

 Received:
 2020.03.04

 Accepted:
 2020.09.12

 Available online:
 2020.10.01

 Published:
 2020.11.27

Authors' Contribution: Study Design A Data Collection B Statistical Analysis C Data Interpretation D Manuscript Preparation E Literature Search F Funds Collection G Role of Exosomes in the Progression, Diagnosis, and Treatment of Gliomas

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Haozhe Piao, e-mail: piaohaozhe@cancerhosp-ln-cmu.com, Xi Zhao, e-mail: xizhaocancerhos@163.com This study was supported by the National Science Foundation of Liaoning Province (No. 20180530059), the National Science Foundation of Liaoning Province (No. 2020-ZLLH-37), Liaoning Revitalization Talents Program (No. XLYC1902023), the Key Research and Development Project of Liaoning Province (No. 2018225040), and the Tumor Mass Spectrometry Project of Liaoning Cancer Hospital and Institute (ZP202002)

Gliomas are the most common primary malignant brain tumors associated with a low survival rate. Even after surgery, radiotherapy, and chemotherapy, gliomas still have a poor prognosis. Extracellular vesicles are a heterogeneous group of cell-derived membranous structures. Exosomes are a type of extracellular vesicles, their size ranges from 30 nm to 100 nm. Recent studies have proved that glioma cells could release numerous exosomes; therefore, exosomes have gained increasing attention in glioma-related research.

Recent studies have confirmed the importance of extracellular vesicles, particularly exosomes, in the development of brain tumors, including gliomas. Exosomes mediate intercellular communication in the tumor microenvironment by transporting biomolecules (proteins, lipids, deoxyribonucleic acid, and ribonucleic acid); thereby playing a prominent role in tumor proliferation, differentiation, metastasis, and resistance to chemotherapy or radiation. Given their nanoscale size, exosomes can traverse the blood-brain barrier and promote tumor progression by modifying the tumor microenvironment. Based on their structural and functional characteristics, exosomes are demonstrating their value not only as diagnostic and prognostic markers, but also as tools in therapies specifically targeting glioma cells.

Therefore, exosomes are a promising therapeutic target for the diagnosis, prognosis, and treatment of malignant gliomas. More research will be needed before exosomes can be used in clinical applications. Here, we describe the exosomes, their morphology, and their roles in the diagnosis and progression of gliomas. In addition, we discuss the potential of exosomes as a therapeutic target/drug delivery system for patients with gliomas.

MeSH Keywords: Biological Markers • Exosomes • Glioma

Full-text PDF:

https://www.medscimonit.com/abstract/index/idArt/924023





e924023-1

e-ISSN 1643-3750 © Med Sci Monit, 2020; 26: e924023 DOI: 10.12659/MSM.924023

Background

A glioma is the most common primary tumor of the central nervous system (CNS); it is characterized by high aggression, relapse, and mortality [1]. Gliomas account for 24.7% of all primary brain tumors and 74.6% of malignant brain tumors [2]. Approximately half of all the gliomas are glioblastoma multiforme (GBM). Patients with gliomas generally undergo conventional treatment, including surgical resection combined with radiotherapy and temozolomide (TMZ)-based chemotherapy. Although the molecular etiology of gliomas has been elucidated and advances have been made in glioma diagnosis and treatment, the prognosis of these patients undergoing treatment remains poor [3].

Extracellular vesicles (EVs) are membrane-bound subcellular organelles that bud from the cell surface and participate in transferring cytoplasmic or membrane-bound biomolecules to the neighboring cells, or the extracellular space [4]. EVs contain proteins, lipids, deoxyribonucleic acid (DNA), messenger ribonucleic acid (mRNA), and multiple noncoding RNAs [4]. EVs are released by several cells and play a key role in intercellular communication by transmitting their molecular contents to different cell types [5]. EVs are currently divided into 4 types: exosomes, microvesicles (microparticles), membrane particles, and apoptotic vesicles; while the terms "ectosomes" and "exosome-like vesicles" have been abandoned because there was insufficient evidence to support their existence [6]. Given their potential as a therapeutic target or a drug delivery system, this review describes the exosomes, their morphology, and their role in the diagnosis and progression of gliomas.

In 1983, Harding et al. [7] and Pan et al. [8] first described peptide-containing vesicles capable of mediating intercellular communication in sheep reticulocytes. These vesicles were termed "exosomes" by Johnstone et al. in 1989. They were eventually isolated and purified from reticulocytes [9]. For a long time, exosomes were considered to be cellular waste [9]. However, they are nanovesicles that are 30 nm to 100 nm in diameter and originate from endosomes. Exosomes are formed from multivesicular bodies and released extracellularly after fusing with the cell membrane. They exist in reticulocytes and other cell types, and can be found in various body fluids, including blood, cerebrospinal fluid (CSF), lymphatic fluid, urine, and pleural and abdominal effusions [10]. They recognize target cell(s), participate in signal transduction, and can transfer their contents into cells by direct fusion with the recipient cell membrane [11]. The biological functions exerted by exosomes vary according to the physiological conditions and their cellular origin [12]. The contents of some exosomes are similar, whereas exosomes from different subclasses specialize in specific biological functions [13]. As exosomes are rich and diverse in their biomolecular contents, can be obtained by minimally invasive techniques, and monitored dynamically, the detection of exosomes and their contents in the peripheral fluids (including blood and cerebrospinal fluid) of patients is an efficient way to diagnose tumors using a fluid biopsy [14]. Tumor cells secrete more exosomes than normal cells [14]. Recent studies have revealed that exosomes bind to receptors on the surfaces of target tumor cells via surface markers [15,16]. Moreover, given their nanoscale size, exosomes can freely pass through the blood-brain barrier (BBB). Therefore, exosomes have gained increasing attention in the diagnosis and treatment of malignant gliomas. In this review, we describe the contents of exosomes and their role in the diagnosis, progression, and potential therapeutic intervention (target/delivery system) of gliomas.

Exosomes

Formation and secretion of exosomes

The formation of exosomes begins with cell membrane invagination forming vesicles that contain cytoplasmic material called early endosomes [17]. Early endosomes mature into late endosomes when the microsomal membrane "sprouts" inward to form intraluminal vesicles that coalesce to form the multivesicular endosome [17]. Subsequently, the multivesicular endosomes employ multiple mechanisms to fuse with the cell membrane after which they are released into the extracellular matrix as exosomes [17]. The secretion of exosomes requires the concerted action of numerous families of proteins. Rab proteins, including Rab11, Rab27, and Rab35, and a soluble N-ethylmaleimide-sensitive-factor (NSF) attachment protein receptor play a key role in exosome secretion [18]. There are other factors, including changes in the intracellular pH and potassium concentration that regulate the formation and secretion of exosomes [19].

