



Published in final edited form as:

*Adv Exp Med Biol.* 2019 ; 1147: 65–91. doi:10.1007/978-3-030-16908-4\_2.

## Pericytes in Glioblastomas: Multifaceted Role Within Tumor Microenvironments and Potential for Therapeutic Interventions

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### Abstract

Glioblastoma (GBM) is an aggressive and lethal disease that often results in a poor prognosis. Unlike most solid tumors, GBM is characterized by diffuse infiltrating margins, extensive angiogenesis, hypoxia, necrosis, and clonal heterogeneity. Recurrent disease is an unavoidable consequence for many patients as standard treatment options such as surgery, radiotherapy, and chemotherapy have proven to be insufficient in causing long-term survival benefits. Systemic delivery of promising drugs is hindered due to the blood-brain barrier and non-uniform perfusion within GBM tissue. In recent years, many investigations have highlighted the role of GBM stem cells (GSCs) and their microenvironment in the initiation and maintenance of tumor tissue. Preclinical and early clinical studies to target GSCs and microenvironmental components are currently underway. Of these strategies, immunotherapy using checkpoint inhibitors and redirected cytotoxic T cells have shown promising results in early investigations. But, GBM microenvironment is heterogenous and recent investigations have shown cell populations within this microenvironment to be plastic. These studies underline the importance of identifying the role of and targeting multiple cell populations within the GBM microenvironment which could have a synergistic effect when combined with novel therapies. Pericytes are multipotent perivascular cells that play a vital role within the GBM microenvironment by assisting in tumor initiation, survival, and progression. Due to their role in regulating the blood-brain barrier permeability, promoting angiogenesis, tumor growth, clearing extracellular matrix for infiltrating GBM cells and in helping GBM cells evade immune surveillance, pericytes could be ideal therapeutic targets for stymieing or exploiting their role within the GBM microenvironment. This chapter will introduce hallmarks of GBM and elaborate on the contributions of pericytes to these hallmarks by examining recent findings. In addition, the chapter also highlights the therapeutic value of targeting pericytes, while discussing conventional and novel GBM therapies and obstacles to their efficacy.

### Keywords

Glioblastoma; GBM Pericytes; Microenvironment

## Introduction

### Glioblastoma: Incidence and Histological Characteristics

GBM is the most common and aggressive primary malignant brain cancer with a dismal prognosis (Sattiraju et al. 2017a; Van Meir et al. 2010). The median overall survival rate is currently 20.8 months for patients undergoing treatment with adjuvant chemotherapy (temozolomide) following maximum safe resection (Roh et al. 2017). It is estimated that about 13,000 patients are newly diagnosed with GBM annually with an estimated 5-year survival rate of about 10% for adults and about 40% for children (Rostomily et al. 2005; Song et al. 2010; Stupp et al. 2009).

Histopathological characteristics of GBM include anaplasia, macrophage and microglial infiltration, extensive angiogenesis and regions of severe hypoxia, resulting in pseudopalisading structures around necrosing neural tissue (Bissell and Radisky 2001; Van Meir et al. 2010). At a cellular level, GBM is characterized by the rapid proliferation of malignant cells which invade diffusely into the surrounding normal brain parenchyma accompanied by extensive proliferation of endothelial cells which assemble into highly torturous, disorganized and leaky blood vessels which often result in hypoxic microenvironments within tumor tissues (Van Meir et al. 2010; Zong et al. 2012). Invading GBM cells tend to displace preexisting astrocytes and pericytes that are otherwise tightly wrapped around endothelial cells, thereby disrupting the blood-brain barrier (BBB), resulting in leaky blood vessels (Dubois et al. 2014; Watkins et al. 2014). The cell of origin (COI) which is hypothesized to be the first malignant cell that sets the eventual formation of aberrant tissue into motion is thought to exist within the tumor mass in areas of severe hypoxia. GBM is a highly heterogenous disease indicated by the high degree of genetic variations and existence of multiple subclones within tumor tissues. This high degree of heterogeneity has also prompted some researchers to believe that there may be multiple cells of origin, giving rise to a mosaic of aberrant cells which we consider as the tumor tissue (Bradshaw et al. 2016; Cabrera et al. 2015; Dalerba et al. 2007; Fidoamore et al. 2016; Friedmann-Morvinski and Verma 2014; Heddleston et al. 2011).

### Hypoxic Microenvironment Within GBM

GBM is an aggressive disease which requires abundant supply of oxygen and nutrients for the survival of highly proliferating cells. The high rate of cellular proliferation within the tumor tissue causes severe hypoxia (0.1–0.5%) within the tumor core and mild hypoxia (0.5–2.5%) in peripheral regions of the tumor (Evans et al. 2004). Within these hypoxic regions, cells residing far away from preexisting blood vessels that cannot adapt to their hypoxic microenvironment undergo apoptosis or coagulative necrosis (Brat and Van Meir 2004; Martínez-González et al. 2012). This results in the outward movement of cells in the surrounding areas towards the periphery of the hypoxic region. These outward moving cells form palisading structures called “Pseudopalisades” and these events are thought to enhance the invasiveness of GBM cells, which in turn stimulate endothelial proliferation and angiogenesis through secretion of VEGF and other factors. Pseudopalisades are a distinct feature of GBMs and a marker of aggressive disease, often distinguishing them from low-

grade astrocytomas (Brat et al. 2004; Heddleston et al. 2009; Rong et al. 2006; Sattiraju et al. 2017a).

## Hallmarks of GBM

### Glioblastoma Initiation and Maintenance

Recent investigations have pointed towards progenitor cells present within the subventricular zone (SVZ) and in peripheral white matter of the cortex as potential COIs for GBM (Alcantara Llaguno et al. 2015). These investigations had employed cell population-specific knockdown of proto-oncogenes and in vivo transduction of progenitor cells expressing fluorescent labels in the cortex of the brain of transgenic mice to show that primitive progenitor cells give rise to aberrant populations of rapidly proliferating cells in a hierarchical fashion. *Llaguno et al.* pointed to the mutation of proto-oncogenes as a prerequisite for the transformation of otherwise normal progenitor cells and subsequent altered migration and production of aberrant glial cells. *Assanah et al.*, on the other hand, showed that the overexpression of growth factors receptors without mutations to proto-oncogenes was sufficient to alter the function of progenitor cells and cause the growth of tumors within brains of mice (Assanah et al. 2006).

