Glioma Cell Death: Cell-Cell Interactions and Signalling **Networks**

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Abstract The prognosis for patients with malignant gliomas is poor, but improvements may emerge from a better understanding of the pathophysiology of glioma signalling. Recent therapeutic developments have implicated lipid signalling in glioma cell death. Stress signalling in glioma cell death involves mitochondria and endoplasmic reticulum. Lipid mediators also signal via extrinsic pathways in glioma cell proliferation, migration and interaction with endothelial and microglial cells. Glioma cell death and tumour regression have been reported using polyunsaturated fatty acids in animal models, human ex vivo explants, glioma cell preparations and in clinical case reports involving intratumoral infusion. Cell death signalling was associated with generation of reactive oxygen intermediates and mitochondrial and other signalling pathways. In this review, evidence for mitochondrial responses to stress signals, including polyunsaturated fatty acids, peroxidising agents and calcium is presented. Additionally, evidence for interaction of glioma cells with primary brain endothelial cells is described, modulating human glioma peroxidative signalling. Glioma responses to potential therapeutic agents should be analysed in systems reflecting tumour connectivity and CNS structural and functional integrity. Future

insights may also be derived from studies of signalling in glioma-derived tumour stem cells.

Keyword Glioma · Cell death signalling · Mitochondria

Abbreviations

PDGF

Plase A2

PKAII

PUFA

roi

PDR

AAarachidonic acid Akt serine-threonine protein kinase CHOP/GADD153 CCAAT/enhancer binding protein homologous transcription factor **CNS** central nervous system COX cyclo-oxygenase $\Delta \Psi m$ transmembrane mitochondrial potential ER endoplasmic reticulum **ERK** extracellular signal-regulated kinase **GLA** gamma linolenic acid GF growth factors **GFAP** glial fibrillary acidic protein GRP78/BiP ER glucose-regulated protein 78 H_2O_2 hydrogen peroxide iNOS inducible nitric oxide synthetase LO lipoxygenase MEK-1/2 mitogen-activated protein kinase kinase 1/2 **MPT** mitochondrial permeability transition pore pol(ADP-ribosyl) polymerase **PARP PBR** peripheral benzodiazepine receptor

platelet-derived growth factor

phospholipase A2

type II protein kinase A

polyunsaturated fatty acids

reactive oxygen intermediates

peripheral benzodiazepine receptor

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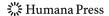
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TLR Toll-like receptor-2



TNF tumour necrosis factor

TRAIL TNF-related apoptosis-inducing

ligand

roi reactive oxygen intermediates **VEGFR** vascular endothelial cell growth

factor receptor

Introduction

Despite recent advances in treating malignant gliomas, prognosis remains poor. The median current survival in patients with glioblastoma multiforme, the most common, malignant human glioma, is less than 15 months in randomised controlled trials [1]. A better understanding of the complex biology of glioma cells and their interaction with the tumour micro-environment is essential for identification of molecular and signalling targets that can be exploited for therapeutic purposes. Gliomas differ from other tumour types in aetiology and localisation in the lipidrich tissue of the brain. Glioma cells are unusually resistant to apoptotic cell death, and a central question in glioma treatment is the identity of the survival/death decision points and signals amenable to therapy. In this context, recent studies in glioma have identified signalling pathways that mediate stress responses in glioma cells [2], and stem cell and metabolic studies have shed light on glioma development [3, 4].

This review will evaluate recent signalling pathways of stress signalling and membrane responses [4-29] identifying and characterising their temporal activation and compartmentalisation [4-15, 17-19, 21-24, 26-36] and supporting a role for lipid signalling in glioma cell death [2, 11, 13, 16, 18–21, 33, 37–45]. Current questions concerning cell death decision points will be raised, together with glioma network responses. Earlier events in triggering cell death signalling will be emphasised, rather than downstream, effector signals (caspase signalling is discussed by Simon Brown and Andrew Wyllie in this edition, [46, 47]). Novel concepts defining mitochondrial stress responses in primary glioma cell preparations and their interaction with brain endothelial cells will be presented. It is proposed that advances in therapy will come from understanding glioma signalling in systems which more closely resemble in vivo networks of interconnected cells. Such advances will depend on imaging, glioma stem cell analysis and identification of signal transduction elements leading to glioma cell death. Glioma mitochondrial and lipid metabolism are reviewed by Whittle and Colquhoun in this edition [2, 48] and brain endothelial cell function by Rizzo [49].

