroup of patients with metastatic ERBB2⁻ (also known as HER2⁻) breast cancer who had not received prior chemotherapy. Under these particular conditions, a regimen of bevacizumab in combination with the cytotoxic drug paclitaxel extended progression-free survival (but not overall survival)⁸⁶. By contrast, trials in patients pretreated with chemotherapy showed no benefit from bevacizumab as second-line therapy, alone or in combination with additional chemotherapy (see [New approval for bevacizumab \(Avastin\)](#) URL in Further information). The latter result suggests that the pretreated tumours may have already activated mechanisms that coincidentally convey intrinsic resistance to subsequent anti-angiogenic therapy. What, then, could be the basis for the modest albeit demonstrable value of bevacizumab when combined in a first-line setting with chemotherapy, which led to its approval by the FDA? Certainly bevacizumab might be transiently blocking angiogenesis in these tumours. Another possible explanation is that VEGF blockade in the context of these late-stage breast cancers is serving primarily to transiently normalize the permeability of tumour vasculature, reducing VEGF-induced haemorrhaging and oedema and therefore relieving the physiological effects of tumour burden. In addition, one cannot exclude the possibility that the induction of vascular normalization by VEGF pathway inhibitors serves to improve blood flow, consequently facilitating intratumoural bioavailability of efficacious levels of the attendant cytotoxic chemotherapy⁶⁶. This latter effect could also explain the selective clinical benefits of anti-VEGF therapy only in combination with chemotherapy in metastatic colon and lung cancer. This view of VEGF inhibitors as 'chemosensitizers' does not imply that *bona fide* anti-angiogenic therapy has no potential benefit in treating such late-stage cancers, but rather implies that we must find ways to finesse pre-existing resistance by targeting the alternative pro-angiogenic signals in addition to the vascular normalization (and contributions to angiogenesis inhibition) afforded by VEGF inhibition. Here again, there is clear need to histologically analyse treated human tumours in response, relapse and progression phases (in this case with bevacizumab plus chemotherapy), to ascertain the temporal ontogeny of effects on the tumour vasculature (for example, vascular dropout, revascularization and normalization).

Pre-existing inflammatory cell-mediated vascular protection. A recent preclinical study found that a subset of murine transplant tumours growing in mice showed no responsiveness to an anti-mouse VEGF monoclonal antibody that mimicked bevacizumab³⁴. The non-responsive tumours, which had not previously been treated with chemotherapy, were characterized by a pre-existing infiltration of inflammatory cells, principally CD11b⁺Gr1⁺ myeloid cells, which were shown to express a number of pro-angiogenic factors. By contrast, the responsive tumour types had comparatively low levels of such inflammatory cells. Pharmacological impairment of myeloid cell

recruitment rendered the otherwise resistant tumours responsive to the VEGF blockade. Although the experimental design and data cannot discriminate between rapid adaptation and pre-existing resistance, there is reason to suspect that these pro-angiogenic myeloid cells, when sufficiently abundant as a consequence of chemoattractants produced by such tumours, will convey pre-existing resistance. Certainly this result is not definitive of the possible general principle, but it motivates both preclinical and clinical studies to further assess this prospective mechanism.

Characteristic hypovascularity and indifference toward angiogenesis inhibitors. Pancreatic ductal adenocarcinoma (PDAC) may display another class of pre-existing resistance, manifest as a tumour type that is characteristically hypovascularized with a massive (largely avascular) desmoplastic stroma⁸⁷. The fact that treatment-naïve PDAC tumours are intrinsically poorly vascularized and nevertheless not widely necrotic suggests that PDAC tumorigenesis involves adaptation to survive and prosper in the harsh, presumably hypoxic microenvironment that results from the sparse vascularity. It is unclear why these tumours fail to induce a dense neovasculature and instead evolve to flourish in the absence of prominent angiogenesis. Irrespectively, the PDAC tumour physiology may render it intrinsically indifferent to angiogenesis inhibitors. For example, ~75% of PDACs carry inactivating mutations in the *TP53* tumour suppressor gene^{88,89}, loss of which in another tumour type has been shown to improve survival in hypoxic conditions, including ones induced by angiogenesis inhibition^{90,91}. Consistent with this line of reasoning, it is notable that bevacizumab showed no efficacy in a clinical trial of patients with late-stage PDAC, alone or in combination with the standard-of-care chemotherapeutic agent gemcitabine²⁴. One can infer, therefore, that normalization is either not induced or lacks the ability to augment the limited utility of gemcitabine. Preliminary studies in a mouse model of PDAC with another VEGF pathway inhibitor also showed no efficacy alone or combined with gemcitabine and no alterations in vascularity, consistent with this clinical observation (P. Olson and D.H., unpublished observations). Although this class of intrinsic indifference to VEGF and other angiogenesis inhibitors may be applicable to some or all cases of PDAC and perhaps to rare subtypes of other organ-specific cancers, appreciation of its potential existence may provoke further investigation and, if validated, will probably prove influential for evaluating potential therapeutic trials with angiogenesis inhibitors in PDAC and other similar tumour types.