Even though the question of which cell population(s) initiate GBM is still being hotly debated (Safa et al. 2015; Soda et al. 2011), studies examining the effect of therapeutics on established GBM models have reported the existence of stem-like, plastic cells present within GBM tissue that tend to survive therapeutic exposure and later cause disease relapse (Bao et al. 2006; Baskar et al. 2012; Chen et al. 2012; Jackson et al. 2015; Mannino and Chalmers 2011; Murat et al. 2008; Weller et al. 2012). These tumor cells which show the genetic expression and functional characteristics of stem cells have been termed as glioblastoma stem cells (GSCs) (Bradshaw et al. 2016; Cabrera et al. 2015; Calabrese et al. 2007; Dalerba et al. 2007; Gilbertson and Rich 2007; Hanahan and Weinberg 2011; Heddleston et al. 2009; Lathia et al. 2015; Liebelt et al. 2016; Singh et al. 2004a, b). In recent years, further studies into the role of GSCs have indicated their importance in the maintenance of GBMs and reconstitution of tumors post therapy (Chou et al. 2012; Dewhirst et al. 2008; El Hallani et al. 2010; Heddleston et al. 2009; Jhaveri et al. 2016; Lathia et al. 2010; Nakada et al. 2013; Ogden et al. 2008; Ricci-Vitiani et al. 2010). Although several investigations have elucidated the role of GSCs and explored potential ways to target them to enhance the efficacy of current and future therapies, questions regarding the ontology of GSCs have not been conclusively answered (Brooks et al. 2013; Chen et al. 2014; Fan et al. 2010; Huang et al. 2012; Mendez et al. 2010; Persano et al. 2012; Sattiraju et al. 2017a). In addition, *Segerman et al.* *Chaffer et al.* and others showed that cancer cells are highly plastic and that terminally differentiated cancer cells dedifferentiate into cancer stem cells (CSCs) in response to stressors and therapy (Chaffer et al. 2013; Niklasson et al. 2017; Segerman et al. 2016). These studies highlighted the importance of tissue microenvironment in the regulation of cancer cell state, but further investigations are necessary to understand the importance of cellular plasticity in tumor maintenance and therapeutic resistance.

## Angiogenesis and Perfusion with GBM

GBM is characterized by extensive angiogenesis to allow the growth and survival of rapidly proliferating cells (Das and Marsden 2013; Jain et al. 2007). Hypoxic microenvironments cause GBM cell invasion and stimulate vascular and perivascular cells to produce pro-angiogenic factors (Brat et al. 2004; McCord et al. 2009; Rong et al. 2006). Vascular endothelial growth factor (VEGF) and its receptors, platelet-derived growth factor (PDGF), PDGF receptor-beta (PDGFR $\beta$ ), angiopoietins (Ang1 and Ang2), Tie2, matrix metalloproteinases (MMP-2 and MMP-9), bone morphogenic proteins (BMPs), etc. have been shown to be involved in this process (Jackson et al. 2017; Ribeiro and Okamoto 2015). The central nervous system (CNS) vasculature consists of tightly packed endothelial cells that are wrapped around by pericytes which provide structural support. Additionally, these vessels are further wrapped around by astrocytes that extend across endothelial cell tight junctions and by interneurons which together form the BBB. BBB is a protective vascular barrier that only allows the passive diffusion of water, oxygen, carbon dioxide, and highly lipophilic molecules. Glucose, amino acids, hormones, and larger molecules are actively transported across the BBB (Abbott 2002; Abbott et al. 2010). Co-opted vasculature and the newly formed angiogenic blood vessels within the GBM tissue show greater vascular permeability than normal CNS vasculature due to their disorganized architecture. This altered, often leaky vascular barrier within the tumor tissue is termed as the blood-tumor barrier (BTB) and results in regions of edema which hinder effective drug delivery (Agarwal et al. 2013; Dubois et al. 2014; Sattiraju et al. 2017b). The presence of BBB in peritumoral regions and a leaky BTB within the tumor results in ineffective perfusion of GBM tissue which further contributes to necrosis and hypoxia while certain parts of a tumor might escape exposure to systemic therapies, thereby resulting in recurrent disease (Pardridge 2005, 2012).

## Immune Microenvironment with GBM

A bulk of GBM tissue consists of infiltrated microglia and macrophages, but their response to tumor cells is often suppressed. Immune suppressive factors such as interleukin-10 (IL-10), IL6, transforming growth factor-beta (TGF- $\beta$ ), prostaglandin E2 (PGE2) suppress immune response against GBM cells, promote transformation of dendritic cells (DCs) into a regulatory phenotype and promote the activation of FOXP3+ regulatory T cells (Tregs). Hypoxia and subsequent expression of HIF-1 $\alpha$  and VEGF production have also been reported to cause Treg activation and immune suppression. Macrophages isolated from GBMs tend to polarize towards M1 (pro-inflammatory) and M2 (anti-inflammatory) phenotypes. Tumor-activated macrophages (TAMs) which are similar to M2 macrophages in function and cell surface marker presentation have been shown to play a vital role within the GBM microenvironment by promoting invasion and growth of tumor cells. TAMs have been reported to cause matrix degradation and enhance survival of GBM cells and GSCs. Hypoxic microenvironments within GBM have been reported to stimulate activation of microglia into TAMs (Jackson et al. 2017; Nduom et al. 2015; Razavi et al. 2016).

Activation of naïve T cells within the GBM microenvironment not only requires contact with major histocompatibility complexes (MHCs) on antigen-presenting cells (APCs) but also activation of co-stimulatory receptors (Driessens et al. 2009; Sharpe and Abbas 2006).

Based on significant findings in the past decade that shined the spotlight on the role of immune cells and immune-related mechanisms within tumor microenvironment that effect tumor evasion and clearance, a lot of efforts are currently being made to exploit these mechanisms to deliver therapeutics which could suppress inhibitory signals and allow immune cells to identify and clear tumor cells (Rosenberg and Restifo 2015; Wang et al. 2014; Yang 2015). Inhibiting immune checkpoints is currently the most commonly exploited mechanism for therapeutic purposes. Immune checkpoints prevent naïve T cells from causing autoimmune responses during infections by producing inhibitory signals. Programmed death-1 (PD-1) is an immune checkpoint co-stimulatory receptor expressed on the cell surface of T cells. Normal cells of the body express ligands for PD-1, namely PDL-L1 and PDL-L2, to prevent activation of naïve T cells and subsequent cytotoxicity. GBM cells exploit this mechanism and overexpress PDL-L1 on their cell surface to evade immune recognition and attack. Monoclonal antibodies that inhibit PD-1-PDL-L1 binding have shown to boost immune response against tumor cells and enhance survival in pre-clinical studies (Iwai et al. 2017). Promising results in clinical trials resulted in the approval of immune checkpoint inhibitors (ICIs) such as Nivolumab (approved for NSCLC, melanoma and renal cell carcinoma), Pembrolizumab (approved for melanoma, lung cancer and head and neck cancer), and Atezolizumab (approved for Urothelial and lung cancers) for the treatment of cancer patients (Alsaab et al. 2017; Hamanishi et al. 2016; Kang et al. 2016).

Cytotoxic T cell antigen-4 (CTLA4) is a co-stimulatory receptor, similar to PD-1, which negatively regulates T-cell activation. Inhibiting the binding of CTLA4 to its ligand had been shown to cause increased immune response against tumors. Monoclonal antibodies that inhibit CTLA4 binding such as Ipilimumab were approved for use in the clinic for melanoma patients after they showed promising results in preclinical and clinical trials (Alsaab et al. 2017; Hamanishi et al. 2016; Kang et al. 2016). The delivery of ICIs to GBMs is hindered, however, by the presence of BBB and the ineffective perfusion within tumor tissue due to angiogenic blood vessels (Hodges et al. 2016; Lyon et al. 2017; Rolle et al. 2010).

