ORIGINAL PAPER



The Unwanted Cell Migration in the Brain: Glioma Metastasis

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Received: 3 February 2017 / Revised: 12 April 2017 / Accepted: 17 April 2017 / Published online: 6 May 2017 © Springer Science+Business Media New York 2017

Abstract Cell migration is identified as a highly orchestrated process. It is a fundamental and essential phenomenon underlying tissue morphogenesis, wound healing, and immune response. Under dysregulation, it contributes to cancer metastasis. Brain is considered to be the most complex organ in human body containing many types of neural cells with astrocytes playing crucial roles in monitoring both physiological and pathological functions. Astrocytoma originates from astrocytes and its most malignant type is glioblastoma multiforme (WHO Grade IV astrocytoma), which is capable to infiltrate widely into the neighboring brain tissues making a complete resection of tumors impossible. Very recently, we have reviewed the mechanisms for astrocytes in migration. Given the fact that astrocytoma

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shares many histological features with astrocytes, we therefore attempt to review the mechanisms for glioma cells in migration and compare them to normal astrocytes, hoping to obtain a better insight into the dysregulation of migratory mechanisms contributing to their metastasis in the brain.

Keywords Cell migration · Astrocyte · Glioma · Astrocytoma · Glioblastoma multiforme · Metastasis

Introduction

Cell migration is a highly orchestrated process which is known by now as a fundamental and essential phenomenon underlying tissue morphogenesis, wound healing and

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immune response. When its orchestration was disrupted, it would induce unwanted cell migration such as metastasis. Metastasis is one of the "Hallmarks of Cancer" and the leading cause of cancer mortality. When cancer has metastasized, the possibilities of curative treatment would greatly reduce. The latest data showed that there were 14 million new cases and 8.2 million cancer related deaths in 2012 [1]. The number is expected to rise by about 70% over the next two decades in the lowest-income countries.

Surgery is one of the main treatments for cancer. However, when Michael Baum asked the question "Does surgery accelerate or disseminate cancer cells?" [2], the effectiveness of surgery as a cancer treatment had already been questioned. There are reports on surgery significantly stimulating the malignant growth of tumor mass [3] and promoting micrometastases [4]. These suggestions chime perfectly with our previous findings that a physical scratch in culture or a stab wound in brain simulating surgery would induce the astrocytes along the wound to be reactivated into highly migratory cells [5, 6]. Some of these migratory cells also acquired many oncogenic properties [7]. These observations supported the potential of surgery in inducing unwanted cell reactivation, movement and migration, thus might result in metastasis after tumor removal.

Cancer in the brain is the most difficult among all cancers to deal with. Astrocytoma is the most common glioma arisen from astrocyte, and accounts for ~75% of all glioma [8]. World Health Organization (WHO) classified gliomas into four grades of ascending malignancy [9]. Glioblastoma multiforme (GBM) is grade IV that comprises 55% of all gliomas [10]. Their cells diffuse and infiltrate widely into brain tissues making a complete surgical resection impossible [11]. Migration and infiltration of astrocytes and glioma cells appear to be regulated by some very similar mechanisms [7]. This review addresses some interesting and promising mechanisms related to glioma cell migration, including metabolism, epithelial-mesenchymal transition (EMT) and mesenchymal-epithelial transition (MET), β-catenin, integrins, cell polarity, cytoskeleton, epidermal growth factor receptor (EGFR) signaling, aquaporins and so forth. Meanwhile, some of the similarities and differences between migration of injured astrocytes [12] and metastasis of glioma were compared (Table 1), with a hope to further delineate the multiple cues in GBM metastasis the unwanted cell migration.

Glioma metabolism

In the adult mammalian brain, the primary energy substrate is glucose [120]. An increase in glucose metabolism was observed in the glioma periphery. It would up-regulate aquaporin 1 (AQP1), lactate dehydrogenase (LDH), and cathepsin B. These up-regulations would contribute to the acidification of the extracellular milieu to enhance invasiveness of the glioma cells [16]. Tumor growth is usually faster than angiogenesis, thus leading to many portions of the tumor under a hypoxic microenvironment. The hypoxic condition would stimulate Warburg anaerobic glycolysis of GMB in both aspects of glucose consumption and lactate production [121–123]. Warburg effect, hypothesized by Nobel Laureate Otto Heinrich Warburg in 1924, was believed to be the root cause of cancer [124]. It stated that cancer, malignant growth, and tumor growth were caused by the fact that tumor cells mainly generate energy by non-oxidative glycolysis with lactate secretion even in the presence of oxygen. The changes of glucose concentration in tumors would induce carbohydrate-response element (ChoRE) expression. The hypoxia would lead to an increase in production of hypoxia-inducible factor (HIF-1) that acts as a key regulatory transcription factor responsible for adaptive cellular changes. HIF-1 and ChoRE interact to induce changes with particularly notable in genes on metabolism. These genes are glucose transporter (GLUT1), hexokinase (HK), phosphoglucose isomerase (PGI), phosphofructokinase (PFKL), fructose-bisphosphate aldolase (ALDO), glyceraldehyde-3-phosphate dehydrogenase (GAPDH), phosphoglycerate kinase (PGK), phosphoglycerate mutase (PGM), enolase 1 (ENOA), pyruvate kinase (PK), pyruvate dehydrogenase kinase (PDK1) and lactate dehydrogenase A (LDH-A) [125, 126]. Under stressful and pathological conditions, lactate and ketone bodies can be used as a substitute for glucose [127]. High concentration of lactate was known to induce glioma cell migration through its strong association with TGF-beta2-dependent regulation of MMP-2 and integrin $\alpha_{\nu}\beta_{3}$ receptors [17].

