Brain Barriers

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Barriers between the brain and systemic circulation are dynamic and highly specialized to strictly regulate the access of a wide variety of molecules to the brain. These barriers allow for the delivery of nutrients and other molecules necessary for neuronal functioning, but often limit the permeation of xenobiotics, including drugs. In brain tumors, these barrier functions may be disrupted or altered. However, this disruption is often heterogeneous and not reliable to guaranteee the delivery of efficacious concentrations of antineoplastic agents to brain tumors.

brain barriers

blood-tumor barrier

neurovascular unit

1. Introduction

The blood-brain barrier (BBB) remains one of the greatest obstacles to effective pharmaceutical interventions in the treatment of central nervous system (CNS) disease, including brain tumors. While it is true that some loss of neurovascular and barrier integrity may occur in and around brain tumors, the magnitude of this change is not consistent, and new pharmaceutical strategies for the treatment of brain tumors have yet to show significant efficacy in the clinic^{[1][2][3]}. This lack of efficacy is largely attributed to insufficient drug delivery due to the presence of the BBB. The dense vascular network of the brain works to strictly regulate the transport of substances into and out of the brain parenchyma in order to maintain ionic homeostasis, nutrient supply, and removal of waste for optimal neuronal function. In recent decades, research has revealed that the BBB is composed of specialized endothelial cells (ECs), which are surrounded and supported by pericytes and astrocytes and are regulated by neuronal signaling, forming what is referred to as the neurovascular unit (NVU)^[4]. A lack of vesicular transport across these specialized ECs and the presence of active efflux proteins help to further restrict the access of drugs to the CNS^[5]. Currently, treatment for the majority of brain tumors involves maximal surgical resection, if possible, followed by radiation, and in the case of glioblastoma multiforme (GBM), concomitant temozolomide (TMZ)^[2]. However, these treatments often prove to be palliative, and malignant brain tumors are nearly always fatal within five years of initial diagnosis^{[6][Z]}.

2. Barriers and Boundaries in the Brain

Rational drug delivery to any organ requires a thorough understanding of the structures and properties of the target tissue. This section includes detailed features of the dynamic NVU model that is rapidly supplanting the former static BBB concept. Furthermore, the role that these features serve in guiding the molecular basis for current therapies for CNS tumors is illustrated.

2.1. CNS Blood-Tissue Barriers

Any strategy for blood-borne drug delivery into the CNS must consider several structural obstacles related to blood-tissue interfaces^{[2][9]}. First, the brain is covered by layers of cells collectively described as the duraarachnoid-pia membranes. The dura separates peripheral vessels within the cranium from the cerebrospinal fluid (CSF) in which the brain resides. Several compact layers of epithelial cells, with tight junctional contacts and relatively low surface areas, prevent materials from traversing this boundary. Within the CSF compartment, the pial layer contains vessels that penetrate the brain parenchyma. These vessels also exhibit very low permeability and surface area. Other major barriers include relatively small regions of CNS circulation and include the choroid plexus, circumventricular organs (CVOs), and ependymal cells. Each of these has specialized epithelial cells with properties that highly restrict water-soluble chemicals from penetrating the mass of parenchymal tissue. The choroid plexus that produces CSF, for example, contains specialized epithelial cells with tight junctions (TJs) encasing a fenestrated vascular endothelium that allows the exchange of blood constituents with the extracellular space. The CVOs are also composed of fenestrated endothelial cells, but their location is confined by surrounding tanycytes, a specialized epithelial cell that also has tight junctional contacts that localize and restrict the interstitial fluid. Direct exchange between the CSF and interstitial fluid is restricted by ependymal cells that line the ventricular surface and form a selectively permeable cellular barrier.