Exosome contents

Exosomes are small vesicles (30 nm to 100 nm) that contain lipids, proteins, mRNAs, and micro ribonucleic acids (miRNAs) [20]. The commonly found exosomal proteins include cytoskeletal proteins (tubulin and microfilament-binding proteins), membrane fusion proteins (Rab proteins, ALIX, and flotillin), metabolic enzymes, and channel proteins. Exosomes also contain unique proteins depending on their biological source [21]. Some proteins that are overexpressed in tumor cells can be present in exosomes too, including Fas ligand proteins, tumor necrosis factor (TNF)-related apoptosis-inducing ligands, tumor antigens, and immunosuppressive proteins (transforming growth factor [TGF]- β] [22]. The lipids contained in exosomes include lysophosphatidic acid, cholesterol, and ceramide [21]. Exosomes encapsulate a large amount of RNA and are effective vectors

for RNA transport into target cells; thereby, playing a regulatory role. This encapsulation ensures that the RNA molecules are protected from degradation by the intra- and extracellular RNases. Although numerous studies have shown the presence of protein and RNA in exosomes, few studies report the presence of DNA. However, mitochondrial and chromosomal DNA have been found in exosomes [23]. Exosomes have been demonstrated to contain 100 bases to 17 kilobases of doublestranded DNA [24]. Thakur et al. demonstrated that the majority of exosomes from tumor cells contain genomic doublestranded DNA that reflects the tumor mutation status [24]. However, the mechanism by which DNA is transported into the exosomes remains unclear. Some studies have suggested that a deregulated DNA repair pathway in tumor cells allows DNA to accumulate in the cytoplasm (as compared to healthy cells); thus making DNA available for entry into the exosomes [25].

Morphology and physiological functions of exosomes

The exosomes are circular or cup-shaped vesicles, primarily related to the metabolism and exchange of biomolecules between the cells and body fluids [26]. The donor exosomes can fuse with recipient cells and regulate their intracellular metabolism. Exosomes are considered to be signaling mediators. Specific enzymes within exosomes are responsible for converting their contents into signaling molecules that bind to the target-cell receptors and mediate the intercellular transfer of materials. The internalization of exosomes by target cells involves fusion with the plasma membrane, endocytosis, micropinocytosis, phagocytosis, and a lipid raft-mediated internalization [27]. Exosomes play a vital role in the differentiation and development of neurons, and in tumor growth, invasion, metastasis, angiogenesis, and immunity [26]. Therefore, biomarkers within the exosomes can be used in the diagnosis and treatment of certain diseases, such as miR-21, miR-222, miR-124-3p [28,29].

Isolation of exosomes

Exosomes are found in multiple biofluids [30]. The current methods for exosome isolation are mostly based on their physical properties, including particle size and density; and include ultracentrifugation, density-based separation, precipitation, ultrafiltration, and immunoaffinity. Ultracentrifugation is regarded as the criterion standard for exosome separation because of its large extraction capacity, reliable purity, and its ability to completely satisfy the needs for subsequent biochemical and cellular experiments. However, it is difficult to differentiate between exosomes and other EVs [17]. Recently, lbsen et al. [31] developed an alternating current-based, electrokinetic microarray chip device that requires only 30 μ L to 50 μ L of a plasma sample that can be processed in 15 minutes. Cumba Garcia et al. isolated exosomes from the plasma

of GBM patients using convenient density gradient-based ultracentrifugation to analyze the presence of biomarkers [32]. The presence and purity of exosome preparations can be further assessed by western blotting for exosome-specific markers (CD9, CD63, ALIX, or TSG-101) [32].

Role of Exosomes in the Progression of a Glioma

GBM is a complex morphological and molecular disease with a high degree of heterogeneity, complicating its classification and treatment [1]. Studies on the role of exosomes in the development of gliomas have gained momentum in the last 3 years [29]. Glioma-derived exosomes (GDEs) stimulate angiogenesis, migration [33], glioma cell proliferation [34], modulate tumor invasiveness, and activate signaling. In addition, GDEs participate in the occurrence, development, and therapeutic resistance of gliomas (Table 1).

Exosomes affect the microenvironment around gliomas

The resistance of malignant gliomas to various treatments is primarily attributed to their immunosuppressive tumor microenvironment (TME) that comprises tumor cells, fibroblasts, and immune cells [45]. Cellular communication between the tumor and its environment is mainly mediated by the shedding of membranous microbubbles that can fuse with other cells within the TME [46]. Exosomes play an important role in immunosuppression, stimulation of tumor progression, invasion, metastasis, and multidrug resistance [47,48].

Exosome-mediated angiogenesis

Angiogenesis is a critical event in glioma progression and numerous angiogenic factors in GBM-derived exosomes stimulate angiogenesis [49,50]. The tetraspanins selectively supply exosomes with proteins and mRNA to mediate the exchange of information between the exosomes and the vascular endothelial cells, thus promoting angiogenesis [51]. The GDEs carry proteases associated with malignancies, including MMP-9 and active MMP-2 along with a tissue plasminogen activator, and a high molecular weight urokinase-plasminogen activator, which are essential for angiogenesis. A variety of angiogenic factors (mainly epidermal growth factor receptor variant III [EGFRvIII]) are secreted by gliomas. The EGFRvIII can be "shared" between glioma cells by the intercellular transfer of exosomes via a phosphatidylserine-dependent mechanism [52]. When human astrocytes were incubated with exosomes containing EGFRvIII for 24 hours, the EGFRvIII bound the cell membrane and the phosphorylated extracellular protein kinases [52]. The angiogenesis-related genes and proteins play a vital role in pro-angiogenesis [37]. The U251 human glioma Table 1. Studies on the usability of exosomes in gliomas.

Donor cells	Reference	First author (country)	Year	Recipient cells	Mechanism	Model	Result
Hypoxic U87MG and U251	[35]	Qian M (China)	2019	M2 macrophages	miR-1246 mediates H-GDE-induced M2 macrophage polarization, activates STAT3 signaling, and inhibits NF- kB signaling by targeting TERF2IP	In vitro and in vivo	Promotes glioma cell growth, invasion, and migration
GSC cell line	[29]	Sun Z (China)	2019	U251 and U87 glioma cells <i>in vitro</i> , and inoculated into 5-week-old male BALB/c nude mice	Activates Notch1 signaling	In vitro and in vivo	Promotes glioma cell proliferation, invasion, and neurosphere formation; enhances stemness and tumorigenicity of non-GSC glioma cells
Human GBM cells	[36]	Cai Q (China)	2018	Human glioblastoma cell line T98G cells	Activates STAT3 signaling by targeting CADM1	In vitro	Promotes glioma cell proliferation and metastasis
A172 cells	[37]	Lang HL (China)	2017	HBMECs	Upregulates protein synthesis of bFGF, VEGFA, bFGFR, and angiogenin by exosomes loaded with linc-POU3F3	In vitro	Induces angiogenesis
U87MG-GSCs	[38]	Zhang GB (China)	2017	U87MG; HBMECs	Mediates miRNA transport and facilitates the proliferation of epithelial and glioma cells	In vitro	Promotes glioma cell proliferation
GA-hMSCs	[39]	Figueroa J (USA)	2017	GSCs	Exosomal miR- 1587 downregulates tumor suppressive nuclear receptor corepressor 1	In vitro	Enhances GSCs tumorigenicity, and increases GSCs proliferation and clonogenicity
GSCs from U251 cells	[40]	Sun X (China)	2017	ECs	Stimulates miR- 21/VEGF/VEGFR2 signaling	In vitro	Increases the angiogenic potential of ECs
Glioma cell lines (A172, U87-MG, U251, and T98G)	[41]	Lang HL (China)	2017	ECs	Linc-CCAT2 increases Bcl-2 expression and decreases Bcl2- associated X (Bax) and caspase-3 expression	In vitro	Promotes angiogenesis and decreases apoptosis