Invasive (and metastatic) co-option of normal vessels without requisite angiogenesis. Extrapolation from the observations of adaptive resistance by increased invasion and metastasis predicts the intrinsic resistance of tumours that have already switched on highly invasive and/or metastatic capabilities early in their ontogeny, serving to effect local or distant co-option of quiescent normal tissue vessels so as to sustain growth without requisite angiogenic switching and development of neovasculature. Grade II and III astrocytoma may be an example. These diffusely

Oedema

An observable swelling due to an increase of interstitial fluid in any organ.

Desmoplastic stroma

Abnormal and excessive growth of stromal cells that is often associated with invasive cancers.

growing tumours do not form lesions with a prominent neovasculature containing proliferative endothelial cells^{80,92}. We suspect that such tumours will prove refractory to angiogenesis inhibitors and even to inducers of normalization such as bevacizumab, given that they have not acquired the aberrant, angiogenic vasculature that is typical of the more aggressive grade IV lesions (GBM).

Although the rationales posed above for the existence of different forms of pre-existing (intrinsic) resistance to anti-angiogenic therapy may seem speculative based on the current data, we are confident in predicting that intrinsic resistance will prove to be operative in some tumours, with consequent implications for anti-angiogenic therapies. Certainly, the observation that patients bearing the same tumour type respond differently to the same therapy is well recognized. Such a heterogeneous response to anti-VEGF therapy is exemplified in the aforementioned report of a clinical trial of the VEGFR inhibitor cediranib, in which a subset of patients with GBM responded transiently, whereas another group of patients had essentially no response⁴². It remains to be elucidated whether the intrinsic indifference to anti-angiogenic therapy in the latter group is due to one or more of the described intrinsic resistance mechanisms. The results underscore the importance of biomarkers that would predict whether a patient will respond to anti-VEGF therapy. We raise the hypothesis herein to set seeds of future inquiry that should, in due course, establish (or not) its accuracy and prevalence, and clarify the alternative mechanisms that most commonly manifest this proposed mode of intrinsic resistance or indifference to angiogenesis inhibitors.

Conclusions

These considerations, about modes and underlying mechanisms of resistance to anti-angiogenic therapy, present both an agenda for future research and the outlines of a strategy for improving the treatment of human cancer with angiogenesis inhibitors. All of the mechanisms, both of adaptive and of intrinsic resistance, deserve further investigation and rigorous evaluation of their prevalence and significance, both in animal models of cancer and in humans. We also expect that other evasive mechanisms will be uncovered. Additionally, it will be important to clarify the circumstances that elicit particular mechanistic pathways of resistance, alone and in the context of standard-of-care chemotherapy and radiotherapy for different

cancer types, recognizing that the likely future of most anti-angiogenic therapies will be in such combinations. The constraints on clinical investigation pose challenges indeed, but we envision that advancing technologies, in non-invasive imaging for example, and in collecting and analysing tumour biopsies to assess histological parameters of response and evasive resistance, will be instrumental. Another challenge is to identify robust biomarkers for tumour angiogenesis and for vascular normalization⁹³, as well as for the modes of intrinsic and evasive resistance.

As for cancer therapeutics, we foresee several applications of these principles. First and foremost is the emerging role of invasion and metastasis, both in intrinsic and adaptive resistance. The combination of anti-invasive and anti-metastatic drugs with anti-angiogenic drugs would seem to have particular promise. For example, multiple drugs targeting the pro-invasive hepatocyte growth factor (HGF)–MET and insulin-like growth factor 1 (IGF1) receptor pathways are in clinical trials^{94,95} and should be tested in combination with VEGF pathway inhibitors. The HIF regulatory network also holds promise as a target, given its global effects on angiogenesis, invasion and stress-adaptive cell physiology^{96–99}. Another promising avenue involves multi-targeting of parallel pro-angiogenic signalling circuits, aimed to circumvent pre-existing redundancy or evasive mechanisms involving upregulation of alternative angiogenic signalling circuits. Drugs targeting such signals (such as FGFs) are entering clinical trials, now aimed in part to assess their impact on evasive resistance to VEGF blockade^{100–102}.