A recent phase-I study showed the potential for using interleukin-13 receptor alpha 2 (IL13RA2) directed chimeric antigen receptor (CAR) T cells against recurrent GBM. The IL13RA2 redirected CAR T cells in this study were delivered into the resection cavity of a 50-year-old patient using a neurosurgical technique called convection-enhanced delivery (CED) (Brown et al. 2016). CED provides the opportunity to bypass the BBB and enhance the efficacy of drugs by delivering them directly into the bed of the tumor (Bobo et al. 1994; Pardridge 2005; Lidar et al. 2004). Even though results from the study were promising, more efforts are needed to enhance the efficacy of this strategy.

### **Failure of Conventional Therapies Against GBM and Trends for the Future**

Recurrent disease is a major contributor to GBM patient mortality and is almost an eventual outcome for newly diagnosed patients whose tumors regress following adjuvant chemotherapy using temozolomide. Invasive GBM cells residing outside the area of resection that escape surgical debulking, therapy-resistant GBM cells that either attain stem cell-like characteristics through de-differentiation or retain their stem cell state after tumor

initiation (COIs), insufficient diffusion due to the BBB, and resulting accumulation of ineffective concentrations of systemic therapies within GBM tissue are thought to be major reasons for therapeutic failure and disease relapse (Chaichana 2014; Eyupoglu et al. 2013; Lathia et al. 2015; Pardridge 2005; Persidsky et al. 2006; Wolburg and Lippoldt 2002; Wolburg et al. 2012).

In recent years, a lot of attention has been placed on stymieing angiogenic processes using anti-angiogenic therapies such as Bevacizumab (which binds to and inhibits VEGF function) and to target GBM-specific cell surface receptors, often overexpressed by tumors such as the mutated EGFRvIII (Gilbert et al. 2014). These therapies have proven to be ineffective in causing a long-term effect on the progression of tumors and have not dramatically increased the median survival rate. Tumor heterogeneity, barriers to effective drug delivery, and additional factors involved in the promotion and maintenance of angiogenic vessels are thought to have caused the failure of these therapies in the clinic. Latest strategies to modulate the immune activity within GBM microenvironment, redirecting T cells using chimeric antigen receptors (CARs) towards GBM cells and oncolytic viruses have shown promise in preclinical investigations and phase-I clinical trials, but further research is needed to increase their effectiveness in the future. Additionally, preclinical investigations evaluating the efficacy of targeting GSCs, strategies to transiently enhance BBB permeability, therapeutic stem cells-based drug delivery and targeted molecular irradiation are currently underway to deliver systemic therapies more effectively to GBM tissue (Sattiraju et al. 2017b, c) (Fig. 2.1).

### Multifaceted Role of Pericytes Within GBM

#### Stemness and Tumor Initiation

Pericytes were previously thought to mainly play a role in supporting vascular architectures within the brain as part of the neurovascular unit (NVU) and to regulate blood flow within capillaries, but recent investigations have shed light on their role in tissue homeostasis and disease pathologies (Jackson et al. 2017; Sweeney et al. 2016). Using transgenic mouse models and cell surface receptor expression analysis, pericytes have been shown to be plastic multipotent perivascular cells which have the capability to differentiate into vascular smooth muscle cells, adipocytes, primary osteocytes, chondrocytes, fibroblasts, myofibroblasts, and neural cell lineages (Birbrair et al. 2014a, 2017a; Ribeiro and Okamoto 2015). Even though studies have shown pericytes to be stem cell-like, not all pericytes are plastic. *Birbrair et al.* had previously discovered two distinct subpopulations of pericytes in Nestin-GFP/NG2-DsRed double-transgenic mice, of which only Nestin+/NG2+ “Type-2” pericytes were observed to be involved in tumor angiogenesis while Nestin-/NG2+ “Type-1” pericytes were not, indicating that pericytes are heterogenous in their function (Birbrair et al. 2011, 2013a, b, 2014b). *Birbrair et al.* and others had discovered that pericytes (Type-2) can also be differentiated into neural and myeloid lineages to give rise to neural-like stem cells (NLSCs) and macrophages (Armulik et al. 2011; Birbrair and Frenette 2016; Birbrair et al. 2015).

In later studies, *Birbrair et al.* also observed that NLSCs derived from type-2 pericytes tend to migrate to the subventricular zone (SVZ) of healthy mice when implanted intracranially (Birbrair et al. 2013a). Importantly, *Birbrair et al.* also observed that in mice bearing



orthotopic GBMs, NLSCs migrate to regions of orthotopic tumors and invade along infiltrating margins when intracranially implanted in the ipsilateral hemisphere. These results indicate NLSCs to behave similarly to mesenchymal stem cells (MSCs) with regard to their migration to GBM sites when implanted intracranially but unlike MSCs which are co-opted by the tumor cells and transform into tumor-associated fibroblasts (TAFs), which assist in tumor growth and expansion, NLSCs did not differentiate into a TAF-like phenotype when co-cultured with GBM cells. In addition, unlike MSCs, NLSCs did not promote angiogenesis when in contact with GBM cells (Birbrair et al. 2017b). These results further contribute to the hypothesis that pericytes could be closely related to MSCs either as their precursors or as a specialized subpopulation of MSCs within the brain. Further adding fuel to this line of thought are reports showing that pericytes and MSCs share cell surface marker expression such as NG2, CD44,  $\alpha$ SMA, pDGFR $\beta$ , CD90, CD73, CD105, and Sca-1 (Crisan et al. 2008; Ribeiro and Okamoto 2015). The ability of pericytes to differentiate into neural and myeloid lineages could point to their role of pericytes in responding to injury and neurological diseases (Dore-Duffy et al. 2000; ElAli et al. 2014).

In GBM, GSCs have been reported to transdifferentiate into tumor pericytes, which form the majority of pericytes found within tumor tissue and assist in GBM cell proliferation and GSC self-renewal (Caspani et al. 2014; Cheng et al. 2013; Jackson et al. 2017). Neural stem cells (NSCs) have previously been shown to harbor the capacity to transdifferentiate into pericytes in normal brain tissue, indicating that GSCs could exploit such mechanisms to give rise to perivascular niche components (Cheng et al. 2013; Goldberg and Hirschi 2009). Due to the tendency of GSCs to localize near vasculature and due to their ability to give rise to malignant multipotent pericytes, researchers have suggested the possibility for such GSC-derived CD133+ “malignant pericytes” to drive tumor progression. *Appaix et al.* suggested an alternative “Cancer Pericytes” model for GBM initiation and progression where they hypothesized malignant CD133+ pericytes to act as COIs (Appaix et al. 2014). According to this hypothesis, existing pericytes behaving as MSCs could attain a neural stem cell-like phenotype by detaching from basement membrane, thereby forming a cancer stem-cell pool. These malignant pericyte-derived CSCs could then proliferate extensively, giving rise to a tumor mass which results in a hypoxic microenvironment. Subsequently, endothelial cells recruited through the CXCR4/ CXCL12 pathway could initiate angiogenesis. Malignant pericyte-derived CSCs could then differentiate into an aggressive mesenchymal phenotype, transdifferentiate into other cell types and drive tumor heterogeneity within the tumor mass or migrate to co-opted blood vessels to initiate another cycle of tumor formation. The authors speculated that such malignant pericytes could gain and lose CD133 expression during different stages of tumor formation and could thus explain the existence of CD133<sup>-</sup> stem cells within GBMs.