On the other hand, Professor Ursula Sonnewald has described that cultured astrocytes and glioblastoma C6 cells do not show any difference in lactate production in the presence and absence of glucose [14]. Instead, they are different in their ketone metabolism. Astrocyte utilizes ketone bodies to produce glutamine and glutathione. However, glioblastoma also utilizes ketone bodies, but could not perform anaplerosis. They could only produce an extremely small amount of glutamine, but a lot of glutamate and release them into the extracellular space. This glutamate would not only excitotoxically damage and kill many neurons in the tumor neighborhood via the activation of glutamate receptors and changes in the microenvironment [14], but also facilitate the invasiveness of the glioblastoma cells by stimulating cell proliferation and motility [128, 129]. Therefore, further studies on these specific characteristics and changes of metabolic pathways in tumor cells might lead us to identify some direct targets for the future development of metabolic therapy in blocking or manipulating glioma growth and metastasis.

Factors			Astrocyte migration	Glioma metastasis
Metabolism	Glutamate		Regulation of astrocyte synaptic cleft invasion via Cx30 [13]	Stimulation of proliferation and motility of glioblastoma cells via activa- tion of glutamate receptors and enhancement of invasion of surround- ing tissue by causing excitotoxic damage to normal brain and necrosis in glioblastoma [14]
	Glucose		Impact significantly on astrocyte proliferation, adhesion, and migration by increased production of inflammatory cytokines and oxidative stress [15]	Providing a scenario to the invasive potential of glioma cells in perivas- cular space [16]
	Lactate		Undetermined	Induction of glioma cell migration through TGF- $\beta 2$ -dependent regulation of MMP-2 and integrin $\alpha_{\alpha}\beta_{\beta}$ receptors [17]
Extracellular	ECM	Fibronectin [§] *	Involvement in astrocyte adhesion and migration [18]	Promotion of glioma cell migration [19]
Environment		Laminins	Laminins containing $\beta 2$ and $\gamma 3$ chains: indispensable for migration and spatial organization of astrocytes [20]	Prevention the invasion of the tumor spheroid into the target aggregate and inhibition of glioma cell migration [21]. Laminin-8: promotion the spread of glioma [22]
		TN-C*	Regulation of migration in cultured astrocytes [23]	High expression in glioma tissue and playing a crucial role in cell migration or invasion [24]
		Brevican	Undetermined	Consistent expression by human glioma and playing a role in glioma invasion [25]
		§N40	Participation in astrocyte migration after focal ischemia [26]	Link with NECL1 in inhibiting migration and invasion of glioma cells [27]
		Collagens* [‡]	Collagen I and VIII: stimulation of astrocyte migration [28, 29]	Collagen I: suppression of malignant glioma invasion [30]. Collagen IV: mediation of MMP-9 depletion induced diminishment of migration speed and reduction of invasion in glioma cells [31]
		Versican [§]	Undetermined	Correlation with invasive phenotype of low-grade astrocytoma [32]
	IL-1 β		Enhancement of astrocyte migration [33]	Induction of glioma cell invasion [34]
	IL-18		Undetermined	Enhancement of glioma cell migration through NO/cGMP pathway [35]
	5-HT		Undetermined	Positive modulation of glioma cell proliferation, migration, and invasion [36]
	MMPs ^{\$*II}		MMP-2: regulation of astrocyte motility through interplay with integrins and actin cytoskeleton [37] MMP-9: positive correlation with astrocyte migration [38]	MMP-2: promotion of glioma cell invasion [39] MMP-9: positive correlation with glioma cell adhesion, migration, and invasion through collagen IV [31]
Cell Mem- brane Proteins	Channels	AQPs	AQP4: enrichment in astrocyte endfoot membrane domains facing microvessels and pia [40]; with free mobility in plasma membrane and rapid diffusion into extending lamellipodial regions for astro- cyte migration support [41] AQP5: promotion of astrocyte process elongation via its up-reg- ulation and polarization to the migrating processes and plasma membrane in the leading edge of the scratch [6]	AQP1: enhancement of glioma cell growth, migration and invasion [42] AQP4: promoting glioma cell migration [43] AQP9: high expression in astrocytoma and positive correlation with the pathological grade of gliomas [44]
		ASIC-1	Undetermined	Regulation of glioma cell proliferation and migration through interactions with integrin- β 1, mediated by α -actinin [45]
		TRPV1	Mediation of extracellular Ca^{2+} influx and cytoskeletal changes induced by mechanical stress in astrocyte mobilization [46]	Undetermined