Finally, although the transitory efficacy of the groundbreaking VEGF pathway inhibitors might be construed as disappointing, the results must be considered in the context of the current standards of care for most of the major human cancers, which typically have transitory efficacy, inevitable progression and/or resistance, toxicity and poor quality of life effects. Angiogenesis inhibitors, despite their evident limitations, are an important milestone in cancer therapeutics, where they are becoming components of standard-of-care therapy, for example for colorectal and renal cancers. Moreover, the growing knowledge about their effects and efficacy, and about the existence and mechanistic basis for adaptive evasive resistance and abject failures (intrinsic indifference), presents an exciting future of opportunity for improving and sustaining the benefits of anti-angiogenic therapy.

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DATABASES

Entrez Gene:

<http://www.ncbi.nlm.nih.gov/entrez/query.fcgi?db=gene>
[Angpt1](#) | [COL18A1](#) | [COL4A3](#) | [CSF3](#) | [CXCL12](#) | [CXCR4](#) | [EfnA1](#) | [EfnA2](#) | [ERBB2](#) | [Egfr1](#) | [Egfr2](#) | [HGF](#) | [HIF1a](#) | [IGF1](#) | [IL8](#) | [ITGAM](#) | [KDR](#) | [KIT ligand](#) | [MET](#) | [MMP9](#) | [osteopontin](#) | [PDGFA](#) | [PGF](#) | [PTPRC](#) | [TEK](#) | [TP53](#)

National Cancer Institute: <http://www.cancer.gov/>
[breast cancer](#) | [colon cancer](#) | [glioblastoma](#) | [lung cancer](#) | [non-small-cell lung cancer](#) | [renal cancer](#)

National Cancer Institute Drug Dictionary:
<http://www.cancer.gov/drugdictionary/>
[bevacizumab](#) | [gemcitabine](#) | [paclitaxel](#) | [semaxanib](#) | [sorafenib](#) | [sunitinib](#)

FURTHER INFORMATION

G. Bergers' homepages: <http://www.ucsf.edu/bms/faculty/bergers.html>; http://neurosurgery.medschool.ucsf.edu/neurosurgery_research/btrc/bergers_lab.html

Angiogenesis Inhibitors Therapy: <http://www.cancer.gov/cancertopics/factsheet/Therapy/angiogenesis-inhibitors>

Clinical trials in patients with brain cancer:
<http://www.cancer.gov/search/ResultsClinicalTrialsAdvanced.aspx?protocolsearchid=4816487>

Clinical trials in patients with breast cancer:
<http://www.cancer.gov/search/ResultsClinicalTrials.aspx?protocolsearchid=4586362>

Clinical trials in patients with gastrointestinal stromal tumours: <http://www.cancer.gov/search/ResultsClinicalTrials.aspx?protocolsearchid=4586407>

Clinical trials in patients with kidney cancer:
<http://www.cancer.gov/search/ResultsClinicalTrials.aspx?protocolsearchid=4586416>

Clinical trials in patients with liver cancer:
<http://www.cancer.gov/search/ResultsClinicalTrials.aspx?protocolsearchid=4586421>

Clinical trials in patients with lymphoma:
<http://www.cancer.gov/search/ResultsClinicalTrials.aspx?protocolsearchid=4586451>

Clinical trials in patients with lung cancer:
<http://www.cancer.gov/search/ResultsClinicalTrials.aspx?protocolsearchid=4586453>

Clinical trials in patients with pancreatic cancer:
<http://www.cancer.gov/search/ResultsClinicalTrials.aspx?protocolsearchid=4586434>

FDA approval of bevacizumab: <http://www.cancer.gov/cancertopics/druginfo/fda-bevacizumab>

New approval for bevacizumab (Avastin): <http://www.fda.gov/cder/Offices/OODP/whatsnew/bevacizumab200802.htm>

ALL LINKS ARE ACTIVE IN THE ONLINE PDF