Cell Death Signalling in Glioma Pathogenesis

Mitochondrial and lipid metabolics have led to important insights in biology: the chemiosmotic analysis of oxidative phosphorylation [50] demonstrated that links between ion flux and cell signalling extended beyond the central nervous system (CNS). Advances in calcium signalling linked ion flux to phospholipase and membrane metabolism [18, 32, 51–53]. The concept of cell death signalling, initiated by Wyllie and colleagues [47, 54] was developed in seminal studies linking lipid signalling to CNS pathophysiology [30, 55].

Death Signals and Signalling Pathways: Intrinsic and Oxidative Systems In glioma, a range of environmental and pathological signals trigger cell death pathways [Table 1]. Such signals include stress signals, signalling to the intrinsic system, and intracellular reactive oxygen intermediate formation [2, 4, 11, 12, 14, 18, 19, 21, 22, 24, 27, 40], and we have proposed that lipid signalling networks are important in glioma pathophysiology [18, 41, 42, 45]. Salient questions in glioma signalling relate to the complex cellular interactions and signals within glioma tissue, which determine prognostic features for poor outcome. These prognostic properties include glioma proliferation, invasion, vascularisation, necrosis and resistance to radiation and chemotherapy (Fig. 1).

An important question is the extent to which hypoxia in glioma may cause selection of cells resistant to oxidative stress [4, 10, 12]. Signalling elements in oxidative stress in C6 glioma cells include transient activation of transcription factor nuclear factor kappa B and increased c-Jun N-terminal kinase and p38 kinase activity [24]. The intrinsic mitochondrial pathway may play a role in signalling oxidative cell death in rat C6 glioma cells, via the mitochondrial permeability transition pore and depolarisation of the mitochondrial transmembrane potential $(\Delta \Psi m)$ [5, 56]. Also, mitochondrial depolarisation, ERK activation and cell death associated with H₂O₂ treatment of the human glioma cell line A172 was inhibitable using catalase and MEK-1/2 inhibitor [22], and we observed mitochondrial depolarisation in response to oxidative stimuli in primary human glioma cell preparations (Fig. 2). Additionally, oxidative stress and reactive oxygen intermediates may act as mediators in radiotherapy [57].

Death Signals and Signalling Pathways: Extrinsic and Facilitative Signals In addition to intrinsic mitochondrial and endoplasmic reticulum (ER) pathways of cell death, extrinsic cell death signalling and sphingolipid signalling have been identified in glioma: TNF-alpha stimulates sphingolipid release and apoptosis [23], and ceramide and TNF-related apoptosis-inducing ligand

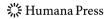


Table 1 Pathophysiological stimuli associated with polyunsaturated fatty acid signalling in glioma multiple pathways of cell death are involved, including intrinsic, extrinsic, endoplasmic reticulum (ER) and extrinsic cytokine-associated signalling pathways

Ischaemia, hypoxia,	Proliferation, roi	4, 12, 29
Oxidative stress	NFκB, c-JNK, p38, PARP, $\Delta\Psi$ m	21, 24, 58
	ERK, H ₂ O ₂ , Akt, PTEN	2, 11, 22, 49, 56
	Cardiolipin	16, 44, 48
Irradiation	ROS, Bcl-XL, JNK1/2/3	39, 41, 46, 58
ER stress signals	GRP78, ubiquitin, JNK	5, 14, 28
Cytokines	Rodent brain, astrocytes, microglia	9, 36
[TNF, IL-1β,TRAIL]	COX-2, iNOS	13, 23, 65
TLR	Dendritic cell migration	8
PGE2	proliferation, PKA II	33, 34
PUFA	roi, lipoperoxidation	11, 38, 18–20, 29, 41–45
	Glioma regression, MTP, Akt	2, 40
GF, PDGF, melatonin	Progression, PLC-Υ1	26, 59, 62
HGF, VEGFR	Adhesion, migration, AA, LO	26, 35, 61, 64
PBR	Intrinsic, $\Delta\Psi$ m	60, 62
Heavy metals	Intrinsic, roi, ER	27, 28
Cisplatin	Plase A2	58