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Indexed in: [Current Contents/Clinical Medicine] [SCI Expanded] [ISI Alerting System] [ISI Journals Master List] [Index Medicus/MEDLINE] [EMBASE/Excerpta Medica] [Chemical Abstracts/CAS]

Donor cells	Reference	First author (country)	Year	Recipient cells	Mechanism	Model	Result
U87 cells	[42]	Yang JK (China)	2016	SHG-44 cells	miR-221 inhibits gene DNM3 expression	In vitro	Promotes cell proliferation, migration, and TMZ resistance
UPN933	[43]	Hellwinkel JE (USA)	2016	PBMCs	Reduces T cell activity and migration efficiency	In vitro	Suppresses immune responses
Murine- derived GL26 Cells	[44]	Liu ZM (China)	2013	CD8+ T cells; 6-week-old female C57BL/6 mice	Reduces the number and function of CD8+ T cells	In vitro and in vivo	Promotes GBM growth
U87 cells	[33]	Kucharzewska P (Sweden)	2013	HUVECs, HBMECs, GBM cells; 8-week- old female NOD/SCID mice	Secretes multiple growth factors and cytokines to activate pericyte PI3K/AKT signaling	In vitro and in vivo	Promotes angiogenesis and tumor growth
Human GBM cells	[34]	Skog J (USA)	2008	HBMECs; U87 cells	Modifies the tumor microenvironment, and promotes migration, angiogenesis, and cell proliferation by providing abundant mRNAs	In vitro	Promotes angiogenesis, proliferation, and invasion of glioma cells

Table 1 continued. Studies on the usability of exosomes in gliomas.

bFGF – basic fibroblast growth factor; bFGFR – basic fibroblast growth factor receptor; ECs – endothelial cells; GA-hMSCs – gliomaassociated human mesenchymal stem cells; GBM – glioblastoma multiforme; GSCs – glioma stem cells; HBMECs – human brain microvascular endothelial cells; H-GDE – hypoxic glioma-derived exosomes; HUVECs – human umbilical vein endothelial cells; mRNA – messenger ribonucleic acid; miRNA – microRNA; PBMCs – peripheral blood mononuclear cells; STAT3 – signal transducer and activator of transcription 3; TMZ – temozolomide; VEGFA – vascular endothelial growth factor A.

cells secrete exosomes containing various proangiogenic factors, which promote the proliferation, migration, and lumen formation of endothelial cells (ECs) [47]. Moreover, the GDEs regulate the angiogenic capacity of the ECs by up- or downregulating miRNA expression. Xu et al. reported that the glioma stem cell (GSC)-derived exosomes activate the angiogenic potential of ECs by increasing the levels of miR-21 and the proangiogenic growth factor, vascular endothelial growth factor (VEGF) [40]. Lang et al. reported that glioma cells stimulate angiogenesis in ECs mainly via the linc-POU3F3-containing exosomes [37]. The GDEs contain miR-296, miR-29a, and miR-30e to regulate the ECs-mediated angiogenesis. The exosomes derived from GSCs or glioma cells enhance tube formation and migration of the ECs, indicating that the malignant cells affect the function of the surrounding cells [40]. Skog et al. reported that the tubule length of the human brain microvascular ECs (hBMECs) doubles within 16 hours in the presence of the exosomes derived from GBM, and the ECs in the brain microvasculature internalize the GDEs that are responsible for tubule formation and angiogenesis [34]. Lang et al. found that glioma cells stimulate angiogenesis by transferring the long intergenic noncoding-RNA CCAT2 (linc-CCAT2) to the ECs via the exosomes, suggesting that they are the potential therapeutic targets in gliomas [41]. Monteforte et al. reported that the exosomes from GBM induce angiogenesis during peripheral ischemia, and contain proteoglycans and small RNAs that promote revascularization [53]. Thus, exosomes can be the potential therapeutic targets for multiple cancers associated with enhanced capillary formation.

Exosomes and glioma hypoxia

Hypoxia is a hallmark of the GBM microenvironment; it regulates the transcriptome and proteome of the tumor cells [50]. The exosomes play a prominent role in the adaptation of tumor cells to hypoxia, which is related to angiogenesis, tumor progression, and metastasis. Hypoxic stress modulates the physiological activity of the surrounding or distant cells; thus,

qualitatively and quantitatively altering the protein content of the GBM cell-derived exosomes. A recent study has revealed that hypoxic exosomes regulate the glioma cells by upregulating the transcription of small nucleolar RNA, C/D box 116-21, and downregulating the potassium voltage-gated channel subfamily J member 3 [50]. The protein cargo in exosomes differentially regulates the expression of certain genes. The hypoxic status of the glioma cells is mirrored in their cargo [33]. King et al. reported that in response to hypoxia, cancer cells improve their survival and invasiveness by enhancing the secretion of exosomes into the microenvironment [54]. In addition, the exosomes secreted by the hypoxic cancer cells promote tumor progression in the normoxic areas [48]. The hypoxia regulates the exosomal miRNA content and release [54-56]. A tube formation assay with endothelial progenitor cells indicated that compared to normoxic exosomes, the hypoxic exosomes enhanced more features associated with angiogenesis [50]. The GDEs contain various mRNAs and proteins that regulate hypoxia; this is often associated with poor prognosis, and the resistance of gliomas to radiotherapy and chemotherapy [57]. Kucharzewska et al. revealed that the GBM cell-derived hypoxic exosomes regulate ECs by secreting various growth factors and cytokines that stimulate pericyte PI3K/AKT signaling and migration [33]. The GBM-derived exosomes show enhanced autocrine and promigratory functions under hypoxic conditions. Thus, the hypoxic exosomes play a vital role in tumor vascularization, pericyte vessel coverage, and GBM cell proliferation [33]. Qian et al. found that hypoxic GDEs (H-GDEs) promote glioma progression and facilitate the formation of an immunosuppressive microenvironment by inducing M2 macrophage polarization [35]. The H-GDEs are potent stimulants of myeloid-derived suppressor cells that regulate the formation of an immunosuppressive environment while evading host immunity against tumors. Guo et al. demonstrated that hypoxia upregulates the miR-10a and miR-21 contents in the GDEs, and activates myeloid-derived suppressor cells via the RORα/IκBα/NF-κB and PTEN/PI3K/AKT pathways [58]. miR-29a and miR-92a in the H-GDEs mediate the differentiation of functional myeloid-derived suppressor cells [59]. Thus, the H-GDEs play an important role in the development of gliomas.

