

Treatment of Glioma in the 21st Century: An Exciting Decade of Postsurgical Treatment Advances in the Molecular Era



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CME Activity

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Abstract

The past decade has brought about major changes in the way we classify and have begun to approach patients with high-grade glioma. As we trend toward personalized medicine, we are now able to utilize the molecular characteristics of each individual's tumor in order to tailor their treatment, particularly if the patient is elderly or has a poor performance status at baseline. We address the state of the practice as of 2016 in regard to chemotherapy, immunotherapy, and tumor-treating fields. The goal of this review is to enhance readers' understanding of the nuances that are allowing clinicians to tailor the treatment of high-grade glioma more specifically.

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The past decade has seen remarkable research advances in glioblastoma that have culminated in tremendous improvements in patient outcome, with extensions in overall survival that have never been

seen previously. Whereas 2-year survival for patients with glioblastoma remained stagnant for decades at a dismal 10%, it nearly tripled to 27% with the advent of temozolomide (TMZ) and increased almost 5-fold for patients

whose tumors harbor a specific genetic alteration (O^6 -methylguanine-DNA methyltransferase gene [MGMT] promoter methylation) in a pivotal study just over 10 years ago.¹⁻³ The publication of that pivotal study a decade ago was the beginning of one advance after another, each documenting improvements in survival for patients with gliomas. Through advances in the identification of molecular signatures found in these tumors, new directions have been identified that are guiding treatment options in the field. Such new directions include the key role of genetic alterations that will inevitably replace the standard diagnostic procedures based on microscopy (histology/grade). These genetic alterations help to better match specific treatment modalities with a tumor based on its biologic features. This article will highlight several key advances in chemotherapy that have occurred over the past decade that, in turn, direct us toward an even brighter future for our patients.

CHEMOTHERAPEUTIC EXTENSION OF SURVIVAL IN PATIENTS WITH GLIOBLASTOMA: TMZ AND THE IDENTIFICATION OF MGMT AS A MOLECULAR MARKER

As the field of neuro-oncology was approaching the end of the 20th century, we had amassed a wealth of information about the molecular biology of glioma cells, which, in turn, identified many potential targets for therapeutics. Figure 1 illustrates the signal transduction pathways that drive glioma growth, with several principal components: the growth factor receptor (GFR) at the cell surface functioning as a “docking station” for growth signals; secondary messenger systems within the cells that are activated by GFRs; the DNA as a common convergence point for many signal transduction pathways to activate expression of cancer-associated genes (oncogenes); and the protein products of those oncogenes that then define the malignant phenotype (cell cycle progression/mitosis, angiogenesis, tumor invasiveness).¹ Each component of this molecular diagram is an ongoing or potential target for therapeutics. Despite numerous drugs developed to inhibit the new molecular targets of signal transduction, in the end it was targeting DNA—the archetypal target in most cancers—that led to the first chemotherapy breakthrough with the advent of TMZ.^{2,3}

The benefit of TMZ is summarized in Table 1. For decades, the dogma held that surgery⁴ and radiotherapy (RT)⁵ were the only 2 therapeutic modalities that improved survival of patients with glioblastoma multiforme (GBM), with only 10% of patients surviving 2 years. A pivotal European/Canadian study by Stupp et al² described the addition of a well-tolerated oral chemotherapeutic agent, TMZ, which alkylates DNA (adds methyl group—hence the term *alkylation*—to guanine residue of DNA) (since termed the *Stupp protocol*). The Stupp protocol includes TMZ at 75 mg/m² on days 1 through 42 with concomitant RT, followed by TMZ on days 1 through 5 of 28 for 6 consecutive months as adjuvant therapy at a dose of 150 to 200 mg/m². This regimen led to an improvement in 2-year survival to 27%. Furthermore, the presence of a specific alteration—methylation of the MGMT gene promoter—improved 2-year survival to 47%, a 5-fold increase compared with RT alone.³ The MGMT gene product repairs the DNA modification caused by alkylators such as TMZ, and therefore, silencing of the MGMT gene by promoter methylation is thought to confer increased sensitivity to TMZ. As a result of these 2 back-to-back publications in 2005, RT combined with TMZ became the long-awaited new standard of care in newly diagnosed glioblastoma and MGMT the key molecular marker in our field.

TAILORING THE STUPP REGIMEN FOR THE ELDERLY PATIENT POPULATION

MGMT promoter methylation thus became an important prognostic marker in neuro-oncology. Given that the benefit of TMZ is most apparent when this gene alteration is present, MGMT methylation may also be a marker that guides therapy choices. The role of MGMT in helping to guide therapy was highlighted in a series of reports pertaining to optimization of treatment for the elderly population with high-grade glioma.

Table 2 summarizes the evolution of treatment—in particular, the trimming of the Stupp regimen—for the elderly patient population. The French study⁶ found that a standard 6-week course of RT (6000 cGy over 6 weeks) improves survival compared with palliative care alone. However, the standard RT course was poorly tolerated by

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