Additionally, *Zhang et al.* have reported that overexpression of cytoplasmic GT198 (a DNA repair gene that activates VEGF) within pericytes gives rise to tumors. The authors suggested that malignant pericytes could be derived from GT198 expressing GSCs or from normal pericytes that undergo somatic mutations upon microenvironmental stimuli. GT198+ malignant pericytes could also be resistant to radiotherapy and cause the failure of anti-VEGF therapies (Zhang et al. 2017).

Endothelial cells on co-opted blood vessels within the brain are thought to stimulate the migration of GSCs towards them through SDF-1/CXCR4 pathway and later induce their transdifferentiation into tumor pericytes by secreting TGF- $\beta$  (Cheng et al. 2013). This has been suggested as a mechanism to allow for the proliferation of endothelial cells, as tumor pericytes present within the perivascular niche secrete VEGF and other paracrine factors. In addition, enhanced pericyte coverage of co-opted and angiogenic blood vessels is thought to render resistance to anti-angiogenic therapies (Gabriele Bergers and Hanahan 2008; Ribeiro and Okamoto 2015). The ability of GSCs to give rise to tumor pericytes would also indicate their independence from relying on perivascular cells and their progenitors within the peritumoral region for engineering a suitable microenvironment.

*Caspani et al.* have suggested a “Dual Cell of Origin” hypothesis where pericytes drive tumor diversification upon direct contact with GSCs. In their study, the authors reported that pericytes attain a stem cell-like state upon transfer of cytoplasm from GSCs giving rise to GSC-pericyte cell fusions. The authors reported the existence of GBM cells that expressed labels for both pericytes and GBM cells within orthotopically implanted xenografts, suggesting that aneuploid cells derived from multipolar division of these GSC-pericyte cell fusions could drive GBM diversification (Caspani et al. 2014). The findings mentioned in the above section indicate that pericytes within NVU are plastic and play a critical role in GBM initiation and progression.

### Pericytes Are Involved in GBM Invasion

Pericytes are thought to play a protective role against tumor invasion by acting as physical barriers but poor and disorganized pericyte coverage is often observed within the tumor microenvironments which allows tumor to spread (Xian et al. 2006). Invading GBM cells tend to incorporate existing blood vessels into tumor tissue by a process termed “Vascular Co-option.” These co-opted blood vessels undergo necrosis upon angiopoietin-2 secretion by GBM cells, repeating events of hypoxia and subsequent invasion into surrounding areas of normal brain parenchyma (Liebelt et al. 2016; Reiss et al. 2005). Alternatively, edema caused due to incomplete coverage of angiogenic blood vessels by pericytes within tumor tissue has also been hypothesized to enhance tumor invasion by increasing intratumoral fluid pressure. This increased intratumoral fluid pressure is further thought to suppress surrounding blood circulation, resulting in hypoxic microenvironments, thereby stimulating the invasion of GBM cells into surrounding normal brain parenchyma (Cooke et al. 2012).

Cytoplasmic extensions of invading GBM cells around pericytes that they encounter, termed “Flectopodia,” have been suggested to facilitate the co-option of existing blood vessels by trafficking GBM cell cytoplasm into the cellular cortex of pericytes. GTPase Cdc42 was shown to play a critical role in the formation of such cytoplasmic extensions and pericyte activation (Caspani et al. 2014).

Poor pericyte coverage allows spreading of GBM, as incomplete coverage of angiogenic vessels allows vascular invasion by GBM cells. In addition, poor pericyte coverage also enhances the metastatic potential of other solid tumors that tend to migrate to the brain. Brain metastatic lung and melanoma cells that extravasate through capillaries were shown to survive and proliferate for long periods of time only when in contact with the abluminal



endothelial cells of capillaries in a pericyte-like position. This mechanism where tumor cells position themselves similar to pericytes when in contact with the abluminal endothelial cells has been extensively studied in melanomas and is termed as pericyte mimicry or angiotropism (Bentolila et al. 2016; Lugassy et al. 2014; Scott et al. 2015). Pericyte mimicry not only allows for the survival and proliferation of tumor cells but has also been reported to be exploited by tumor cells for extravascular migratory metastasis. This mechanism allows tumor cells to spread to local and distant sites by avoiding vascular invasion. GBM cells have been well documented in in vivo and in vitro studies to invade surrounding normal brain parenchyma along abluminal side of capillaries through the mechanism of pericyte mimicry (Scott et al. 2015).

As mentioned in the previous section, the ability of GSCs to attain a pericyte-like phenotype and their similarities to pericytes within the GBM microenvironment, with regard to cell surface marker expression, could be a result of pericyte mimicry of tumor cells and could support the hypothesis that pericyte-like cells within the GBM microenvironment could also act as initiating cells.

### Pericytes and the BBB

Pericytes have been shown to play a vital role in regulating BBB permeability by controlling the expression and alignment of tight and adherens junction proteins along with the transcytosis of molecules across the BBB (Sweeney et al. 2016). The importance of pericytes for the structural stability of CNS vessels and for maintaining the BBB was observed in PDGF $\beta$  knockout mice where pericyte depletion resulted in enhanced CNS vascular permeability due to BBB disruption (Armulik et al. 2011; Bell et al. 2010; Daneman et al. 2010; Sweeney et al. 2016). PDGF-BB(ligand)-PDGFR $\beta$  signalling pathway is critical for pericyte-endothelial cell interactions to maintain BBB stability and to regulate extravasation across the barrier into brain parenchyma. Transgenic mice with both PDGF and PDGFR $\beta$  null mutations have shown embryonic lethality due to the development of microvascular instability and hyperplasia, aneurysms, and microhemorrhages (Sweeney et al. 2016). In studies where PDGF-BB-PDGFR $\beta$  signalling pathway was partially disrupted, age-dependent BBB disruption was observed (Armulik et al. 2011; Bell et al. 2010; Daneman et al. 2010; Sweeney et al. 2016). Similarly, *Abramsson et al.* showed that the lack of PDGFR $\beta$  expression on pericytes resulted in their poor vascular coverage and enlargement of blood vessel diameter, resulting in increased leakiness (Abramsson et al. 2002). In addition, TGF $\beta$ , Ang1, and Notch signaling have been reported to play an important role in maintaining the integrity of the BBB (Bruna et al. 2007; Cheng et al. 2013; Liebner and Plate 2010; Reis and Liebner 2013; Sweeney et al. 2016). TGF $\beta$ -TGF $\beta$ R2 signaling promotes pericyte maturation, proliferation, and attachment to endothelial cells. Aberrant signaling of downstream effectors of TGF $\beta$ -TGF $\beta$ R2 pathway such as Smad1, Smad2, and Smad4 results in vascular destabilization and brain hemorrhages (Goumans and Mummery 2000; Li et al. 2011; Maddaluno et al. 2013; Sweeney et al. 2016). Additionally, forkhead transcription factor (Foxf2) which affects TGF $\beta$  signaling also plays a role in maintaining BBB integrity as Foxf2 knockout transgenic mice show BBB disruption, hemorrhages, perivascular edema, increase in luminal endothelial caveolae, and thinning of basal lamina of capillaries (Reyahi et al. 2015).