Role of exosomes in the proliferation and invasiveness of gliomas

Proliferation and invasion are regulated by the exosomes and are important for glioma cell survival and glioma recurrence [60]. The GDEs are indispensable in the regulation of tumor proliferation, survival, and invasion due to their coding and noncoding RNA contents. Cai et al. [36] reported that the GBM progression is promoted by exosomal miR-148a (which targets CAMD1) and increased signal transducer and activator of transcription 3 (STAT3) activation. The exosomes derived from glioma-associated-human mesenchymal stem cells (GA-hMSCs) contain miR-1587, which targets the tumor suppressor gene NCOR1 in GSCs; thereby altering the biological functions of the GSCs and increasing tumor cell proliferation and colony formation [39]. The GSC-derived exosome-mediated miRNA transport facilitates the proliferation of epithelial and glioma cells [39]. Further, exosomes secreted by the GSCs mediate the dedifferentiation of non-GSC glioma cells into GSCs via the Notch signaling pathway, and enhance the stemness and tumorigenicity of non-GSC glioma cells [29,38].

Integrins, adhesion molecules, and other extracellular matrix factors play a prominent role in regulating cell migration. The GDEs contain the adhesion/recognition molecule L1CAM. A disintegrin and metalloprotease 10 (ADAM10) catalyze the proteolysis of L1CAM to release its extracellular domain, which allows an interaction with the fibroblast growth factor receptor, and integrins $\beta 1$ and $\alpha 5$ on the target cells. This results in the activation of the integrin ligase, focal adhesion kinase (FAK); upregulation of PI3K/AKT, and NF-kB signaling to induce cell migration [61]. The NEDD4 family interacting protein 1 (NDFIP1) promotes the cellular uptake of PTEN-containing exosomes; thereby downregulating the PI3K/AKT signaling. The NDFIP1 expression is suppressed in a variety of tumors, including gliomas, resulting in a decrease in the uptake of PTEN-containing exosomes to maintain PI3K/AKT signaling, and promoting tumor survival and proliferation [62]. A study by Pinet et al. revealed that silencing the chitinase protein (YKL-40) led to downregulation of the tyrosine kinase receptor B (TrkB), sortilin, and p75^{NTR}, resulting in low invasiveness [63]. The migration and activation of the YKL-40-inactivated cells could be rescued by the TrkB released from the exosomes secreted by the glioma cells. Arscott et al. reported that therapeutic radiation affects the production of exosomes in a dose- and time-dependent manner; radiation-induced exosomes are more likely to be absorbed in cocultured cells [64]. Moreover, the activation of TrkA and FAK signaling enhanced cell migration by changing the molecular composition of the exosomes. Thus, exosomes regulate intercellular signaling upon stimulation by radiation.

Role of exosomes in resistance to glioma treatment

Exosomes play an important role in the treatment resistance exhibited by gliomas as they can export therapeutics from tumor cells [65]. Exosomes can promote the formation of fibroblasts causing fibroblastic reactions, which form a barrier to antitumor drugs. However, exosomes can also transform drugsensitive tumor cells into drug-resistant tumor cells via biomolecules, such as the miRNA [66,67]. Hypoxia promotes the resistance of gliomas to chemotherapy by inducing specific signaling pathways depending on the uptake of exosomes [68]. Moreover, it has been suggested that the GDEs regulate the surrounding cells and create an optimal environment for glioma cell growth [69]. O-6-methylguanine-DNA methyltransferase (MGMT) and alkylpurine-DNA-N-glycosylase (APNG) are essential for repairing TMZ-induced DNA damage; their expression negatively correlates with therapeutic efficacy [70]. Shao et al. reported that in TMZ-resistant cell lines, the exosomal mRNA levels of MGMT and APNG increased rapidly upon TMZ treatment; whereas in TMZ-sensitive cell lines, they decreased, indicating that the exosomal transcript level of MGMT can predict the efficacy of TMZ in GBM patients [70]. The APNG promotes chemoresistance to TMZ in patients with GBM, and the coexpression of APNG with MGMT further enhances drug resistance [71]. Garnier et al. reported that exosome-transcribed MGMT mRNA is released by the GSCs and reflects the resistance to TMZ [72]. Moreover, Yu demonstrated that the exosomal MGMT mRNA levels of reactive astrocytes are significantly higher than those of nonreactive astrocytes. The reactive normal human astrocytes enhance recipient-cell chemoresistance to TMZ by transferring exosomal MGMT mRNA to the surrounding glioma cells; thereby suppressing apoptosis [68]. Zeng et al. reported that the protein tyrosine phosphatase receptor type Z1 (PTPRZ1)- receptor tyrosine kinase (MET) fusioncontaining GDEs increase the resistance of GBM to TMZ and regulate the microenvironment by targeting multiple pro-oncogenic effectors in the recipient cells. Thus, the PTPRZ1-MET and gene fusions in exosomes play a crucial role in GBM invasiveness [73]. The exosomal miRNA-221 in the blood of glioma patients positively correlates with the malignant grade and resistance of gliomas to TMZ [42,74].

Exosomes mediate immune responses

Exosomes play an important role in the communication between tumor and immune cells. The exosomes secreted by tumor cells carry tumor-specific antigens that activate and inhibit the immune system (under specific conditions) and promote GBM proliferation, invasiveness, and chemoresistance. The differentiation and function of the immune cells can be regulated by miRNAs released from the tumor-derived exosomes [75]. These tumor-derived exosomes have multiple physiopathological roles and act on various immune cells, including effector T cells, naturally occurring Treg cells, and natural killer (NK) cells, which are involved in immune suppression and tumor progression [51,76]. Numerous immunomodulatory molecules can be found in the serum exosomes isolated from GBM patients, including antigen-presenting molecules, TGF-β, tumor antigens, and immune intracellular adhesion molecules [77]; indicating that tumors exert immunosuppressive activity across the CNS barrier. Moreover, Harshyne et al. reported that the serum exosomes from GBM patients drive M2 polarization in normal monocytes, indicating a bias towards the reactivity of T helper 2 cells (Th2). The Th2 responses are considered undesirable in tumor immunotherapy since they alleviate cytotoxic antitumor immune mechanisms and help escape cell-mediated immunity [78]. Moreover, microglia-derived exosomes

mediate immune responses important for inflammation, degeneration, and tumorigenesis in the CNS [79]. Dendritic cells (DCs) are professional antigen-presenting cells. Upon incubation with the GDEs, they induce T cell activation and mediate cytotoxicity against gliomas in vitro [80]. Hellwinkel et al. [43] demonstrated that immunosuppressive phenotypes and high cytokine concentrations in the tumor-derived exosomes lead to a decreased expression of other cytokines, including interleukin 2 (IL-2) and CD69, and T cell activity; consequently hindering lymphocyte migration and inhibiting tumor immunity. The exosomes derived from the GBM GL26 cells reduced the number and function of cytotoxic CD8⁺ T cells, stimulating tumor growth [44]. The GDEs help differentiate peripheral bloodderived monocytes into alternately activated M2 tumor-supporting macrophages [81], and regulate the cytokine output and migratory ability of mitogen-stimulated, healthy, peripheral blood mononuclear cells [43].