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Factors			Astrocyte migration	Glioma metastasis
		KCNH1	Undetermined	Low expression in brain metastasis including GBM patients [47]
		BK channels	Undetermined	Co-localization with CIC-3 chloride channel to the invading processes of glioma cells [48]
	Receptors	EGFR	Guidance of astrocyte migration [49]	Regulation of glioma growth control and metastasis [50]
		CXCR4/ CXCL12	Undetermined	Control of glioma invasion via autologous chemotaxis driven by their activation [51]
		uPAR	Undetermined	Promotion of glioblastoma cell migration and invasion [52]
		PDGFR	Association with MMP-9 expression induced by thrombin, IL-1 β , and LTA in astrocyte migration [33, 53–55]	Overexpression in high-grade poorly differentiated gliomas [56] and promotion of medulloblastoma migration [57]
		NTR	Undetermined	Regulation of glioma invasion in a neurotrophin dependent way [58] and mediation of medulloblastoma invasion via NTR proteolytic process- ing [59]
		EB12	Stimulation of ERK phosphorylation, Ca^{2+} signaling and induces astrocyte migration [60]	Undetermined
		$CysLT_1R$	Mediation of TGF- β 1 induced astrocyte migration [61]	Undetermined
		SIPR	Mediation of phosphorylated FTY720 to promote astrocyte migra- tion [62]	Enhancement of glioma invasion partially through uPA and CCN1 [63]
		P2YR ^{\$}	Interaction with α integrin to mediate astrocyte migration [64]	Mediation of glioma cell mobility and migration [65]
	Others	Integrins#	$\beta_1 \mbox{ integrin } Co-localization and interaction with MMP-2 at the leading edge of migrating astrocytes as a linker between pericellular proteolysis and the actin cytoskeleton [37]$	Regulation of glioma metastasis in the process of cell-cell and cell- matrix contact sites [45]
			$\alpha_{\nu}\beta_{3}$ integrin Induction of astrocyte migration [66]	High expression in glioma cells and contribution to the processes of glioma migration and invasion [67]
			$\label{eq:alpha} \begin{array}{ll} \alpha_5\beta_1 \text{ integrin} & Support of filopodial adhesion to the astrocytic migration template via interaction with fibronectin [68] \\ \end{array}$	Undetermined
			$\alpha_0\beta_1$ integrin Undetermined	Participation in the increase of diffusive properties of glioma cells [69]
		N-cadherin [§] *#	Promotion of a faster and less-directed migration in astrocyte under reduction of expression [70]	Impact on glioma cell polarity and promotion of a non-directed migra- tion [70]
		E-cadherin*#	Contribution to the distinct EMT in reactive astrocytes concomitant with enhanced migration and invasion activity [71]	Inhibition of glioma metastatic behaviors while restoration of its expres- sion [72]
		CD44 [#]	Playing a pivotal role in cytoskeleton activation and astrocyte migration via interaction with Rac1-PKN γ [73]	Involvement in human glioma cell invasion <i>in vitro</i> through its role in cell interactions with extracellular matrix proteins [74]
		Cx30 [#]	Regulation of astrocyte synaptic cleft invasion [13]	Undetermined
		Cx43#	Modulation of phenotypical changes in cell morphology, migratory activity including migration and process formation, and cell adhe- sion in astrocytes [75]	Regulation of p38-mediated cell migration and invasion induced selectively in glioma cells by low doses of γ -radiation in an ERK1/2-independent manner [76]
		Thy-1	Induction of astrocyte migration through the engagement of $\alpha_V \beta_3$ integrin and syndecan-4 via its mediated cell-cell contact [66]	Undetermined

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Table 1 (continued)

Table 1 (contin	nued)		
Factors		Astrocyte migration	Glioma metastasis
Cytoskeletons	Arp2/3 complex	Undetermined	Regulation of glioma cell migration and invasion [77]
	APC	Requirement for the polarization of migrating astrocytes [78]	Undetermined
	GFAP	Regulation of astrocytic hypertrophy and migration occurring after astrocyte activation [79]	A marker for extracranial metastasis of astrocytoma [80]
	β-Tubulin	Undetermined	High expression in astrocytoma and its nitration level positive correla- tion with the grade of astrocytoma [81]
	SEPT7	Undetermined	Involvement in glioma cell migration with the assistance of cofilin phosphor-mediated cytoskeleton locomotion [82]
	Filamin-A	Undetermined	A specific and sensitive marker for high-grade astrocytoma patients [83]
	Nestin	Localization at the margins of the wound and the immediately adjacent region in astrocytes [7]	Dysregulation of its localization within astrocytoma [7]
	Vimentin*	Expression in proximal reactive astrocytes correlation with migra- tion after focal brain injury [84]	Regulation of glioma progression and metastasis [85]
Signaling Pathways and Kinases	PI3K/Akt*	Activation in astrocyte migration [33, 53–55, 86]	Involvement in glioma metastasis [87]
	VEGF/VEGFR [§]	Strong impact on astrocyte proliferation and motility [88]	Requirement in glioma metastasis [89]
	Wnt/β-catenin*	Participation in Cdc42-dependent phosphorylation of GSK-3 β occurring specifically at the leading edge of migrating astrocytes [90]	Regulation of glioma cell migration and invasion [71, 91]
	PLC-PKCα	Promotion of astrocyte migration upon its activation [92]	Undetermined
	PLC-PKC8	Modulation of astrocyte migration through mediation of ERK1/2 phosphorylation [93]	Undetermined
	MEK-ERK	Initiation of astrocytes migration after injury partly upon its activa- tion [94]	Participation in glial tumor metastasis [95]
	GSK-3α	Undetermined	Suppression of cell invasion induced by its down-regulation [96]
	GSK-3β	Promotion of astrocyte proliferation and migration following injury [96]	Requirement for cell polarity formation and glioma cell invasion [97]
	cGMP-protein kinase G	Enhancement of astrocyte migration induced by its stimulation [98]	Undetermined
	ROCK1	Involvement in increased mobility of astrocytes after injury [99]	Regulation of glioma cell proliferation and metastasis [100]
	mTOR	Attenuation of astrocyte migration upon its blockade [101]	Mediation of migration and invasion in glioma cells [87]
	PKCα	Regulation of ERK1/2 phosphorylation and migration of astrocytes [92]	Regulation of glioma cell metastasis via actin cytoskeleton rearrangements [102]
	ΡΚϹζ	Negative association with proper polarization, orientation, and migration of astrocytes [103]	Promotion of glioblastoma cell migration and invasion [104]
	PKCe	Undetermined	Participation in the suppression of glioma metastasis [102]
	CK2	Regulation of migratory ability in astrocytes through reorganization of cytoskeleton [105]	Overexpression in glioblastoma and regulation of glioblastoma metas- tasis [106]