Pericytes have been observed to attach loosely and improperly cover angiogenic vessels within GBMs, yet proper pericyte coverage has been shown to result in stabilization of tumor blood vessels, thereby accelerating GBM cell proliferation and invasion. High TGF- $\beta$ -Smad activity was shown to result in highly aggressive and proliferative GBMs in patients which conferred a poor prognosis. TGF- $\beta$ -Smad pathway has also been shown to activate PDGFB gene expression in primary GBM cells in vitro (Bruna et al. 2007). Majority of pericytes found in GBMs are thought to arise from GSCs that transdifferentiate upon migration to endothelial cells through SDF-1-CXCR4 signaling pathway. TGF- $\beta$  has been shown to play a vital role in inducing this transdifferentiation of GSCs into pericyte-like cells which help promote angiogenesis, stabilize tumor vessels, and contribute to GBM growth (Cheng et al. 2013).

Notch3, expressed by brain pericytes, has been shown to play an important role in interactions with endothelial cells and in maintaining the integrity of BBB. Abnormal TGF- $\beta$ -Notch signaling has been reported to cause cerebral cavernous malformation. Transgenic mice with dysfunctional Notch signaling caused by knockout of RBP-J $\kappa$  transcriptional factor, showed brain hemorrhages (Li et al. 2011; Maddaluno et al. 2013; Sweeney et al. 2016). Like TGF- $\beta$ , Notch1 signaling has also been shown to induce transdifferentiation of GSCs into pericyte-like cells in vitro which promote angiogenesis (Guichet et al. 2015). Angiopoietin-1 (Ang1) expressed by pericytes, which binds to Tie2 receptor tyrosine kinase on endothelial cells, maintains BBB integrity. Like TGF- $\beta$  and Notch1, knockout of Ang1 in transgenic mice resulted in BBB disruption and promotion of angiogenesis (Suri et al. 1996). The major facilitator superfamily domain-containing 2a (MFSD2A), a BBB transporter whose expression is thought to depend on presence of pericytes, has also been reported to play a role in the formation and maintenance of BBB integrity (Ben-Zvi et al. 2014).

Pericytes interact with astrocytes through apolipoprotein E (APOE4)-LRP1 interaction, resulting in the activation of MMP-9 activity by signaling through cyclophilin A (CypA)-NF- $\kappa$ B pathway, which promotes inflammation. Increased APOE4-LRP1 signaling and resulting increase in the activity of MMP-9 causes BBB disruption due to degradation of endothelial tight junctions and basement membrane (Bell et al. 2012; Sweeney et al. 2016). Pericytes also regulate perfusion through brain capillaries and regulate BBB permeability through their contractile nature which is facilitated by synthesis of vimentin, actin and myosin microfilaments, tropomyosin and desmin. As part of the NVU, pericytes are also in contact with interneurons and receive communications from the nervous system as the average distance between a neuron and a brain capillary is 8–23  $\mu$ m. Conditioned media from cultured human brain pericytes has shown the presence of neurotrophic factors and pericytes have been shown to regulate capillary diameter and cerebral blood flow upon signaling from neurons (Hawkes et al. 2011; Lovick et al. 1999). Studies have shown that norepinephrine leads to pericyte contraction (reduction of capillary diameter) while GABA, dopamine, glutamate, and adenosine cause pericyte relaxation (increase in capillary diameter) (Sweeney et al. 2016).

The BBB provides a formidable challenge for delivering drugs into the CNS, especially to brain tumors. In addition to intact BBB in areas around brain tumors, leaky angiogenic

vessels which cause edema in certain areas of tumors result in the ineffective and non-uniform delivery of systemically delivered therapies (Sattiraju et al. 2017a, b). Ineffective drug delivery is thought to be one of the major reasons for GBM relapse, as cells that are not exposed to effective concentrations of systemic therapies reduce therapeutic efficacy. Therefore, efforts are being made to either transiently disrupt the BBB or to target cells involved in regulating BBB permeability, in order to enhance the extravasation of systemic therapies into GBM. As pericyte recruitment and stabilization of blood vessels within brain tumors have been shown to regulate vascular permeability, strategies to disrupt pericyte recruitment by angiogenic endothelial cells are currently being investigated.

In a recent study by *Behling et al.* by targeting monomeric vascular endothelial cadherin, which is expressed on angiogenic vessels and endothelial progenitor cells (EPCs), using a monoclonal antibody E4G10 that was labeled to  $\alpha$  particles, the authors reported enhanced survival in Ntva transgenic mice bearing GBMs. As extensive proliferation of pericytes is usually observed in this GBM model, their higher density was suspected to protect endothelial cells from radiotherapy. In addition to mitigating tumor growth, the authors also observed normalization of the morphology of angiogenic vessels, reduction of edema, and a decrease in pericyte coverage of these vessels. In addition, depletion of regulatory T cells (Tregs) and EPCs was also observed (Behling et al. 2016). In another recent study, *Sattiraju et al.* showed enhancement in BBB permeability in brains of mice bearing orthotopic GBMs by targeting integrin alpha-V beta-3 ( $\alpha_v\beta_3$ ) using an  $\alpha_v\beta_3$  targeted antagonist conjugated to partially polymerized liposomes that was labeled to  $\alpha$  particles. Apart from observing enhanced BBB permeability, the authors also reported tumor cytotoxicity evidenced by nuclear accumulations of  $\gamma$ H2Ax double-strand DNA break repair protein within tumor mass. Overexpression of  $\alpha_v\beta_3$  by GBM cells, especially at invasive ends and the presence of proliferating malignant cells in perivascular regions could explain this effect. The authors in this study also observed enhanced BBB permeability in peritumoral and normal regions of the brain surrounding tumors, indicating that enhanced vascular permeability in these distal regions might not have been caused due to direct a particle-induced cellular effects (Sattiraju et al. 2017c).

Studies by both *Behling et al.* and *Sattiraju et al.* show the feasibility of targeting vascular and perivascular components within the NVU using short-ranged, targeted molecular irradiation to either enhance vascular perfusion or BBB permeability, to better deliver systemic drugs to GBMs and to reduce radio-resistance conferred by perivascular cells within GBM microenvironment. But, as evidenced in previous studies where pericyte ablation caused enhanced invasion and metastasis, further long-term evaluations to assess tumor resistance and remission in abovementioned strategies are necessary (Bentolila et al. 2016; Lugassy et al. 2014; Scott et al. 2015). *Xiong et al.* reported an alternative strategy of enhancing BBB permeability by remotely activating intracranially implanted genetically engineered MSCs using high-frequency focused ultrasound (HIFU) to secrete TNF- $\alpha$ . This study highlights the ability to locally deliver factors which can transiently influence vascular and perivascular cells that regulate BBB integrity and perfusion within normal brain parenchyma and GBMs, thus enhancing the delivery of systemic drugs to GBMs and CNS (Sattiraju et al. 2017b; Xiong et al. 2015).