(TRAIL) mediate apoptosis in invasive glioma cell preparations [7, 9, 13]. Other mediators which affect invasiveness and proliferative capacity are also associated with lipid signalling: myelin-associated glycoprotein in malignant gliomas contains phophatidylinositol specific phospholipase C, which increases migration and adhesion [50], PGE2 affects proliferation, oncogene expression and COX-2 and microsomal prostaglandin E synthetase-1 regulate migration, growth and apoptosis in human glioma cells [33]. In

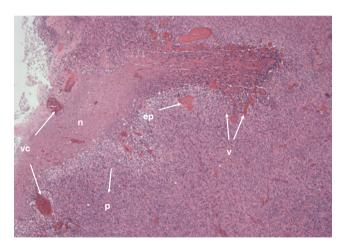


Fig. 1 Signalling of glioma and endothelial cell death and proliferation H&E section through malignant glioma, indicating histological evidence of differences in micro-environment associated with glial and endothelial cell proliferation, differentiation and death. Areas of necrosis (n) and intravascular coagulation (vc) in the *upper left* portion, border regions with rapid glial cell proliferation (p), detected using proliferative markers and evidence of mitosis, endothelial cell proliferation (ep), and increased vascularisation (v). Regional intratumoural hypoxia and ischemia may select cell clones resistant to hypoxia and with different responses to stress signals

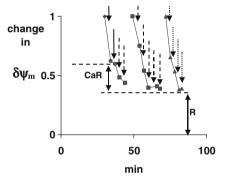
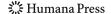


Fig. 2 Effect of calcium ionophore A23187, H₂O₂ and GLA on Δψm in primary glioma cell preparations. Typical experiment in an oligodendroglioma (WHO grade III) glioma cell preparation. Primary human glioma cell preparations were heterogeneous in morphology and in mitochondrial membrane polarisation (Δψm), using JC-1 fluorescence microscopy. The change in Δψm was analysed after exposure to stress stimuli, four to five principal patterns of basal Δψm were detected. The predominant cell types with polarised mitochondria were the large-/ medium-sized glial and astroglial cell populations, which varied in different glioma preparations and WHO grades, and which responded to stress stimuli by decreases in mitochondrial polarity. Depolarisation of mitochondria in previously polarised cells was determined by counting 100 cells, whose mitochondria were predominantly red (polarised) or green (depolarised). The proportion of cells changing from polarised to depolarised is shown on the y axis, and the time after loading probe into glioma cells on the x axis: Increasing agonist concentrations were added to three samples of glioma cell preparations. Basal $\Delta\psi m$ was determined, followed by Δψm in the presence of calcium ionophore A23187, 0.4 and 1.4 µM (circles, solid arrows) and H₂O₂ (squares, dashed arrows, 2.5 and 27.5 µM) was then added to the ionophoretreated preparation and further depolarisation, which had been Ca ionophore-resistant (CaR) was detected; H₂O₂ (0.1, 0.35, 14 and 29 µM) or GLA (triangles, dotted arrows, 10, 34, 133 and 378 µM) were added to two further glioma cell preparations. A residual, peroxidising agonist insensitive cell population (R) was detected in glioma cell preparations, as predicted by the Graeber hypothesis. WHO III and IV tumour mitochondria were more resistant to stress stimuli than WHO II tumours



glioma, it is also necessary to consider cerebral vascular and tumour-associated endothelial cell signalling [26, 47]. Also, gliomas are often located close to crucial neuronal networks which are highly sensitive to neurodegenerative and stress stimuli [5, 7].

Polyunsaturated Fatty Acid Signalling in Glioma Cell Death

Pathophysiological Signals Associated with Polyunsaturated Fatty Acids In glioma, polyunsaturated fatty acid (PUFA) signalling is linked to many pathophysiological stimuli associated with cell death signalling (Table 1). Stimuli associated with PUFA release include ischaemia [29], irradiation [39, 41], heavy metal ions [27, 28, 58] and specific cytokines, including TNF-alpha and TRAIL ligands [9, 23]. Previous work from our laboratory supported a pivotal role of PUFA in glioma cell death. It was proposed that PUFA, as locally released and locally active mediators, may act as network mediators in the pathophysiology of cell death in human primary brain tumours [18]. Therefore, identifying signalling activated early in glioma pathogenesis could provide a more selective and less toxic approach, using PUFA and agents targeting PUFA signalling as pharmacological agents. However, PUFA also plays an important role in neuronal signalling: A close connection between arachidonic acid (AA) and ion flux has been identified, both in recently characterised transient receptor potential and store-operated calcium entry channels and in classical ion channels associated with longterm potentiation and photoreception [32]. This may limit the use of intravenous and dietary agents that affect nontumour PUFA metabolism.