Table 1 (co	ntinued)			
Factors			Astrocyte migration	Glioma metastasis
Others	Rho GTPases	RhoA	Requirement in astrocyte migration [66]	Contribution to Netrin-1-promoted glioblastoma cell invasiveness and angiogenesis via its activation [107]
		Rac1	Requirement in cytoskeleton activation and astrocyte migration via interaction with CD44/hyaluronan [73]	Regulation of glioma cell migration and invasion [108, 109]
		Cdc42	Control of centrosome positioning in front of the nucleus and cell orientation toward the wound in migrating astrocytes [110]	Regulation of medulloblastoma cell migration and invasion via actin cytoskeleton disruption induced by Src [111]
	S100		Regulation of astrocyte migration [112] S100A4: reduction of the migratory capacity of reactive white mat- ter astrocytes in the injured CNS [113] S100B: regulation of astrocyte shape and migration via interaction with Src kinase [86]	S100A4: stimulation of the migration rate of glioma cells by modifying the organization of their actin cytoskeleton [114] S100B: participation in invasive properties of glioma cells [86]
	TWIST1*		Undetermined	Expression in human gliomas and promotion of invasion [115]
	Nm23-R1/	UDPKββ	Undetermined	Interaction with intermediate filaments in glioma metastasis [116]
	IDH1/2		Undetermined	Involvement in the metastasis of gliomas [117]
	Tiam1		Regulation of wound closure in confluent monolayers of primary astrocytes [118]	Undetermined
	PICK1		Undetermined	Negative regulation of neoplastic infiltration in astrocytic tumors [119]
*Epithelial-r \$Molecules i #Cell adhesic	nesenchymal tra n tumor microei n molecules	nsition markers avironment		
[‡] Collagen I [§]	ID3 and MMP0	belong to epithelia	ul-mesenchymal transition markers al-mesenchymmal transition markers	
AP-I activate AP-I activate 1, Cdc42 cel $CysLT_IR$ cys GBM gliobli $IL-I\beta$ Interle kinase kinase kappa-light-c PICKI prote serotonin/5-h and metastas	or protein 1, AP or protein 1, AP l division contra iteinyl leukotriet astoma multifor ukin 1 β , <i>IL-18</i> e, <i>MMPs</i> matrix hain-enhancer (in interacting w iydroxytryptami is-inducing prot	C adenomatous po C adenomatous po ol protein 42 homo are receptor 1, <i>EBI</i> me, <i>GFAP</i> glial fi Interleukin 18, $K($ metalloproteinas of activated B cell rith C Kinase-1, P ne, <i>SEPT7</i> septin ein 1, <i>TIMP</i> tissue	The structure of the second structure of the second secon	big potassium, <i>bHLH</i> basic helix-loop-helix, <i>CCN1</i> CCN family member exin 30, <i>Cx43</i> connexin 43, <i>CXCR4</i> chemokine (C-X-C motif) receptor 4, lermal growth factor receptor, <i>ERK</i> extracellular signal-regulated kinases, <i>K3β</i> glycogen synthase kinase 3β, <i>IDH12</i> isocitrate Dehydrogenase <i>I2</i> , <i>X</i> 5-lipoxygenase, <i>LTA</i> lipoteichoic acid, <i>MEK</i> Mitogen-activated protein hosphate kinase β , <i>βNECL1</i> nectin-like molecule 1, <i>NF-kB</i> nuclear factor eptor, <i>OPN</i> osteopontin, <i>PDGFR</i> platelet-derived growth factor receptor, se <i>C</i> , <i>PKC</i> protein kinase <i>C</i> , <i>ROCK</i> Rho-associated protein kinase, <i>5-HT</i> <i>ħy-1</i> CD90-cluster of differentiation 90, <i>Tiam1</i> T-cell lymphoma invasion ptor potential cation channel subfamily V member 1, <i>uPA</i> urokinase-type

plasminogen activator, uPAR urokinase receptor, VEGF vascular endothelial growth factor, VEGFR vascular endothelial growth factor receptor

EMT and MET

EMT is a process through which epithelial cells differentiate into mesenchymal cells by losing their cell polarity, cell-cell adhesion, and obtaining migratory and invasive properties. EMT is also known to be essential in the initiation of cancer metastasis [130]. MET is the reverse process of EMT [130]. Relatively little is known about the role MET plays in cancer when compared to the extensive studies of EMT in tumor metastasis. MET is believed to participate in the establishment and stabilization of distant metastasis by allowing cancerous cells to regain epithelial properties and integrate into organs [131, 132]. EMT initiates metastasis by breaking the carcinoma cells adhesion and allows them to break through the basement membrane into the bloodstream through intravasation. Later, when these circulating tumor cells (CTCs) exit the bloodstream to form micrometastases, they undergo MET for clonal outgrowth at these metastatic sites. Thus, EMT and MET form the initiation and completion of the invasion-metastasis cascade.