2.2. Neurovascular Unit

By far, the major blood–tissue interface in the brain is the microvasculature network extending from arterioles to capillaries and to venules. By illustration, in one gram of human brain, the length of vessels, if joined end to end, approximates the length of 4.5 football fields and almost a quarter (~23%) of the surface area of a sheet of photocopy paper. The current model of the brain microvasculature consists of several different cell types working collectively to form a functional NVU^{[4][10]}. Endothelial cells joined by tight junctions that block paracellular diffusion are surrounded by pericytes that form gap junctions with multiple adjacent endothelial cells. Astrocytic endfeet also cover >99% of the endothelial–pericyte cell surface (Figure 1). The astrocytes, in turn, extend processes that monitor synaptic activity and react by signaling endothelial cells and pericytes to respond to increased metabolic demands by increasing nutrient delivery. Microglia, the resident immune cells, are extravascular when dormant but react swiftly to remove cellular debris (by phagocytosis) or respond to inflammatory signals associated with disease or injury. Loss of pericytes and their signaling in the NVU by injury or genetic means leads to reduced expression of endothelial tight junction proteins and dysfunction of their permeability barrier^{[11][12]}. These examples illustrate the remarkable dynamics, plasticity, and interdependence of signaling among the NVU cells to maintain functional stability in the contemporary model of the neurovasculature.



Figure 1. The neurovascular unit/blood–brain barrier (NVU/BBB) is composed of specialized endothelial cells and support cells, including pericytes and astrocytes. The cross-sectional view illustrates that the majority of the abluminal surface of the endothelial cell is covered by pericytes and astrocytic foot processes. Paracellular transport across the BBB/NVU is restricted by tight junction proteins, and even small, lipophilic molecules that might diffuse across the BBB may be subject to active efflux by a variety of proteins. Facilitated active transport, receptor-mediated transport, and ion transporters allow the brain to be supplied with nutrients while maintaining strict homeostasis.

2.3. Blood-to-Brain Permeability and Transport

As the brain depends on external nutrients for growth and development and yet must be protected from the influence of circulating toxins or xenobiotics, the specialized endothelial cells of the brain express critical membrane-imbedded proteins that function as transporters. One group of transporters includes facilitated carriers and secondary active transporters for the delivery of energy substrates and essential nutrients (Table 1). The glucose transporter (GLUT1), monocarboxylic acid transporter (MCT1), and amino acid transporters are examples of the >40 transporters detected by functional and transcriptomic analyses^[13].

A second type of transporter that is critical to brain drug delivery is the ABC (ATP-binding cassette) superfamily that uses the energy of ATP hydrolysis to expel endogenous and exogenous xenobiotics from the cell (and from the

brain) and return them to the blood for transformation and excretion (Table 2). The most relevant ABC transporters expressed by brain endothelial cells are P-glycoprotein (P-gp, ABCB1), breast-cancer-related protein (BCRP, ABCG2), and the multiple drug-related proteins (MRP1, -4, -5; ABCC1, -4, -5). Many anticancer drugs are substrates of the ABC transporters and, therefore, may influence their effectiveness as brain cancer drugs (Figure 1). The relevance of ABC transporters in anticancer drug delivery in brain tumors with apparently high permeability is illustrated by the fact that the most permeable tumor vasculatures have influx rate constants several-fold less than the rate constants for other organs such as muscle, heart, lung, kidney, and liver^[14]. Therefore, efflux mechanisms are likely important even in CNS tumors, in which the permeability of the brain vasculature may be compromised and elevated compared to the surrounding tissue.

Transport System	Typical Substrate	SLC Family	Common Name
Carbohydrates			
Hexose	Glucose	SLC2A1	Glut1
Sodium Myo-inositol	Myo-inositol	SLC5A3	SMIT
Monocarboxylates			
Monocarboxylic acid	Lactic acid ketones	SLC16A1	MCT1
Amino Acids			
Large neutral amino acid	Phenylalanine	SLC7A5	LAT1
Small neutral amino acid	Alanine	SLC38A2	SNAT2, -3, -5
Cationic amino acid	Lysine	SLC7A1	Cat1, CAT3

Table 1. Endothelial cell membrane transporters: partial list of common carriers.