## Pericytes Drive Angiogenesis

Angiogenesis is a crucial multi-stage process resulting in the transformation of GBM into an aggressive disease (Jain et al. 2007). Highly proliferating cells within GBM tissue drive up nutrient and oxygen demand and the resulting hypoxic microenvironment stimulates the production of new and often-disorganized angiogenic blood vessels. Pericytes play a major role in this process by forming a scaffold for newly proliferating endothelial cells to form blood vessels and secrete factors that stabilize these newly formed angiogenic vessels (Mancuso et al. 2006; Ribeiro and Okamoto 2015). The process of angiogenesis is tightly regulated and requires direct contact and crosstalk between endothelial cells and pericytes. Pericytes are involved in secreting factors that stimulate endothelial tip sprouting from existing blood vessels in their vicinity, recruit endothelial cells, and aid in their proliferation. Upon endothelial recruitment and formation of a vessel structure, pericytes within GBM microenvironment wrap endothelial cells which results in vessel stabilization and maturation (Caspani et al. 2014).

Endothelial cells are thought to produce PDGF $\beta$  which recruits PDGFR $\beta$  expressing pericytes, which in turn secrete VEGF and Ang-1 to stabilize angiogenic blood vessels (Hellstrom et al. 1999). Studies examining the effect of a pan tyrosine kinase inhibitor (SU6668), which preferentially targets PDGFR $\beta$  in RIP1-Tag2 transgenic mice that generate pancreatic islet carcinomas showed that ablating pericytes from tumor blood vessels caused vascular regression (Bergers et al. 2003). GSC-derived pericyte-like cells within the GBM microenvironment have also been reported to express PDGFR $\beta$  and aid in vessel stabilization and maturation (Cheng et al. 2013). Inhibiting PDGF-BB-PDGFR $\beta$  signaling resulted in regression of tumor vessels, indicating that it affects the survival of endothelial cells, GSC-derived pericyte-like cells and co-opted pericytes (Hellstrom et al. 2001). Overexpression of PDGF $\beta$  by endothelial cells has been reported to increase pericyte coverage and accelerate tumor growth (Furuhashi et al. 2004).

TGF- $\beta$ , expressed by pericytes in a latent form, also plays a critical role in angiogenesis and GBM progression. TGF- $\beta$ -TGF $\beta$ R2 signaling has been reported to result in the stabilization of angiogenic vessels, and high expression of TGF- $\beta$ -Smad has been reported to result in a poor prognosis for GBM patients. TGF- $\beta$ -Smad pathway has been shown to induce expression of PDGF-B in GBM cells and the transdifferentiation of GSCs into pericyte-like cells, which aid in angiogenic vessels stabilization and maturation (Goumans and Mummery 2000; Maddaluno et al. 2013). TGF- $\beta$  signaling alone has been reported to be able to induce genetic and phenotypic changes, often seen in altered vasculature within GBMs.

Notch1-DII4 signaling has been reported to be critical in regulating sprouting angiogenesis. Inactivation of Notch1 or DII4 genes have shown to cause increased endothelial tip sprouting and tip-cell numbers, indicating that Notch1-DII4 signaling at endothelial sprout restricts tip sprouting towards VEGF-A gradients, thereby ensuring proper sprouting and branching patterns (Gerhardt et al. 2003). Pericytes have been reported to enhance the survival of endothelial cells and promote angiogenesis by secreting VEGF-A, which binds to VEGFR2 expressed on endothelial cells. VEGFR2 stimulates the upregulation of survival genes such as Bcl-2, survivin and X-linked inhibitor of apoptosis protein (XIAP) in endothelial cells (Franco et al. 2011). VEGF-A-VEGFR2 autocrine signaling within

endothelial cells also promotes their survival. Vitronectin secreted by pericytes also causes an increase of VEGF-A expression in endothelial cells through integrin  $\alpha_v$ -NF $\kappa$ B signaling, which in turn causes intracellular stimulation of VEGFR2 to promote endothelial cell survival (Franco et al. 2011; Sweeney et al. 2016). Inhibiting angiogenesis by interfering with VEGF-A-VEGFR2 signaling was thought to provide significant benefit to GBM patients as it would result in the collapse of existing angiogenic vessels and prevent further tumor vascularization and growth. Bevacizumab (Avastin), a monoclonal antibody against VEGF-A showed very promising results in preclinical studies and was tested in two phase-III clinical trials (RTOG and AVAGlio) as a stand-alone and combinational therapeutic for patients with recurrent GBM (Friedman et al. 2009; Gilbert et al. 2014). Although patients initially showed decreased tumor growth, their tumors eventually grew resistant to the drug, resulting in a progression-free survival of about 10.6 months in Phase-III clinical trials with no added benefit to overall survival when compared to placebo arm (Friedman et al. 2009; Hamza et al. 2014).

Ephrin receptor EphB4, which controls vascular morphogenesis within the NVU during developmental angiogenesis, and its ligand ephrin-B2, which is expressed on brain pericytes and endothelial cells, play a vital role in pericyte-endothelial cell interactions. Ephrin-B2-EphB4 signaling has been reported to regulate pericyte migration and interaction with maturing blood vessels and could be involved in the vascular remodeling within the GBM microenvironment (Augustin and Reiss 2003). Vascular cell adhesion molecule-1 (VCAM-1), N-cadherin and integrin  $\alpha_4\beta_1$  play critical roles in regulating pericyte migration, vessel maturation, and survival of endothelial and mural cells during angiogenesis (Gerhardt et al. 2000).

Overexpression of endosialin (CD248) has also been linked to pro-angiogenic role of pericytes within the GBM microenvironment (Brady et al. 2004). Angiogenic vessels within GBMs were reported to be composed of two layers of pericytes, an abluminal layer of proliferating cells and an adluminal layer of cells surrounded by basal lamina. By restricting the recruitment of pericytes and their signaling with endothelial cells during angiogenesis, newly formed GBM blood vessels could be disrupted causing reduced GBM invasion and growth.