In gliomas, cells would undergo EMT to obtain the capacity to initiate metastasis and invasion. Glioma tissues have a diverse phenotype which might be caused by various microenvironmental factors and intrinsic genetic alterations. The process of EMT is highly affected by glioma microenvironment such as hypoxia or the enrichment of myeloid cells, indicating epigenetic mechanisms might be more crucial than genetic changes in this process. Interestingly, EMT-inducing factors in gliomas are typically varied from those in other cancers [133]. For example, gliomas rarely expressed the most important cell-cell contact factor E-cadherin [133]. Moreover, Twist-related protein 1 (TWIST1), a bHLH transcription factor orchestrating cancer metastasis through EMT [134], is believed to be a great promising target as a cancer therapeutic [135]. In addition, migrating glioma cells underwent EMT may also undergo MET for metastatic tumor nodules establishment [133]. In recent years, researchers have begun to investigate MET as one of many potential therapeutic targets in the prevention of metastasis [136]. This approach for preventing metastasis is known as differentiation-based therapy or differentiation therapy.

β-Catenin

Catenins are a family of proteins to form complexes with cadherin in cell adhesion. Four catenins have been identified. They are α -catenin, β -catenin, γ -catenin and δ -catenin. Several types of catenins work with N-cadherins to play important roles in learning and memory [137, 138]. Among these subtypes, most studies were focused on α -catenin and β -catenin on their multiple roles in cell adhesion. They were identified in association with cadherins at cell-cell junctions linking their cytoplasmic tails to actin via α -catenin [139].

 β -Catenin is a marker protein of EMT and has recently received a lot of attention. The dysregulation of β -catenin pathway inevitably affects cell growth, proliferation, and metastasis [140]. Our previous work has demonstrated that β-catenin was dissociated from the catenin-cadherin complex in cell membrane after astrocyte injury and the detached catenin was translocated into nucleus during astrocyte reactivation [7]. High level of β -catenin protein would lead to an enhancement in cell migration [141]. The high level is likely resulted from the inhibition of its degradation [142]. Inhibition of β -catenin reduces both the response of astrocytes to injury and the induction of the malignant phenotype of astrocytoma. In astrocytoma, catenin signaling pathways were also found to be activated and dysregulated [7]. Therefore, β -catenin is not only important in astrocytes migration after injury, but also essential in glioma metastasis [142–144].

β-Catenin is also regarded as a key signal transducer of the canonical Wnt signaling pathway in many central developmental and pathologic contexts [145]. The upregulation of many key molecules in the Wnt/β-catenin signaling pathway has been identified in astrocytoma [7, 142, 146]. The Wnt/ β -catenin pathway is involved in the regulation of glioma cell migration through promoting nuclear translocation of β -catenin [91]. The pathway was also shown to be up-regulated by FoxM1 which plays an important role in the development and progression of GBM by regulating factors involved in EMT and tumor cell invasion [91]. Wnt inhibitory factor 1 (WIF1) was found to be down-regulated in numerous cancers [147]. It is conceivable that WIF1 inhibits Wnt/β-catenin signal to induce the reversal of EMT, i.e., MET. Some recent basic and clinical research have provided promising results for treating various catenin-associated cancers [148]. Apparently, the Wnt/ β-catenin pathway plays crucial roles in glioma metastasis and elicits a variety of actions and functions, some of which may possibly even prove to be anti-oncogenic. Before further elucidation of the pathway, its diversity would create lots of difficulties in finding precision cancer therapeutic targets related to catenin [148].

Integrins

When cancer cells undergo metastasis, invasion and migration to a new tissue, they will have to detach, penetrate and attach to the basal matrix of the target tissue. This process allows cancer cell to leave from their primary site and pull itself forward into the new tissue. Integrins are a large family of cell-surface receptors mediating the attachment of tumor cells [149]. They are heterodimers with α and β subunits and with many variants [150]. Integrins interact with major plasma membrane components of the tumor cell and ion channels to play crucial roles in metastasis. Integrins are essential for cell migration and invasion, not only for their direct mediation of cell adhesion to the extracellular matrix (ECM) [150], but also for sending and receiving molecular signals for regulating cytoskeletal organization, force generation and survival.

Integrins express and distribute remarkably higher on the surface of malignant tumors as compared with the same type of pre-neoplastic tumors [151]. Its up-regulation in gliomas has been demonstrated to be closely correlated with tumor metastasis [28]. Tumor cells adhere and migrate along the ECM components in the brain via integrin receptors [152].