An immune system evasion is the hallmark of glioma development and progression. The GSC-derived exosomes promote evasion from tumor immunity via intercellular communication. Yu et al. [82] demonstrated that the GSC-derived exosomes induce peripheral T cell immunosuppression. This involves active monocytes and the conversion of peripheral T cells to the monocytic, myeloid-derived suppressor cell-induced tumorigenic phenotype. Since exosomes promote tumor progression by mediating intercellular communication, they can be used as a tool to program anticancer immunity and restore the effector function of immunosuppressive cells. Konrad et al. demonstrated that mononuclear cells regulate the M2 immunosuppressive phenotype and the expression of the programmed death ligand 1 by specific and rapid uptake and release of the GSC-derived exosomal biomolecules. This includes STAT3; a vital molecule in the tumor-mediated immune suppression [83]. In conclusion, the exosomes derived from GSCs are important regulators of the immunosuppressive TME in patients with GBM [83].

Exosomes in the Diagnosis and Prognosis of Gliomas

Neuroradiology has an important role in the diagnosis and treatment of gliomas. Unlike other tumors, the circulating biomarkers (circulating tumor cells and cell-free nucleic acids) are of little use in the diagnosis of brain tumors because of the BBB. Other potential biomarkers have been identified in the serum and CSF of glioma patients; however, no substantial progress has been made in their clinical application [84]. Since the GDEs carry abundant genetic information from the tumor cells, they are promising biomarkers in clinical diagnosis. The GDEs are a better diagnostic sample than other currently used samples (urine or serum) because of the specificity and sensitivity of detection. The type, origin, differentiation, and genotype of a malignant glioma can be determined by studying exosome profiles and predicting the therapeutic response and prognosis [38]. Furthermore, exosomal cargo (mRNAs, miRNAs, proteins, and DNA) has helped identify potentially important biomarkers applicable in clinical diagnosis. Since it is difficult to obtain sufficient tissue samples for diagnosis, combining the clinical features and the genetic information from exosomes can provide a more accurate diagnosis.

mRNA

Tumor-specific RNA in serum exosomes can be used to detect disease progression, and predict the prognosis of tumors in the CNS [20]. Chen et al. [85] detected mutant IDH1 mRNA transcripts in exosomes from the CSF of glioma patients, which also revealed that the exosomes from the CSF of glioma patients contain mutant IDH1 transcripts. Exosomal mRNA levels of MGMT and APNG can be used to monitor the therapeutic efficacy of TMZ in glioma treatment [70]. Yang et al. found that the genes p65 and dynamin 3 (DNM3) were upregulated in exosomes from the brain and blood of patients with primary and recurrent GBM by a xenograft orthotopic mouse model, suggesting that the exosomal levels of Specific transcripts may be potential clinical diagnostic markers for GBM [30].

miRNAs

miRNAs are small (~22 nucleotide long) noncoding singlestranded RNAs involved in the regulation of post-transcriptional gene expression. The dysregulated miRNAs affect cellular physiology and development, leading to disease states [86]. Wei et al. showed that the exosomes isolated from GBM patients contain a large pool of miRNAs [87]. Since miRNAs are relatively stable within the exosomes and exhibit differential expression based on the tumor type, they can be used as biomarkers for the diagnosis and prognosis of tumors [88]. Santangelo et al. reported that compared with healthy controls and low-grade glioma patients, the serum exosomes from high-grade glioma patients contained significantly higher levels of miR-21, miR-222, and miR-124-3p, which decreased rapidly after neurosurgical therapy [89]. The GDE contents of miR-21, miR-222, and miR-124-3p can be determined using minimally invasive methods that can help diagnose brain tumors, predict the pathological grading of gliomas, and preoperative metastases. However, there is a need for more prospective studies to confirm whether the changes in serum exosomal-miRNA concentrations in glioma patients reflect tumor recurrence after surgery, and help devise personalized treatment options.

In 2014, Manterola et al. [90] analyzed the miR-574-3p, miR-320, and RNU6-1 contents in the serum exosomes isolated from GBM patients and healthy individuals; RNU6-1 alone or in combination with numerous miRNAs could specifically diagnose GBM. The expression of miR-301a was markedly increased in the serum exosomes from GBM patients when compared with those of healthy individuals, and correlated with the pathological grade and Karnofsky performance score of GBM patients [91]. The postsurgical levels of miR-301a secreted from the serum exosomes of glioma cells were remarkably decreased and associated with an increase in tumor recurrence. This indicates that exosomal miR-301a levels reflect the actual status and pathological changes of the glioma. Transcription polymerase I and releasing factor (PTRF, also known as Cavin-1) is stably detected in glioma tissues, serum and CSF samples. PTRF overexpression induced malignancy in the surrounding cells by regulating the TME and leading to a poor prognosis for the glioma patients; whereas downregulation of PTRF reversed the malignant phenotype of GBM, indicating that PTRF is an ideal diagnostic and prognostic indicator, and a promising therapeutic target against GBM [92]. Murgoci et al. reported high levels of miR-221 in the exosomes derived from the serum and CSF of GBM patients, suggesting its potential as a GBM biomarker [79]. miR-21 is the most consistently overexpressed miRNA in malignant gliomas and regulates glioma cell invasiveness, migration, and the grades of gliomas [55,93,94]. The amount of miR-21 in the exosomes isolated from the CSF of GBM patients was 10-times higher when compared with that in the exosomes from healthy subjects [93]. The miR-21 levels in the serum exosomes from GBM patients were 40-times higher than those from the serum exosomes of healthy individuals [34]. The receiver-operating characteristic curve analysis revealed the high efficacy (area under the curve: 0.927, 95% confidence interval, 0.865 to 0.985) of using exosomal miR-21 in the diagnosis of gliomas [94]. Together, these findings indicate that only a small amount of serum or CSF is required for the extraction of exosomes to determine the diagnosis, grading, and prognosis of gliomas based on exosomal miR-21. Recent research has shown that miR-1246, the most abundant miRNA in H-GDEs mediates H-GDE-induced M2 macrophage polarization, and the miR-1246 levels decreased after surgery. The exosomal miR-1246 in the CSF of GBM patients activates STAT3 signaling and inhibits NF-kB signaling by targeting the telomeric repeat-binding factor 2-interacting protein (TERF2IP), and serves as an innovative biomarker for the diagnosis of GBM [35]. The exosomal levels of miR-10b and miR-143/145 also correlate with the prognosis of gliomas [95]. Some miRNAs, such as miR-128 and miR-34a, negatively correlate with the increasing malignancy of gliomas [96]. Therefore, the exosomal miRNAs are excellent candidates as biomarkers because of their stability, ease of collection and detection, and tissue and cell specificity. They can be used for early diagnosis of malignant gliomas with accurate stratification. These features make exosomal miRNAs suitable biomarkers for patients who cannot undergo surgery or are suspected of relapse. Fluid biopsies using exosomes can be a promising approach for the development of precision medicine in the future.