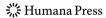
Exogenous Polyunsaturated Fatty Acids and Glioma Cell Death Signalling In human glioma cell preparations, exogenous PUFA increased development of apoptosis and necrosis [41, 45], and this was also observed in rat and human cell lines [11, 37]. Moreover, in five in vivo and in vitro models of glioma, PUFA [AA and gamma linolenic acid (GLA)] elicited glioma regression and cell death when applied intratumorally at appropriate concentrations [41]. These data suggested the potential to modulate death signalling pathways in glioma cells via alteration of membrane-associated or intracellular content of PUFA. Evidence for selective glioma targeting was obtained, comparing normal brain tissue with tumour, and PUFA signalling pathways via reactive oxygen intermediate (roi) signalling, including changes in basal, compared with PUFA-stimulated roi; increased roi response at higher tumour grade; comparison of normal brain tissue and tumour roi response [11, 19, 41]. The kinetics of intracellular roi generation measured in these studies indicated the balance of anti-oxidant status and the rate of pro-oxidant release within glioma cells. Other studies involving GLA effects on ATP depletion [11], modulation of mitochondrial hexokinase and palmitoyltansferase I and mitochondrial ultrastructure suggested that PUFA had specific effects on tumour mitochondria [2]. Glioma mitochondrial lipid abnormalities may be associated with increased susceptibility to lipoperoxidation and apoptosis [16, 38, 39]. However, both dose and localisation are crucial, as tumour proliferation has been reported at lower doses, and PUFA are involved in normal brain function [25, 41].

Endogenous PUFA and PUFA Metabolites in Glioma Pathogenesis Indirect PUFA involvement in glioma signalling was detected in phospholipase C and growth factor signalling. AA and other PUFA may mediate homeostatic signalling, affecting proliferation, migration and tumour growth [26, 59] via cycloxygenase and PGE2 pathways [33]. There is evidence of PUFA release in proinflammatory signalling [9, 23]. In glioma-derived microvascular endothelial cells, there was also evidence of lipoxygenase stimulating the formation of tubule-like structures [60].

Stress Signalling in Glioma Pathogenesis and Cell Death

The aggressiveness of human glial brain tumours is correlated with the level of tissue hypoxia [4, 12]. These areas of hypoxia were associated with necrosis, which is diagnostic of tumour malignancy, and it has been proposed that extreme anoxic conditions favour the development of cells with heightened resistance to cell death signals [10].

Glioma Stress Signalling via the Intrinsic Pathway of Cell Death Glioma stress signalling has been studied in mitochondria of human primary glioma cell preparations and in primary brain endothelial cells. A typical experiment, in which stress signals were added to glioma cells, is shown in Fig. 1. Glioma cells were exposed to calcium ionophore, hydrogen peroxide (H_2O_2) or GLA. Basal $\Delta\psi m$ and stimulated $\Delta\psi m$ were determined by fluorescence microscopy. Increasing concentrations of stress agonists being added until maximal depolarisation was achieved. Stress signals elicited immediate, dose-dependent depolarisation of glial mitochondria with differing efficacy: calcium ionophore A23187 and elicited partial (20–35%) depolarisation, compared with greater depolarisation in response to H_2O_2 and GLA. A residual, calcium ionophore-insensitive,



 $\rm H_2O_2$ -sensitive cell population was detected. Additionally, a residual resistant cell population with high $\Delta \psi m$ was detected after maximal stimulation with oxidative stress stimuli, consistent with the Graeber hypothesis [10]. Also, over longer time periods (4–48 h), other stress signals, such as serum deprivation and anoxia-elicited decreases in glioma cell $\Delta \psi m$.

Glioma and Endothelial Cell Stress Signalling via the Intrinsic Pathway Different responses of brain endothelial cells and other glioma cell types suggest that it is necessary to study signalling in multicellular systems. The stress response of primary glioma and endothelial cells to H₂O₂ resembled that of PUFA, indicating potentially similar roi-associated mechanisms, which may be important in cell death signalling, see also Colquhoun, this edition, [2]. However, the sensitivity of brain endothelial cells to peroxidising agents was higher than that of glial cells. Co-culture of glial and endothelial cells resulted in morphological evidence of cellcell communication and amplification of the primary stress response.