As mentioned above, integrin $\alpha_{\nu}\beta_{3}$ works closely with lactate in glioma metastasis. Now with more details, integrin $\alpha_{\nu}\beta_{3}$ is part of matrix metalloproteinase (MMP)-activating complex [153] which modulates glioma cell migration [154]. The up-regulation of integrin $\alpha_{\nu}\beta_{3}$ is a needed factor for transforming growth factors β (TGF- β) to promote glioma cell migration [155]. The interactions of integrins with ECM and the signaling pathways triggered could be negatively regulated by tumor suppressor PTEN through their direct dephosphorylation of two key tyrosine-phosphorylated proteins [156]. Moreover, integrin- β 1 could interact with an amiloride-sensitive nonselective cation channel through α -actinin to regulate glioma cell migration and proliferation [45]. This ion channel complex composed of acid-sensing ion channel (ASIC)-1 and epithelial Na⁺ channel (ENaC) α - and γ -subunits and has not been identified yet in normal astrocytes.

Integrins could indirectly contribute to tumor metastasis by activating other oncogenes/mitogens such as HIFs. In addition to the metabolic effects mentioned above, HIFs could contribute to the modulation of the tumor microenvironment through secretion of growth factors such as erythropoietin (EPO) and vascular endothelial growth factor (VEGF) [157, 158]. Paulus and Tonn identified that integrins are also involved in basement membrane invasion preceding meningeal dissemination and metastasis of glioma cells [159]. Taken together, integrins appear to play important roles in the penetrative growth, migration and invasion of glioma cells. Therefore, they have the potential to be attractive tumor therapeutic targets for GBM [160].

Cell Polarity

for directional cell translocation. Perturbation of cell polarity is a distinctive characteristic of cancer cells. Glioma cell polarization, migration and invasion are known to be affected by glycogen synthase kinase 3 (GSK-3), Arp2/3 complex, N-cadherin and Rho-mDia1 pathways [7, 77, 96, 161].

GSK-3 is encoded by GSK-3a and GSK-3b. The phosphorylation of GSK-3ß at the Ser9 (pSer9-GSK-3ß) was enriched and localized at the leading edge of scratched glioma cells [96]. The enrichment and polarized localization of pSer9-GSK-36 involved with PKC and MAPK pathways and were critical for glioma cell invasion [96]. On the other hand, the down-regulation of GSK-3a and 3b by specific small interfering RNAs inhibited glioma cell invasion [96]. Actin-related proteins (Arps) play a major role in the regulation of actin cytoskeleton. Two of its subunits, Arp2 and Arp3, closely resemble the structure of monomeric actin. Arp2/3 complex is a seven-subunit protein complex that binds to actin networks to rearrange actin cytoskeleton, an important process for cell locomotion, phagocytosis, and intracellular motility of lipid vesicles. Inhibition of Arp2/3 complex would demolish lamellipodia and cell polarity. This would seriously de-escalate the ability of glioma cell migration and invasion [77]. However, loss of N-cadherin in EMT would contribute to the loss of cell polarity and promote the invasive capacity with a faster but less-directed migration of astrocytoma [70]. Such effects were acquired through the modulation of focal adhesions and subsequent integrin dependent cell division cycle 42 (cdc42)-mediated polarity pathway [70]. mDial, a member of the formin protein family, is a Rho effector. Rho-mDia1 pathway works critically to direct cell migration in gliomas through regulating polarization and focal adhesion turnover by aligning cytoskeleton actin [161]. In addition, glioma cell polarization and directional cell migration were also found to be actively regulated by interstitial flow through a CXC chemokine receptor type 4 (CXCR4) dependent mechanism [51]. Although the above findings indicated an essential involvement of cell polarity in gliomas, the amount of data is still not enough to fully elucidate the underlying mechanism. Therefore, it is still too early to consider the translation of these polarity information into therapeutics in prevention of unwanted cell migration in cancer.

Cytoskeleton

Cytoskeleton is the network of filaments and tubules interconnecting filamentous bridges which give shape, structure and organization to the cytoplasm. Cytoskeletal filaments compose of microfilaments, intermediate filaments, and microtubules. Dysregulation of these filaments has been identified in various types of tumors and the locomotion of these filaments is a crucial contributor of cancer cell spreading.

Microfilaments, also called actin filaments [162], are filamentous structures in the cytoplasm and form part of the cytoskeleton, which are involved in the movement of all mobile cells. The importance of actin regulation has been underscored by inhibition of Rho kinase. Among many regulators of actin filaments, we shall focus more in discussion of the Rho GTPases family. Rho GTPases have been shown to regulate many aspects of intracellular actin dynamics in cell adhesion, migration and invasion. Rho GTPases belong to the family of small signaling G proteins and also are a subfamily of the Ras superfamily. Rho GTPases members have been identified in astrocytoma and melanomas [163]. The Rho family contains 22 members, among which Cdc42, Rac1, and RhoA are being studied in detail [163].

Activation of Rho or Rac has been shown to associate with the proliferative and migratory phenotype of glioma cells [82, 164]. Abnormalities of Rho and Rho-associated coiled-coil-containing protein kinase (ROCK) enhance glioma cell migratory phenotype and induce the local spread of GBM [165]. Overexpression of Rac1N17 in glioma cells has been shown to promote cell migration [108]. Deletion of Rac1 inhibited the medulloblastoma cells migration and invasion through decreasing the cross-linked actin network and pseudopodia [109]. Inhibition of Rho kinase leads to the down-regulation of matrix metalloproteinases (MMPs) and VEGF expression in glioma cells. These down-regulations would also reduce the migration of adjacent endothelial cells and angiogenesis. Moreover, Src-induced disruption of actin network would lead to a decrease of cell migration and invasion [111]. Again, the Src effect is also known to be mediated by the inactivation of Rho-Rac-Cdc42. Thus, Rho kinase might end up playing a key role in establishing a tumor microenvironment [166]. The upregulation of TWIST1, a great promising target as a cancer therapeutic [135], has been found to significantly promote actin cytoskeletal re-organization and enhance cell migration, adhesion and invasion in GBM [135, 167]. SEPT7 is documented as a cytoskeletal protein with GTPase activity and with markedly decrease level in various gliomas [82]. SEPT7 functions in gliomagenesis and in the suppression of glioma cell growth. Its expression is decreased in astrocytomas with different grades and plays a tumor suppressor role. It was found to reduce glioma cell migration by promoting the phosphorylation of cofilin, a widely distributed intracellular actin-modulating protein to cause actin depolymerization [82].