Beta amino acid	Taurine	SLC6A6	TauT
Ala-Ser-Cys	Ala, ser, cys	SLC1A4	ASCT1, -2
Excitatory amino acid	Glutamic acid	SLC1A2	EAAT-1, -2, -3
Glycine	Glycine	SLC6A9, A5	GT-1
Others			
Fatty acids	Essential FA LPC-PC (DHA)	SLC44A1/2 Mfsd2A	FATP-1, -4 Mfsd2A
Nucleoside	Adenosine	SLC29A1 SLC28A1	ENT-1, -2; CNT1–3
Hormones	Thyroid T3	SLC16A2	MCT8
	Thyroid T4	OATP1C1	OATP1C1
Biotin, pantothenic acid	biotin	SLC5A6	SMVT
Folic acid	Folinic acid	SLC46A1	PCFT
Copper	Cu ⁺	SLC31A1	CTR1

Table 2. Brain endothelial cell transporters of xenobiotics/drugs. Members of the ABC (ATP-binding cassette)superfamily of transporters demonstrated in brain endothelial cells and non-ABC transporters of organic chemicalspotentially present are listed.

Transport System	Common Name	Typical Substrate
ATP Binding Cassette Transporter (ABC)		

ABCB1	P-gp	Broad-spectrum, xenobiotics
ABCG2	BCRP	mitoxantrone anthracycline xenobiotics
ABCC1	MRP1	GSSG, leukotrienes
ABCC5	MRP5	Thiopurines, cyclic nucleotides
ABCC4	MRP4	Organic anions
Non-ABC Transporters		
SLC22A7	OAT2-3	
SLC22A8	OATP1A4	
SLC20A2	OATP2B1	Organic ions
SLCO1A4	OCTN2	
SLCO2B1	OCT1-3	

3. Heterogeneous Blood–Tumor Barrier Permeability

The understanding of the BBB's physical and biochemical barrier functions, including the expression of tight junction proteins, restricted paracellular transport, and active efflux mechanisms, has been well established. However, determining the integrity of the NVU/BBB in and around tumors and how this affects tumor treatment has been less straightforward. In the case of both primary and metastatic tumors, the NVU/BBB is subject to changes due to tumor growth and signaling, and these alterations in NVU/BBB integrity and physiology result in what will hereafter be referred to as the blood–tumor barrier (BTB). The BTB may be characterized by an inflammatory environment with increased numbers of activated astrocytes, vascular endothelial growth factor (VEGF)-induced reduction in the expression of tight junction proteins like claudin-5, breakdown of the basal lamina, and tumor cell interference in associations between endothelial cells and astrocytic endfeet (Figure 2)^{[15][16][17]}. There is also

evidence for a change in the phenotype of BTB-associated pericytes, which may show decreased platelet-derived growth factor receptor-b (PDGFR-b) expression in addition to increased desmin expression^[18]. As a result of these changes, the BTB can be, on average, somewhat "leakier" (more permeable) than the normal NVU/BBB in the absence of disease^{[17][19][20]}. The predominant question with regards to BBB breakdown and the treatment of brain tumors has therefore been, is the breakdown of the NVU/BBB in the case of brain tumors significant and uniform enough to allow for the accumulation of efficacious drug concentrations?

Figure 2. The blood-tumor barrier (BTB) is characterized by increased cytokine and VEGF signaling from the tumor, which may lead to decreased expression of tight junction (TJ) proteins like claudin-5. Alterations in pericyte phenotype and disruption of astrocytic associations with endothelial cells may contribute to decreased barrier integrity. However, this is not a uniform phenomenon within or among tumors, and the expression of efflux transporters limits drug permeation into the tumor. Evidence exists showing decreased permeability of the BTB in regions distant to the core of the tumor, which more closely resemble "unaffected" brain.