Stress Signal Transduction in Glioma Stress signalling mediators and systems in glioma are being identified and characterised: H_2O_2 -induced apoptosis is mediated by activation of ERK, via growth factor receptor/Ras/Raf/MEK signalling upstream of mitochondrial signalling in human primary glioma cells [22]. Downstream, ER stress activates JNK, ASK, the survival regulator GRP78/BiP, pro-apoptotic factor CHOP/GADD153, ATF6, IRE1 α and PERK/eIF2 α [14, 17]. Furthermore, peroxidative signal transduction in glioma involves PARP as well as lipid peroxides [21], and many stress signals involving intrinsic mitochondrial pathways have been identified: in rat cortical synaptosomes, lipid peroxidation affected mitochondrial function [15] and neuronal calcium dependent AP-1, caspase and then JNK [5].

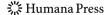
Therapeutic Options Associated with Stress Signalling The evidence for roi involvement in cancer and degenerative disease has provoked much debate about the potential therapeutic role of roi in cancer therapy [25]. Many experimental strategies have been developed, but these have not yet translated into therapies: Glioma cells have receptors, such as the peripheral benzodiazepine receptor, closely coupled to mitochondrial intrinsic signalling [60]. Mitochondrial targeting involving gene transfer and stimulation of oxidative metabolism has resulted in glioma regression in vivo and in vitro in a dog model [56]. Networks of interacting pathways and signals are crucial to cell death signalling in glioma; for example, a combination of calcium and ubiquitin signalling was more effective than either pathway alone [14].

Cell Networks: The Role of Functional Plasticity

Understanding the networks regulating cell death is a central problem in glioma pathophysiology. In primary human glioma cell preparations, waves of apoptosis were detected [45], suggesting coordinated pro-apoptotic signals in glioma cells, and in infused C6 implantation gliomas undergoing GLA-induced regression in vivo, apoptosis and necrosis were largely confined to the tumour and did not pass into the surrounding normal tissue [41]. This confirmed in vitro studies and human infusion data, which indicated higher sensitivity of glioma than normal cells to PUFA [19, 38, 40, 42]. There is evidence that coordinated cell-cell signalling is important in glioma cell death, e.g., enhanced immune recognition, involving both increased dendritic cell infiltration and tumour recognition [8]. Additionally, recent advances in stem cell analysis, coupled to in situ imaging, have provided important information on glioma development and neovascularisation.

The Impact of Stem Cell Analysis on Glioma Aetiology and Development Questions about glioma aetiology and development have been investigated using stem cell and genetic analysis [3, 55]. Glioblastomas contain transformed, multipotent, tumour-initiating cancer stem cells, which have radically changed our perspective of glioma pathology. These studies indicate that glioma cells may develop from transformed neural stem cells, but may develop from different precursor types, thus multiple triggers and selection pressures contribute to glioma development. The discovery that new neurones and glia are produced throughout life opens up new approaches to determining the origin and function of CNS tumours. The large astroglial cells expressing GFAP, which responded to stress signals by changes in roi and $\Delta \Psi m$ [19, 20] and Fig. 1, were found to proliferate in vitro and have been proposed as a candidate potential adult neural stem cell [3]. The functional character of these cells includes migratory activity and endothlial cell association when stimulated with appropriate growth factors and microenvironments [26, 33]. New suspension culture approaches [3, 37, 61] have facilitated the selection of these undifferentiated (or de-differentiated) glioma cell populations.

The Impact of Stem Cell Analysis on Glioma-Associated Endothelial Cell Function Microvascular proliferation and dysregulated endothelial cell development is characteristic of high-grade glioma. Endothelial cell signalling is discussed in more detail by Rizzo in this edition [49]. Stem cell research has indicated that the functional capacities of tumour-associated endothelial cells differ from normal endothelial cells in exhibiting greater differentiation capacity [62]. It has been proposed that glioma cells produce



diffusible growth factors that influence endothelial cell growth and neovascularisation [65] and brain endothelial cell migration [26]. Also, glioma-derived microvascular endothelial cells exhibited higher VEGFR and formed intercellular junctions and tubule-like structures [64]. These studies show increased structural connectivity in co-culture. This is reflected in functional evidence of connectivity, for example, in up-regulation of stress responses.