There are also other actin related regulators involved in glioma metastasis but without known relationship to Rho GTPases. For example, plasma filamin-A is an actin-binding protein crosslinking actin filaments to membrane glycoproteins, and was also determined to be a specific and sensitive marker for high-grade astrocytoma [83]. Extracellular S100 protein A4 interacts with both intracellular and extracellular signaling proteins to speed up astrocytic tumor cell migration through modification of actin cytoskeleton [114]. Blocking the PKC-mediated actin cytoskeleton rearrangements using co-treatment of As_2O_3 and berberine would significantly inhibit glioma cell metastasis [102].

Intermediate filaments GFAP and nestin expression in astrocytoma appeared to be dysregulated for their localizations in contrast to their organized pattern in normal reactive astrocytes [7]. Moreover, intermediate filaments interact with nucleoside diphosphate kinase β (Nm23-R1/ NDPK β), an enzyme functional in cell proliferation, differentiation, tumor progression and metastasis, in cAMPinduced differentiation of rat C6 glioma cells [116]. In glioblastomas, cells strongly express GFAP, vimentin and nestin. In subependymal giant-cell astrocytoma (SEGA), the majority of tumors were GFAP positive and cell processes were filled with intermediate filaments [168]. Furthermore, intermediate filaments were also abundant in gangliogliomas [169]. However, in oligodendrogliomas, intermediate filament proteins are barely observed [170]. Apparently, the direct involvement of intermediate filaments in glioma metastasis remains undefined.

Like in other tumors, glioma metastasis also requires a flexible adaptation of cell shape and cell plasticity. Microtubules associate with centrosome, and together they regulate cell structure and shape [171]. Microtubule β -tubulin shows elevated expression in astrocytoma, and the nitration of β -tubulin to form nitro- β -tubulin which functions in cytoskeleton and cell migration, is positively correlated to the grade of astrocytoma [81]. γ -Tubulin, a core centrosomal protein essential for microtubule nucleation, co-immunoprecipitated with tumor metastasis suppressor Nm23-R1, a constituent of the centrosome, indicating a possible role involved [172].

Although studies with consistent differences in cytoskeletal structure between gliomas and astrocytes have not been well identified so far, the presence of more subtle biochemical alterations in the cytoskeletal structure of glioma cells that contributes to metastasis indicates the important roles cytoskeletal network plays. Further work need to be directed at identifying the various alterations of cytoskeletons involved in glioma metastasis in a more systematic way.

EGFR Signaling

Epidermal growth factor receptor (EGFR) is a transmembrane receptor with intrinsic tyrosine kinase activity. It is a potent mitogen receptor but normally not detectable in the mature CNS. Astrocytes in the cerebral cortex of mouse express a low level of EGFR [173, 174], but with completely undetectable amount in human brain [175–178]. However, EGFR level in astrocytes is substantially induced and enhanced under mechanical injury, ischemia and human glial tumors [175–178]. Ligand binding to these EGFR causes receptor phosphorylation and activates several important signaling pathways such as PI3K, Akt and mTOR. These subsequently result in changes of cell morphology, motility, and gene expression leading to cell migration [179].

EGFR signal dysregulation is a very common phenomenon in gliomas [180]. The gene encoding EGFR is frequently mutated, amplified, and/or rearranged in malignant astrocytoma. EGFRvIII is a constitutively active form of truncated EGFR. GBM cells expressing EGFRvIII release soluble urokinase receptor to promote glioblastoma cell migration and invasion via activation of ERK1/2 pathway [52]. Dysregulation of EGFR pathway by the deficient in endogenous inhibitory elements such as microRNA-7 eventually leads to the failure in control of glioma growth and causes tumor metastasis [50]. Nevertheless, EGF is an essential factor to maintain the self-renewal of glioma stem cells in vitro, indicating EGFR signals play a role in maintenance of the stem cell population in gliomas [179, 181]. It is now believed that targeting EGFR with micro-RNA-7 might be a promising plasmid-based antitumor and anti-metastasis gene therapy for human malignant glioma treatment [50].

Aquaporins

Cell preparation for migration in its confined 3D microenvironments requires cell volume regulation via water permeation. Aquaporins (AQPs), a family of water channels, constitute the principal pathway for water movement across the plasma membranes. AQPs were identified to play important roles in tumor cell migration, angiogenesis, cerebral edema and cell-cell adhesion in the brain [182–184].