In a study by *Svensson et al.* using a RGS5-GFP transgenic mouse model, the authors showed that PDGFR $\beta$  and neuroregulin-2 (NG2)-expressing pericytes were activated and recruited to blood vessels within orthotopically implanted GBMs from peritumoral and distal regions within ipsilateral and contralateral hemispheres, including from the rostral SVZ. Activation and recruitment of pericytes from distant regions of the brain towards tumor tissue has been attributed to parenchymal diffusion of paracrine factors such as hypoxia inducible factor-1 $\alpha$  (HIF-1 $\alpha$ ) through cerebrospinal fluid or cerebral edema. These tumor trophic pericytes showed CD13 MSC cell surface marker expression in areas surrounding tumor tissue, but did not show CD13 expression within tumor tissues, indicating a phenotypic shift as pericytes enter paracrine signaling networks within the GBM microenvironment. The results from this study indicate that pericytes migrate and integrate into the vasculature within GBMs, stabilize angiogenic blood vessels, and promote further angiogenesis (Svensson et al. 2015). In another study by *Huang et al.*, NG2 was identified to

be critical in integrin  $\beta 1$ -dependent pericyte-endothelial cell interactions and ablation of NG2 resulted in two-fold reduction in vessel ensheathment by pericytes. NG2 ablation was also reported to cause reduced collagen IV deposition due to loss of collagen VI anchorage. Impaired vessel maturation and stabilization within GBM tissues of mice lacking NG2 expression resulted in decreased tumor progression, highlighting the importance of NG2 for successful angiogenesis (Huang et al. 2010).

### Pericytes Contribute to Immune Microenvironment

GBM cell survival, proliferation, and invasion into peritumoral regions are thought to be facilitated by their evasion of immune surveillance by using multiple immunosuppressive mechanisms. Pericytes have been shown to regulate immune cell activity by secreting chemokines and other factors in addition to differentiating into macrophages, thus playing a critical role in immune surveillance and tumor clearance within the GBM microenvironment (Valdor et al. 2017). Brain pericytes have been reported to exhibit phagocytic properties and express macrophages cell surface markers such as CD11b, CD68, and MHC class II. Stimulation by IL-1 $\beta$  has been reported to result in the upregulation of iNOS and COX-2 within porcine brain pericytes (Balabanov et al. 1996). In addition, Pieper et al. reported that stimulation by IFN- $\gamma$  or TNF- $\alpha$  resulted in an antigen-presenting activity within brain pericytes (Pieper et al. 2014). Pieper et al. also showed that stimulation by LPS, TNF- $\alpha$ , and IL-1 $\beta$  resulted in chemotactic recruitment and transmigration of neutrophils into the brain parenchyma (Pieper et al. 2013).

During GBM establishment and expansion, pericytes have also been reported to be co-opted by tumor cells through direct cell-cell contact to facilitate immunosuppression and eventual immune evasion of GBM cells (Caspani et al. 2014). Upregulation of PD-L1, CD90, PDGFR $\beta$ , CD248, and *Rgs5* which inhibit CD4 $^{+}$  and CD8 $^{+}$  cytotoxic T cell activity was reported in pericytes derived from within-tumor microenvironments (Bose et al. 2013; Ochs et al. 2013; Ribeiro and Okamoto 2015). *Valdor et al.* recently reported that pericytes conditioned in vitro by GBM cells are characterized by high levels of anti-inflammatory cytokines, suppress the function of cytotoxic T cells and reduce antigen presentation within perivascular regions of orthotopically implanted GBMs. GBM conditioned pericytes (GBM-PCs) showed high levels of IL-10 and TGF $\beta$  anti-inflammatory cytokine expression (100–400 pg/mL) and low levels of TNF $\alpha$  expression (45 pg/mL) in vitro when compared to control native pericytes, indicating an immunosuppressive phenotype. GBM-PCs also showed upregulation of *Ii4ra* (encoding interleukin-4 receptor alpha), *Ii1m* (encoding interleukin-1 receptor inhibitor), and increased expression of angiogenic cytokine IL-6. However, expression of CD80 and CD86 co-stimulatory molecules was reported to be reduced (Valdor et al. 2017).

The exact mechanisms through which pericytes influence immune activity within the tumor microenvironment are not yet clearly understood and further research is required to gain a pristine understanding of the multifaceted role that pericytes play within tumor microenvironments and the way in which their role changes during tumor progression and in response to therapy (Fig. 2.2).



**Conclusions: Opportunities for Therapeutic Intervention**—The median overall survival of GBM patients has been improved to ~16 months using surgical resection followed by adjuvant temozolomide, but recurrent disease remains a major contributor to patient mortality (Roh et al. 2017). Novel therapeutic strategies such as immune checkpoint inhibitors, alternating electric fields, CAR T cells, stem cell-based drug and oncolytic viral delivery and strategies to enhance delivery of systemic therapies show the potential to increase patient survival and to reduce GBM recurrence in the future. As highlighted in this chapter, microenvironmental components play a vital role in the survival and proliferation of GBM cells. It is therefore critical to appreciate the complex relationship between GBM and their microenvironment and to design therapeutic strategies that would not just target one component of their microenvironment, as it would only serve to partially impede GBM survival and progression. The studies and their finding mentioned in this chapter highlight the critical role that pericytes play at various stages of GBM development.

The failure of multiple anti-GBM therapeutics in phase-III clinical trials has been attributed to tumor heterogeneity, ineffective drug delivery, invasive GBM cells, and therapy-resistant GSCs. As this chapter details, pericytes have been reported to significantly contribute to stemness, angiogenesis and altered perfusion, altered BBB permeability, GBM cell invasion and suppression of immune activity within the GBM microenvironment. It is therefore important to further elucidate the role of GSC-derived pericytes and native brain pericytes that are either present within or recruited into the tumor microenvironment. As previous studies have shown, targeting and ablating pericytes might not always stymie the growth and survival of GBM cells. This could be due to the multifaceted role of pericytes within the tumor microenvironments, their cellular plasticity and the existence of different subtypes of pericytes that contribute to various events at various stages of GBM establishment and expansion.

Knowledge of the role that pericytes play within the GBM microenvironment would therefore allow us to design therapies in the future such that they can circumvent cellular events which could otherwise compromise their efficacy and to also possibly design strategies to mitigate the role played by pericytes and other perivascular components co-opted by GBMs in treatment resistance.