As this question has been repeatedly investigated, various preclinical tumor models have routinely led to conflicting results. In some cases, tumor vascular permeability, assessed by the accumulation of fluorescent tracers, has been previously correlated with growth patterns, tumor size, or peripheral tumor of origin^[21]. In other cases, including a variety of brain-trophic metastatic breast cancer models developed at the National Institutes of Health (NIH), no correlation between tumor size and permeability has been found^{[14][19]}. These studies also found that the variability of BTB permeability among tumors in the same animal and even among regions of the same tumors, as assessed by the accumulation of fluorescent tracers and small molecules like paclitaxel, doxorubicin, and lapatinib, could be as much as 100-fold^{[14][22][23]}. More recent studies in HER2+ brain-trophic breast cancer metastasis models have shown a poor correlation between drug accumulation and tracer accumulation, as well as inconsistent drug uptake and variable efficacy of biologics like trastuzumab and other antibody-based therapies^{[22][24][25]}. Another model of lung cancer brain metastases found two-fold increases in permeability to small molecules like 3H-mannitol but concluded that this small relative increase in addition to functional P-gp was still a significant limitation to systemic drug therapy^[26]. In addition, a number of studies utilizing transporter-knockout mice and patient-derived xenograft (PDX) models of GBMs and brain metastases have shown that the efficacy of systemic administration of various small molecules is consistently limited by the presence of the NVU/BBB and BTB, active efflux, and the fact that vascular permeability is widely variable within and around the tumor region^{[20][27][28][29][30][31][32][33]}. This heterogeneity in permeability at the BBB leads to wide variability in drug/tracer accumulation and has also been confirmed by elegant correlated ultramicroscopy and MRI techniques in preclinical tumor models^[34]. These studies point to the conclusion that relying on the potential for increased BTB permeability is unlikely to result in efficacious treatment through the systemic administration of novel therapies and their subsequent regulatory approval for such applications.

Although the aforementioned evidence has been largely preclinical, it agrees with clinical observations when considered in the appropriate context. Increased permeability of the BTB, relative to normal brain, is observed clinically, as increased uptake of tracers in magnetic resonance imaging (MRI) and positron emission tomography

(PET) imaging allows for definitive diagnosis of brain tumors and informs many aspects of their treatment^[35]. However, especially in the case of diffuse and invasive tumors like GBM, it has also been shown that nonenhancing, infiltrating regions of brain tumors often exist outside of the region of T1-weighted contrast enhancement^{[36][37]}. This indicates that some portions of the malignant tumor are protected by a relatively uncompromised NVU/BBB. The patterns of treatment failure are strongly correlated with and attributed to these nonenhancing regions, and maximal resection that includes these regions improves survival^{[38][39][40]}. Increasingly, early-phase studies, in which patients receive drugs prior to tumor resection and biopsy, are being utilized to determine the real extent of antineoplastic drug permeability to the BTB^{[1][2]}. Although fold-increases in drug concentrations relative to normal brain may be observed at the core of the tumor, this may still not be adequate to cause cell death. As has been evidenced in many of the aforementioned preclinical models, it is unlikely that these drug concentrations are representative of concentrations in the entirety of the tumor. In the case of GBM, the infiltrative boundaries of the tumor are likely to have a more competent and intact BTB, closer to that of "unaffected brain"^{[20][31][41][42]}. This heterogeneous drug distribution among different regions of the tumor is also clinically evidenced in drug concentrations from biopsies of non-contrast-enhancing tumor regions^[43].

As there has been a great success with novel treatments of peripheral disease, the culmination of decades of brain tumor research has led to the conclusion that it is imperative that molecules and delivery strategies be designed foremost with an intact NVU/BBB in mind. As an example, GNE317, a small molecule that was designed specifically to avoid active efflux, showed significantly higher activity in a model of brain metastases of lung cancer than another counterpart PI3K inhibitor not designed to penetrate the BBB/NVU^{[44][45]}. Other brain-penetrant inhibitors like osimertinib, an EGFR inhibitor, have also shown better preclinical and potential clinical efficacy^{[46][47]}. While designing small lipophilic molecules in an attempt to optimize tumor penetration and minimize active efflux is certainly one potential method towards effective treatments for brain tumors, there are a vast number of other drug delivery strategies and novel molecules in development for this application.

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