AQPs on tumor cell membrane could create a net inflow of water and ions to the leading cell protrusions to affect their polarization, total number and size [43, 185, 186]; meanwhile they could allow a net outflow at the trailing edge leading to the cell displacement [187]. Modification of the expression and localization of different isoforms of AQPs correlate well with glioma cell migration [42]. AQP1 and AQP4 are detected in all biopsies from glioma patients with their expressions associated with histological subtype and tumor location [42, 182]. High levels of AQP1 and AQP4 are detected in astrocytoma [182]. AQP5 is also detected in some of the biopsy samples. Furthermore, the level of AQP9 in astrocytoma is significantly higher than in the normal brain tissues and its expression levels are positively correlated with the pathological grade of tumor [44].

The level of AQP1 was reported to be extremely high in GBM [42, 188]. The high level of AQP1 was found to enhance glioma cell growth, migration and invasion [42, 189]. Up-regulation of AQP1 contributes to acidification of the extracellular milieu and to the invasive potential of glioma cells in perivascular space [16]. The exact subcellular localization of AQP1 in the CNS is still not clear and its functions remain to be elucidated.

An increase of AQP4 expression was found in GBM. AQP4 is the most studied AQP in the brain. It was believed to be the major water channel in maintaining water and ion homeostasis in CNS [182]. AQP4 expression was shown to be slightly higher than AQP1 in astrocytoma, and it was involved in promoting cancer cell migration [182]. GBM cell migration and invasion was significantly impaired after AQP4 was deleted [43]. AQP4 was reported to undergo rapid rearrangements of their localization in glioma, leading to changes of the actin and the osmolality of the cytoplasm [43, 185, 186]. Therefore, AQP4 might mediate water flux, thus facilitates rapid modification of cell volume and shape in order to accelerate cell movement [190]. PKC activation could inhibit the AQP4 mediated water permeability and at the same time the rate of glioma invasion [191]. However, the underlying molecular mechanisms of these AQP4 effects are far from understood. Therefore, further research on AQPs is urgently required to provide more insight into how water influences the unwanted migration and infiltration of glioma cells.

Other Factors in Glioma Metastasis

Tumor metastasis is a very complex process. Up to today, we have acquired a great amount of knowledge on its mechanism and markers, but unfortunately it is still far from identifying precise metastatic factors and mechanisms for therapeutic purpose. Primary tumor would create a favorable microenvironment to promote tumor cell metastasis. Many mediators and cellular effectors were accumulated in the local microenvironment that would contribute to tumors metastasis. Some of these key mediators and effectors include primary tumor-derived factors VEGF, CD44, N-cadherin, versican and osteopontin (OPN), local stromalderived components fibronectin, and tumor/stromal-derived factors CXCL12 and MMPs. Among these factors, VEGF, N-cadherin and MMPs have been addressed above. Some not described in details were summarized in Table 1.

In order to further understand this undiscovered life's mystery-unwanted cell migration, we have reviewed a variety of cellular and molecular components involved in glioma metastasis and compared them to normal astrocyte migration. In Table 1, we classified them into six categories based on their migration/metastasis-mediating characteristics. Of course, not all of these factors were being researched thoroughly. We have selected 69 factors and addressed the most interesting and promising ones with known details. Table 1 has shown that about 60% (41 factors) of the reviewed factors participated in both normal astrocyte migration and glioma metastasis. Therefore, astrocyte migration and glioma cell infiltration in the brain might be regulated by many common mechanisms [7]. However, the 18 factors specifically participated in glioma metastasis but not in normal astrocyte migration might be better candidates for future development of metastatic therapies.

In this review, we did not describe possible contributions from other cell types in the CNS to tumor metastasis. For example, glioma-derived ECM activates microglia to secret IL-18 to enhance the migration of glioma cell through NO/ cGMP pathway [35]. In addition, metastatic spreading of glioma could also be linked to their extracellular vesicles (EVs) released as exosomes to transfer tumor cell derived genetic materials and signaling proteins [192, 193]. The mechanistic environment caused by suspended and aligned nanofibers might also influence glioma cell migration and membrane blebbing dynamic, indicating the possible roles of biophysical components in glioma metastasis [194].

Furthermore, there must be many more unknown factors among cell and cell interactions waited to be discovered and identified. Moreover, cell drinking is known to be involved in glioma metastasis [193, 195, 196]; however, whether this drinking process a preparation of tumor cell to enter metastasis still requires future elucidation.

Conclusion

We have previously reported that physical scratch simulating surgery would result in astrocyte reactivation and a huge amount of cell migration along the scratch [5, 7, 12]. Among these reactive astrocytes, many of them acquired tumor cells characteristics through the dysregulation of β -catenin signaling pathway [7, 140]. These findings arouse our vigilance whether the well-established surgical removal of cancer as a choice of glioma and cancer treatment [7]. If this is valid, operation on tumor will well-induce a storm of cell reactivation and migration along the surgical wound - a perfect source of unwanted metastasis. The search for answers to the questions of why reactive astrocytes and glioma cells are highly migratory after injury, what factors lead to or facilitate metastasis, and what are the underlying migration mechanisms becomes urgently necessary so as for us to assess whether surgery is one of the causes of metastasis and still a proper treatment for cancer.

Acknowledgements This work was supported by the Beijing Natural Science Foundation (7091004); the National Basic Research Program of China (973 program) (2011CB504400); the National Natural Science Foundation of China (30870818, 31070974, 31171009 and 81471253 and 81671218); and the Foundation for Innovative Research Groups of the National Natural Science Foundation of China (81221002).

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