Current Questions in Cell Death Signalling in Glioma

Current knowledge of glioma pathogenesis suggests that diverse pathways and networks are involved in glioma cell death (Table 1). Attention has recently focused on the adaptive processes that gliomas use to evade cell death, with increased understanding of death checkpoints and with stem cell studies defining the functional plasticity of glioma cell interactions. Signalling processes, including ischemic and stress signalling, necrosis and intrinsic and extrinsic pathways, ER and cytokine-associated pathways, all input into critical cell death decision points. These pathways have provided fruitful avenues to investigate novel therapeutic strategies, and the increased sensitivity of mass spectroscopy and imaging techniques has improved our ability to identify transient signalling molecules. But, the cell biology of glioma is still only partially understood, and therapy is less advanced than in many other neuronal pathologies and systemic tumour types, and many basic questions remain to be answered. Some salient questions in glioma signalling relate to glioma cell proliferation and necrosis and endothelial cell dysfunction (Table 2). But, how these signalling mechanisms translate to the prognostic features for glioma malignancy and poor outcome remains a challenge.

Oxidative and Stress Signals and Mitochondria Evidence is accumulating for oxidative and stress signalling pathways as important components in triggering cell death in glioma (Table 2: question 1). Current research in mitochondrial cell death signalling, see Colquhoun [2] and Whittle [48], this edition, and the role of oxidative

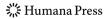
stress in selecting malignant glioma cells [4, 10] leads to the question as to whether intrinsic signalling and glioma sensitivity to PUFA may lead to useful therapeutic approaches [14, 16, 18, 44] (Table 2: questions 2 and 3); however, anti-angiogenic clinical trials in glioma [1, 25, 66] have achieved limited success and have emphasised the importance of understanding glioma cells and tumour-associated vascular endothelial cells in the context of their signalling networks.

Signalling Networks and Stem Cell Analysis Networks of interacting cells, including vascular endothelial and microglial signalling, provide survival and death signals affecting glioma fate [9, 36, 63]. The role of endothelial cell signalling in glioma pathogenesis is an important area of glioma research (Table 2, question 4). It is unclear whether disorders of endothelial growth and vascularisation are largely reactive to changed growth factor signalling in glial cells or whether endothelial cell signals are necessary in pathogenesis [64-66]. However, in brain endothelial cell studies, novel culture and imaging techniques, together with in vivo imaging are beginning to elucidate transcellular system signalling systems influencing glioma function [2, 3, 36, 63]. These approaches are also useful in determining the role of microglia and the local immune response (Table 2, question 5), where increasing immunogenicity and immune recognition may be important in therapy [8, 9, 36, 66]. Recent stem cell studies have given important information on functional plasticity and signalling networks in glioma (Table 2, question 6). The identification of multipotent glioma stem cells [3], characterised by capacity for self-renewal, multilineage differentiation and changed sensitivity to irradiation and chemotherapy [55], has focussed attention on microenvironmental factors that sustain glioma development and growth.

In conclusion, recent advances in lipid signalling have identified mediators of cell death in glioma, the intrinsic mitochondrial stress response and extrinsic adaptive responses. Effective cell death signalling often involves multiple interacting signal transduction pathways and networks of cells and signals, even when a single anticancer agent, such as PUFA or radiation is used. The brain tumour stem cell hypothesis provides a powerful model, which may

Table 2 Current questions in cell death signalling in glioma

- 1. Is oxidative stress involved in selecting malignant glioma cells [4, 10, 12]?
- 2. Is glioma tissue PUFA deficient in specific signalling locations [2, 7, 16, 38]?
- 3. Is glial cell sensitivity to roi/PUFA/stress signals a potential therapeutic option [11, 18-20, 24, 25, 39, 41-45]?
- 4. Is neovascularisation necessary for glioma survival [62, 63] and is anti-angiogenic therapy effective in glioma [1, 49, 66]?
- 5. What is the role of microglia and the resident inflammatory response in glioma cell death signalling and therapy [8, 9, 36, 46, 67]?
- 6. Will stem cell and epigenetic approaches identify cell death decision disorders in glioma [3, 66, 67]?



elucidate basic processes of glioma development. But, equally important is the identification and characterisation of interacting signals capable of promoting specific and selective glioma cell death.

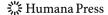
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