Protein

The protein contents in exosomes correlated with the malignancy of gliomas [97], indicating their importance in the diagnosis and prognosis of gliomas. Shao et al. [98] reported that 90% of patients with GBM had at least 1 protein from either EGFR, EGFRvIII, podoplanin, and IDH1 differentially expressed in their exosomes. EGFRvIII exists in 25% of glioma patients and can be detected with sensitively and specificity [34]. However, EGFRvIII cannot distinguish between the different grades of gliomas, since it is found only in high-grade gliomas. The sensitivity of EGFRvIII detection depends on the tumor size, location, serum or CSF content, and method of RNA extraction [34]. Setti et al. reported that the chloride intracellular channel 1 detected in exosomes promotes the proliferation and invasiveness of GBM, and is related with poor prognosis [99]. Phillip et al. found that glioma patients release CD9+/GFAP+/SVN+- and CD9+/SVN+-containing exosomes into circulation; the quantities decrease significantly following anti-survivin immunotherapy. This could be related to longer progression-free survival [100]. Currently, the detection of changes in the protein contents of body fluids or tissues is the most commonly used diagnostic method for the diagnosis, treatment, and prognosis of gliomas.

DNA

DNA has been detected in the exosomes obtained from the serum of pancreatic, and prostate cancer patients. García-Romero et al. successfully detected mutations in gene IDH1 (R132H) in the peripheral blood exosomes of patients with high-grade gliomas; consistent findings were obtained from the immunohistochemistry of surgical specimens [101]. However, there are only a few reports on the presence of DNA in exosomes from the body fluids of glioma patients, and the accuracy of exosomal DNA detection in disease prognosis remains to be determined [101].

Exosomes in the Treatment of Gliomas

Compared with the rapid developments in the diagnosis and characterization of gliomas, there has been very little progress in the therapeutic interventions for glioma treatment. Therefore, it is imperative to develop new treatment strategies. Currently, the clinical application of exosomes can be divided into 3 categories: exosomes as therapeutic agents, exosome-mediated drug delivery, and exosome-based immunotherapy. Exosomes have several advantages as therapeutics: Good specificity (exosomes can transport their contents to specific targets via their surface molecules and homing properties); safety (self-derived exosomes are not toxic to the immune system, and are histocompatible); stability (exosomes are bilayered phospholipid membrane vesicles circulating in the blood, their contents are protected by RNase III and prions, and can be transported to remote target cells); and their nanoscale size reduces their clearance from the mononuclear phagocyte system [102]. These characteristics allow the use of exosomes as carriers for drugs and vaccines; this is important for the treatment of malignant gliomas (Table 2). There have been multiple *in vivo* and *in vitro* studies on the use of exosomes in the treatment of gliomas, which are discussed below.

Exosomes as therapeutic agents

Multiple studies have shown that cell-derived exosomes have potential in disease treatment. NK cell-based immunotherapy is a low-toxicity alternative to chemotherapy and radiotherapy. The NK cell-derived exosomes target GBM cells and play an antitumor role. However, the underlying mechanism remains unclear [104,112]. Hao et al. [107] found that exosomes from the human umbilical cord-derived mesenchymal stem cells (hUC-MSCs) exert partial antitumor activity by regulation of the miR-10a-5p/PTEN signaling. Therefore, this axis may be a possible target for early diagnosis and treatment of gliomas. Combining gene-engineered exosomes with other drugs, including temozolomide, can improve the therapy against GBM cells and make it more effective. The exosomes derived from the embryonic stem cells that were conjugated with a tumortargeting c(RGDyK) peptide and loaded with paclitaxel showed enhanced targeting and better curative efficacy of the paclitaxel towards the GBM cells [113]. To obtain glioma-targeting exosomes, Gang et al. loaded superparamagnetic iron oxide nanoparticles with curcumin and conjugated them with a neuropilin-1-targeted peptide by click chemistry for enhanced stability and biocompatibility [114]. This approach yielded good exosomal targeting towards glioma cells, and could be used in the early diagnosis and estimation of drug efficacy against GBM. Numerous studies are being conducted on the chemical modification and genetic engineering of exosomes to improve the BBB permeation; however, the practicality of this approach remains unclear.

Focused ultrasound (FUS) is a noninvasive technique that transiently and reversibly disrupts the stability of the BBB. Chen et al. demonstrated that the BBB could be "opened" by FUS to increase the delivery of IL-12 to brain tumors [115]. Given the lack of specific targeting, the number of natural exosomes that can cross the BBB are inadequate. Bai et al. developed a natural and secure transportation system using FUS to increase the targeted delivery of exosomes for glioma treatment [116]. FUS can be used with blood serum-derived exosomes to inhibit the growth of a glioma without obvious side effects, indicating that this combination is potent for developing glioma therapeutics. Murgoci et al. used a 3D spheroid glioma model to show that microglia-derived exosomes Table 2. Exosomes in the treatment of gliomas.

Donor cells	Reference	First author (country)	Year	Recipient cells	Exosomal content	Mechanism	Model	Result
HEK-293T cells	[103]	Monfared H (Iran)	2019	U87-MG, C6 and rat model	miR-21 sponge construct	Downregulates miR-21, and upregulates miR-21 target genes (PDCD4 and RECK)	<i>In vitro,</i> and stereota- xically injected into a rat model	Reduces tumor volume
hBMSCs	[104]	Yu L (China)	2019	U251	miR-199a	Downregulates AGAP2	In vitro	Suppresses tumor proliferation, invasion, and migration
hBMSCs	[105]	Wang B (China)	2019	GBM cells (T98G, LN229 and A172) and nude mice	miR-34a	Downregulates MYCN	<i>In vitro</i> , and subcuta- neously injected into a rat model	Suppresses GBM cell growth, invasion, migration, and tumorigenesis, and enhances chemosensitivity of the GBM cells to TMZ
hBMSCs	[106]	Deng SZ (China)	2019	Glioma cells (SHG44, C6, U87, and U251) and nude mice	miR-375	Suppresses SLC31A1	<i>In vitro</i> and a rat model	Promotes apoptosis and suppresses proliferation, migration, and invasion
hBMSCs	[107]	Hao SC (China)	2019	U87 cells	Long noncoding RNA PTENP1	Regulates PTENP1/miR- 10a-5p/PTEN signaling	In vitro	Induces cell apoptosis and inhibits glioma cell proliferation
Rat microglia cells	[79]	Murgoci AN (France)	2018	C6 cells	Not available	Not available	In vitro	Suppresses the invasiveness of glioma cells
DCs carried CRCLs with GL261 glioma cells	[108]	Bu N (China)	2015	6-week- old female C57BL/6 mice	CRCLs	Modulates Cbl-b and c-Cbl signaling	In vivo	Prolongs the survival of mice with tumors and suppresses tumor proliferation
MSCs	[109]	Katakowski M (USA)	2013	9L glioma cells; male Fischer rats	miR-146b	Inhibits EGFR expression	<i>In vitro</i> and <i>in vivo</i> rat models	Inhibits glioma growth
hBMSCs	[110]	Munoz JL (USA)	2013	U87 cells; T98G cells	Anti-miR-9	Reduces miR- 9 content in GBM cells and expression of the drug transporter MDR1	In vitro	Reduces the chemoresistance of GBM cells to TMZ

