

## **Glioblastoma Biology, Genetics and Possible Therapies**

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Glioblastoma is the most aggressive intracranial tumor. Current treatment consists of surgery, radiotherapy, and chemotherapy (temozolomide). Resistance to radiotherapy and chemotherapy are frequent. Temozolomide is preferably used in patients who do not express MGMT. Temozolomide induces the formation of O6-methylguanine in DNA. MGMT repairs this DNA damage. Therefore, glioblastoma cells become resistant to temozolomide when MGMT expression ensures the repair of damaged DNA by temozolomide. On the contrary, if MGMT is not expressed, the DNA damage caused by temozolomide will not be repaired, and glioblastoma cells will die, that is, glioblastoma cells will be sensitive to temozolomide.

Lack of MGMT expression is associated with the hypermethylation of the MGMT promoter. Therefore, in clinical practice, an immunohistochemical approach is used to detect patients who do not express MGMT in glioblastoma biopsies, being these patients the ones who could benefit the most from MGMT expression laboratory assays [1].

Due to chemotherapy resistance, various combinations of drugs with temozolomide are being tested. The sensitization to temozolomide in previously resistant cells can be seen, thanks to the intervention of a second compound [2].

Bioactive compounds are also tested in order to prove their possible inhibitory activity of cell growth or migration in glioblastoma [3–5]. Sun et al. [3] demonstrated the antimetastatic potential of corosolic acid in glioblastoma cells by inhibiting the JAK2/MEK/ERK axis.

Kapoor-Narula and Lenka [4] demonstrated the anticancer effect of Oltipraz, a synthetic dithiolethione present in many vegetables, by decreasing the glioma cancer stem cells content in favor of differentiating GFAP+ glioma cells, together with the suppression of neurospheres formation. Even in vivo treatment with Oltipraz ectopically suppressed glioblastoma tumors xenografted in mice.

Several articles have revealed a preliminary positive effect of cannabinoids against glioblastoma [6–8]. Hohmann et al. [5] did not reach the same clear conclusions: on the one hand, they saw that cannabinoids increased the size of glioblastoma spheroids, but on the other hand, migration was unaffected.

Another way of intervention against glioblastoma is the direct attack of its brain tumor stem cells, trusting that such cells are the initiators and/or maintainers of the tumor, as well as being the cells that make the tumor resistant to chemotherapy and radiotherapy. In this sense, there are promising findings, such as those published by Lacore et al. [9], who blocked the M6a glycoprotein by siRNA, leading to a decrease in cell proliferation and invasion, as well as to an increase in radiosensitivity in glioblastoma stem cells.

Also, trying to target the stem cell component of this tumor, Essien et al. [10] assayed a combined treatment of an HDAC and an MEK inhibitor, together with radiation, detecting a bigger decay in the expression of stem cell markers Nestin and SOX2 than with the standard treatment of temozolomide and radiation. Other studies have shown the efficacy of epigenetically inhibiting HDAC in glioblastoma cells [11–15], even preferentially targeting the cancer stem cell compartment [16,17].



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**Copyright:** © 2023 by the authors. Licensee MDPI, Basel, Switzerland. This article is an open access article distributed under the terms and conditions of the Creative Commons Attribution (CC BY) license (https:// creativecommons.org/licenses/by/ 4.0/). Tumor cell dormancy complicates cancer therapy [18]. Cells that have metastasized to other organs remain in a quiescent state even for years, after which the cells might be newly activated and capable of originating the true metastatic disease. Every effort to understand the life cycle of tumor cell dormancy [19,20], the possible similarities found between tumor dormant cells and tumor stem cells [21,22], and, even more important, the sensitization of dormant cells to chemotherapy [23] are of great importance to overcome metastatic disease. Glioblastoma almost never metastasizes outside of the brain. Rather, it invades into the brain, but also develops tumor cell dormancy [20,22,24] capable of producing recurrent tumors several months after complete surgical resection, radiotherapy, and temozolomide treatments. Therefore, research has also been conducted on how glioblastoma dormant cells induced by temozolomide treatment can develop stem cell characteristics: Kubelt et al. [24] reported in this Special Issue about a possible connection between temozolomide-induced glioblastoma cell dormancy and the development of stem-like characteristics in glioblastoma cells.

Several other articles of this Special Issue concentrate on inhibiting specific targets with the idea of inhibiting or reducing cell proliferation, migration, and invasion. Then, Pai et al. [25] revealed that the inhibition of FABP6 (a bile acid carrier protein) reduced invasion and angiogenesis in glioblastoma cells by decreasing MMP-2 and VEGF. Secondly, the expression of XRN2, a 5'-3' exoribonuclease, was shown to be associated with cell migration and the invasion of glioblastoma cells [26]; therefore, inhibition of XRN2 expression might be a strategy to treat glioblastoma. And thirdly [27], it was revealed that the role of the Warburg effect in cancer cells, which turns on aerobic glycolytic processes and methylglyoxal synthesis, finally provokes a general glycation pattern that leads to the invasion of glioblastoma cells, a mechanism that might be disrupted by deglycating agents.

Another approach to combat glioblastoma might be the possibility of targeting specifically well-known pathways like Sonic Hedgehog, Wnt, Notch, TGFbeta, and others [28]. In such a way, single-cell studies [29], transcriptome analysis [30], and organoid models [31] are good approaches to define a holistic picture of glioblastoma.

Two reviews dealing about the epigenetic role of miRNA [32] and of circular RNA [33] in relation with different pathways that promote glioblastoma lead us into the last known category of epigenetic control, apart from histone methylation, histone acetylation, and DNA gene promoter methylation, all of them playing a role in the genesis of glioblastoma [34,35].

Finally, a new way of leading glioblastoma cells to differentiation is proposed by Hide et al. [36] based on ribosomes and ribosomal protein S6 administered to glioblastoma cells. Those cells might then be differentiated into reprogrammed glioblastoma stem cells with the possibility of the further differentiation of normal cells.

In summary, this second Special Issue on the Molecular and Cellular Mechanisms of Glioblastoma presents 16 articles dealing on the biology, genetics, and possible treatments against this devastating disease. Pathways to gliomagenesis and new targets have been explored, together with epigenetic possibilities like the inhibition of HDAC and the role of miRNA and circular RNA, tumor cell dormancy, cancer stem cells, and other approaches, to try to better understand and possibly combat glioblastoma.

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