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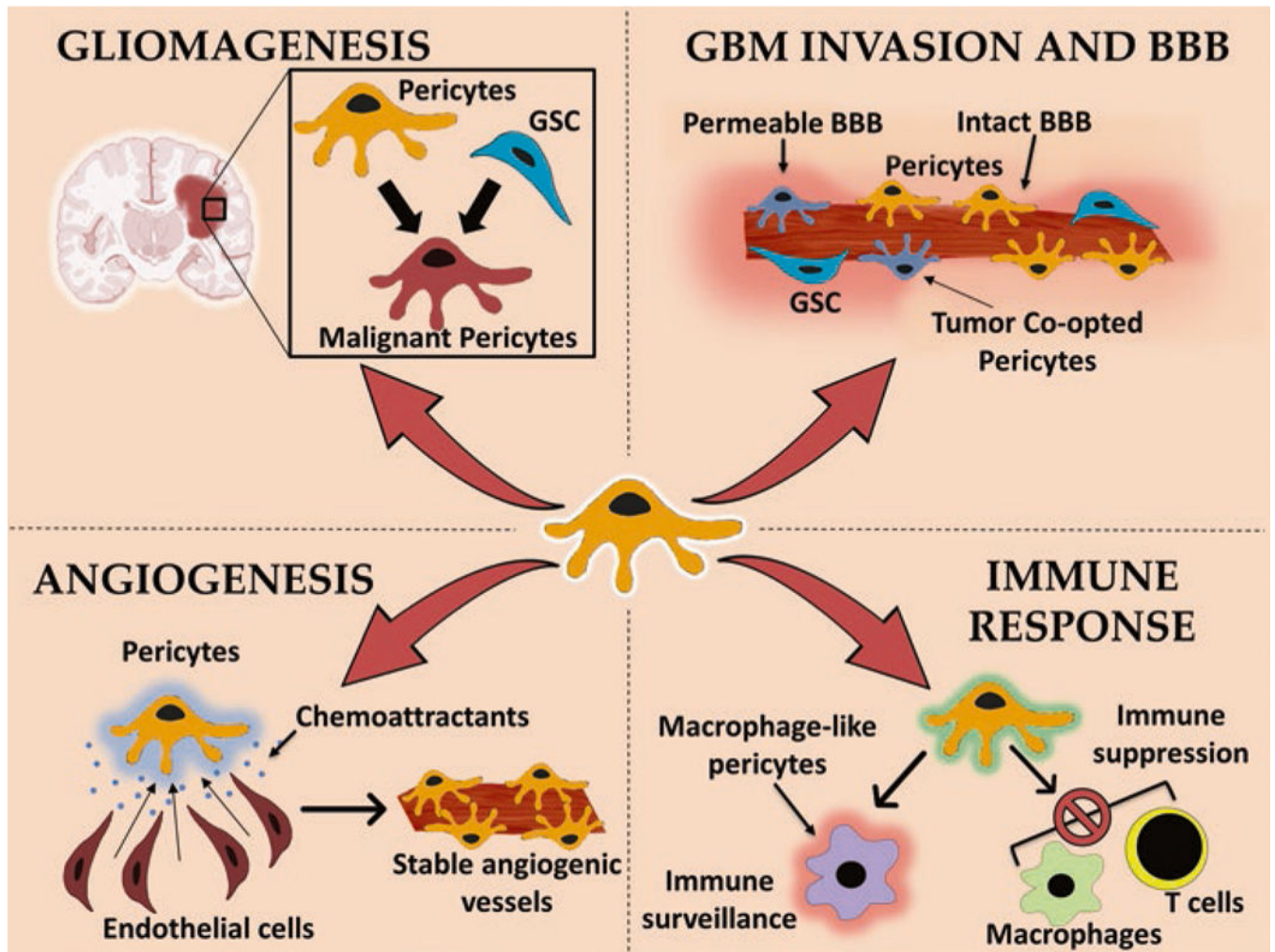


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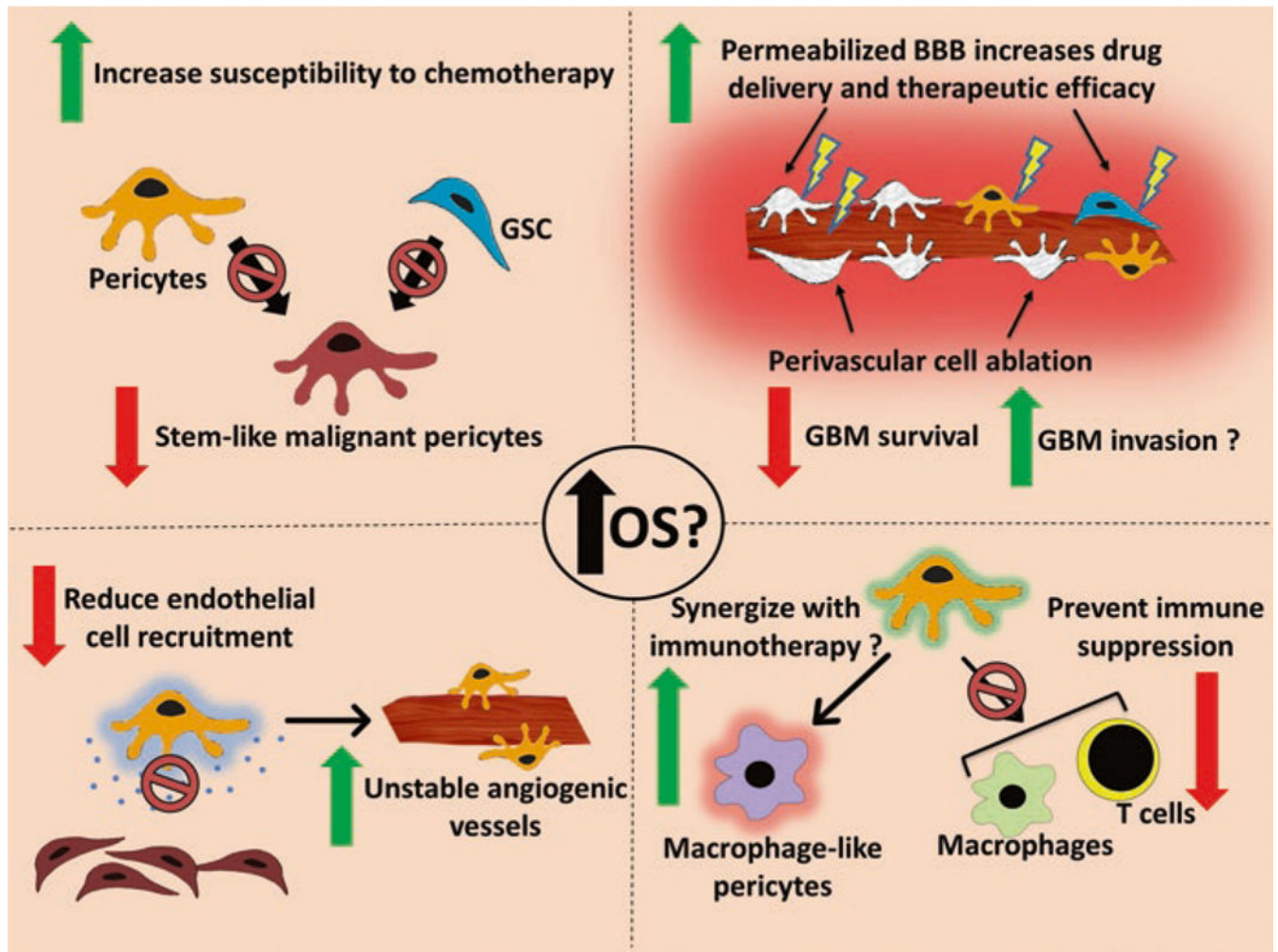
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**Fig. 2.1.** Diagram showing the multifaceted role of pericytes in various critical events of GBM initiation, establishment, maintenance, and progression. *GSC* glioblastoma stem cells, *BBB* blood-brain barrier





**Fig. 2.2.** Diagram showing potential therapeutic interventions which could limit the contribution of pericytes to GBM overall survival and progression. *OS* overall survival, *GSC* glioblastoma stem cells, *GBM* glioblastoma, *BBB* blood-brain barrier