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Donor cells	Reference	First author (country)	Year	Recipient cells	Exosomal content	Mechanism	Model	Result
Human GBM cells	[80]	Bu N (China)	2011	CTLs obtained from PBMCs	Tumor antigen	Activates glioma- specific CD8+ CTLs	In vitro	Kills the autologous glioma cells
GL26 cells	[111]	Zhuang X (China)	2011	Microglial cells	Cucurbitacin I	Selectively inhibits the activity of STAT3 and reduces the expression of IL-1β and IL-6	<i>In vivo</i> by the intranasal route	Promotes tumor cell apoptosis and inhibits tumor cell growth
SMA560vIII	[77]	Graner MW (USA)	2009	VM/Dk mice	None	Produces immune memory, improves T cell activity, and stimulates antibody production	<i>In vivo</i> by the subcuta- neous route	Exosome vaccines enhance immune responses; reduces tumorigenesis and prolongs survival

Table 2 continued. Exosomes in the treatment of gliomas.

CRCLs – chaperone-rich cell lysates; CTLs – cytotoxic T lymphocytes; DCs – dendritic cells; EGFR – epidermal growth factor receptor; GBM – glioblastoma multiforme; hBMSCs – human bone marrow-derived mesenchymal stem cells; MSCs – mesenchymal stem cells; MYCN – basic helix-loop-helix protein 37; PBMCs – peripheral blood mononuclear cells; RNA – ribonucleic acid; STAT3 – signal transducer and activator of transcription 3; TMZ – temozolomide.

suppress tumor invasion in a time-dependent manner, and suggested their potential as nanotherapeutic agents against glioma cells [79].

Exosome-mediated drug or gene delivery

To exploit the exosome potential in clinical applications, an increasing number of studies have focused on utilizing the nanoscale characteristics of exosomes to package and deliver bioactive substances, such as small molecules (anticancer drugs), proteins, and miRNAs [117].

Drug delivery

As exosomes are EVs that originate from multivesicular bodies, they do not trigger adverse immune responses, and are not recognized by the complement system. They are regarded as an innovative delivery system because of their properties of low toxicity, low immunogenicity, excellent biocompatibility, and transport of specific molecules and surface markers [118]. Previous studies have demonstrated the exosome-mediated transfers of lipophilic and hydrophilic drugs, including curcumin [119] and doxorubicin [120]. The current obstacles seem to be the lack of understanding about penetration of the BBB for delivering drugs into the CNS and striking a balance between the blood concentration of the brain parenchyma, and effective therapeutic drug concentration. As exosomes are nanoscale in size, they can efficiently penetrate the BBB via surface proteins that have a donor cell signature [121]. This has been achieved using the exosomes containing metalloproteinases in combination with tumor necrosis factor alpha (TNF- α), interferon γ (IFN- γ), and IL1- β , which promoted the degradation of the extracellular matrix and nitric oxide production, participated in leukocyte or T cell-mediated permeation, and disrupted the BBB [122]. Grapp et al. showed that fusion of the exosomal surface folate receptor- α with the receptor cell membrane helped exosomes permeate the BBB allowing exosome entry into the brain parenchyma from the CSF [123].

The BBB permeability poses a critical obstacle for the delivery of chemotherapeutic drugs. The properties of exosomes make them suitable for the treatment of CNS tumors, and exosomebased combined drugs can overcome the issues of current treatments for GBM [124,125]. There are 3 techniques used to conjugate exosomes with drugs: incubation, the easiest method involves co-incubation [126]; electroporation, which involves the penetration of exosomes using short, high-voltage pulses [127]; and sonication, drugs (such as paclitaxel) can be loaded into exosomes by effective sonication [126]. Alvarez-Erviti et al. [128] injected rabies viral glycoprotein-targeted exosomes intravenously into wild mice; the exosomes permeated the BBB and released glyceraldehyde-3-phosphate dehydrogenase (GADPH)-targeting carrier small interfering RNAs (siRNAs) to the neurons to specifically knock down neuronal GAPDH. The ECs are an important part of the BBB. The exosomes derived from ECs contain various receptors, including the transferrin receptor (TFRC), low density lipoprotein receptor (LDLR), insulin receptor (INSR), and transmembrane protein 30A (TMEM30A). The EC-derived exosomes possess transmembrane activity that could transport substances across the BBB, and enable signaling between the nerve cells and the extracellular material [129]. The exosomal surface derived from glioma cells comprises the pro-permeability protein semaphorin 3A. It disrupts the endothelial barrier in an Nrp1-dependent manner [130]. Consequently, the exosomes can mediate communication across the BBB. Yang et al. [124] demonstrated a decrease in the fluorescent intensity of xenotransplanted zebrafish cancer models and tumor growth markers upon the delivery of anticancer drugs by the brain ECs-derived exosomes. STAT3 is an important protein in multiple malignant tumors, including GBM. Zhuang et al. [111] intranasally administered exosomes containing the STAT3 inhibitor cucurbitacin I into mice with GBM, which significantly prolonged the survival of these mice from 20 days to 44.5 days without adverse reactions or behavioral abnormalities. Salarpour reported that the loading capacity of exosomes depends on the technique used, and chemotherapeutics exhibit a stronger cytotoxicity when encapsulated into exosomes [118]. Therefore, exosomes are promising agents for delivering drugs against brain tumors.

Gene delivery

miRNAs are promising candidates for GBM therapy. However, the challenges include identifying the most effective miRNAs and the delivery method [131]. Exosomes are good carriers of miRNAs that exert anticancer activity against multiple cancer types. The exosomes secreted by the miRNA-transfected mesenchymal stem cells (MSCs) have potent efficacy in treating gliomas [132]. Lang et al. [133] isolated the exosomes derived from miR-124a-expressing (via lentiviral expression) MSCs that released miR-124a, which could be used as therapeutic agents against gliomas. A study by Yu et al. revealed that the exosomes derived from MSCs transport miR-199a to the glioma cells to downregulate AGAP2 and inhibit glioma cell proliferation, invasion, and metastasis [104]. The overexpression of miR-199a in MSCs improved the chemosensitivity to TMZ, and suppressed glioma cell proliferation. Similarly, Munoz et al. demonstrated that the anti-miR-9-delivering exosomes increased the multidrug transporter expression and sensitivity to TMZ in drug-resistant GBM cells, resulting in higher rates of cell death and caspase activity [110]. The miRNA-146bcontaining exosomes derived from MSCs inhibited GBM cell growth [109]. miR-34a released by the exosomes derived from hBMSCs suppressed MYCN expression inhibiting glioma cell growth, invasion, and metastasis, and improving the sensitivity of GBM to TMZ [105]. miR-375 secreted by the hMSC-derived exosomes suppressed SLC31A1 expression and increased apoptosis, while inhibiting glioma cell proliferation, migration, and invasion [106]. Ma et al. showed that the GDEs increased glycolysis and tumor-like transformation of the hBMSCs, indicating that interrupting the interaction between exosomes and MSCs in the TME shows promise as therapy against gliomas [134]. miRNA inhibition using sponge-like constructs is also used as a treatment strategy. Sponge constructs are designed to bind corresponding miRNA(s) or their seed sequences, thus preventing miRNAs from binding to their biological targets [135]. Hamideh et al. reported that exosomes containing an miR-21 sponge decreased the tumor volume in a rat model of GBM; thus, miR-21 sponges seem to offer a promising therapeutic approach [103]. Exosome-mediated intercellular signaling during tumor proliferation is a potential therapeutic target [38]. The exosomal long noncoding-RNA antisense transcript of the hypoxia-inducible factor-1 α promoted GBM progression and radioresistance by downregulating the HIF- 1α expression; hence, it is a potential therapeutic target in GBM [136]. Immature mouse DCs show promise in the production of exosomes with low toxicity and immunogenicity. Tian et al. found that exosomes modified by targeting ligands improved the tumor targeting of doxorubicin, resulting in a stronger suppression of tumor proliferation without toxicity [120]. Sugahara et al. fused an αv integrin-specific iRGD peptide to the surface of the exosomes derived from engineered immature mouse DCs, which facilitated tumor targeting and the absorption of anticancer drugs [137].

Numerous techniques have been applied to producer cells to secrete exosomes with specific cargo. However, low exosome abundance is one of the main obstacles in developing exosome-based treatment. Li et al. reported that small noncyto-toxic pharmacological molecules, including MOPIPP and vacuolin-1, promote the production of exosomes and can be used to treat GBM [138]. Other current challenges associated with exosome-based treatment are donor-cell selection, drug encasement into exosomes, and drug resistance [14]. Therefore, miRNAs can be efficacious treatment agents for multiple diseases, and for the delivery of different synthetic molecules and biomolecules in cell therapy.

Exosome-based immunotherapy

Lymphocytes could enter the CNS when the BBB was damaged by brain tumors [139]. Immunotherapy as a treatment for gliomas has recently gained importance. Exosome-mediated immunotherapy is an efficient treatment strategy [108]. The DCs are the most efficient antigen-presenting cells that stimulate a strong immune response. The DC-derived exosomes

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can enhance antitumor immunity in vitro and in vivo [140]; tumor vaccines can be designed based on this information [141]. Immunotherapy for gliomas has recently gained importance as malignant glioma cell growth is influenced by the immune system. The GDEs are antigen transporters that once loaded into the DCs activate the CD8⁺ cytotoxic T lymphocytes (CTLs) and kill autologous glioma cells [80]. Positive results for the DCbased vaccine immunotherapy for GBM have been obtained in vitro, in vivo, and in early-phase clinical trials [142,143]. However, the overall efficacy of DC vaccines remains unclear. Therefore, there remains an urgent need to improve immunotherapy [144,145]. Ning et al. found that the DC-derived exosomes containing chaperone-rich cell lysates stimulate stronger and more effective antitumor T cell immune responses by regulating Cbl-b and c-Cbl signaling [108]. Liu et al. demonstrated that the α -Galcer-containing tumor-derived exosomes constitute efficacious DC-based vaccines. Alpha-Galcer-activated invariant NK-T cells were used as a cellular adjuvant, thus breaking immune tolerance and inducing an antigen-specific CTL response against the GBM cells [146]. The GDEs can be used as a storage pool for tumor antigens, which can overcome the limitations associated with immunotherapy against malignant gliomas to some extent. For example, phenotypic changes can be observed in patients with progressing gliomas by extracting exosomes, which helped to develop a vaccine that stimulates antitumor immunity in the host [147]. Using animal experiments, Graner et al. [77] showed that the exosome-based vaccines induced strong antitumor immunity via humoral and cellular pathways, and prolonged the survival of mice with gliomas. Changes in the total exosomal protein and mRNA contents in glioma patients treated with antitumor vaccines can be used to monitor their immune and clinical responses. Muller et al. [148] showed that the exosomal immune-related protein content positively correlates with the different grades of gliomas, indicating the significance of immune-related gene expression in serum exosomes in the response of glioma patients to these vaccines.

Conclusions

Since the discovery of exosomes, their composition, functions, and correlations with various diseases have been widely studied. Their involvement in gliomas has gained attention over the years. Exosomes are involved in the immune evasion, tumor hemodynamic remodeling, and progression of gliomas. Exosomes contain a variety of biomolecules that can serve as biomarkers for the diagnosis [149]. Since exosomes are detected in body fluids such as serum or CSF, they allow for minimally invasive diagnosis and treatment of gliomas. Malignant gliomas secrete an abundance of exosomes that enter the peripheral fluid through the BBB. They contain various bioactive antigens that are tumor-specific and participate in the onset and development of gliomas. Therefore, exosomes have been speculated to be promising candidates for the diagnosis and treatment of gliomas. Exosomes are nanovesicles that exhibit targeted homing, stability, biocompatibility, selective transmissivity, low toxicity, and low immunogenicity, and they can traverse the BBB as a carrier for (immuno) therapeutic or diagnostic drugs.

However, there remain many challenges in the clinical application of exosomes. Their isolation and purification has not been standardized; miRNA quantification remains a limitation; the mechanisms involved in the loading and transport of siRNAs and miRNAs in exosomes remain to be unraveled; the multitarget nature of miRNA and the characteristics of exosomes with immune targets may result in side effects and toxicity *in vivo*; and bottlenecks in screening the donor-cell exosomes and drug imports into exosomes need to be eased. The application of exosomes in the diagnosis and treatment of malignant gliomas is still under research *in vitro* and in animal experiments. With the advances in technology, the clinical use of exosomes is expected in the diagnosis, treatment, and prognosis of gliomas in the future.

Conflicts of interest

None.

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