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Biology Over Monotherapy: Mature Results From European Organisation For Research and Treatment of Cancer 22033-26033 Reaffirm Molecular Stratification, Not Treatment Sequence, Defines Prognosis in Low-Grade Glioma

Authors: Ivan F. Buckland, MD ✉ and Lisa A. Rich, MD, MS, MPH, and 19 others
Author Affiliations & Publication Info
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Accompanying Articles
Original Report
Temozolomide Versus Radiotherapy as First-Line Therapy for Low-Grade Glioma: Mature Results of a Randomized Phase III Trial (EORTC 22033-26033/NCIC CTG/TROG/MRC CTG)

Determining optimal treatment selection and timing for patients with low-grade glioma (LGG) remains one of the most complex clinical decision frameworks in neuro-oncology. This challenge stems from a heterogeneous patient population with generally favorable prognosis requiring careful balance of optimizing therapeutic efficacy and long-term survivorship with minimization of treatment-related toxicity. In parallel, advances in the understanding of glioma biology over the past two decades have prompted successive editions of the WHO classification of CNS tumors. The landmark 2016 WHO CNS classification formally incorporated molecular genetic features—including isocitrate dehydrogenase (*IDH*) mutation and 1p/19q codeletion status—thereby ushering in the molecular era of glioma. The current 2021 WHO classification further integrates molecular stratification through integrated diagnostic criteria that define biologically and prognostically distinct subsets of LGG.[1] This progress has substantially complicated the interpretation of trials designed and conducted before this molecular era.

The Takeaway

In the article that accompanies this editorial, Baumert et al[2] report the mature analysis of European Organisation for Research and Treatment of Cancer (EORTC) 22033-26033, a randomized phase III trial with a median follow-up of 13 years, that examined radiotherapy (RT) versus dose-dense temozolomide as up-front treatment in a cohort of 478 patients with clinical high-risk WHO grade 2 glioma and observed no significant difference in progression-free survival or overall survival in the intent-to-treat (ITT) population but the predictive molecular classification was more prognostic of long-term outcomes than treatment allocation. These findings reinforce the concept that biology—not treatment sequence—defines prognosis in this disease entity.

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European Organisation For Research and Treatment of Cancer (EORTC) 22033-26033, a randomized open label phase III trial comparing up-front dose-dense TMZ versus radiotherapy (RT; 50.4 Gy) among patients with high-risk WHO grade 2 LGG (Fig 1) reflects this dynamic landscape as it was conceptualized in the early 2000s and accrued between 2005 and 2012 during a transitional period in which molecular classification was evolving and adopting chemotherapy was reaffirmed as an active agent in glioma.[3] To this point, EORTC 22033-26033 was the first randomized glioma trial to prospectively incorporate molecular features (ie, 1p deletion) as a stratification factor. With a median follow-up of 13 years and mature survival data now available, Baumert et al[2] provide a unique opportunity for clinical insight as to the relative contributions of up-front treatment sequence and the natural history of clinical high-risk LGG. We contextualize these findings in molecular subgroup and treatment sequence and contrast this approach with the current standard of care (Fig 1) —controlled-toxicity treatment with RT and alkylating chemotherapy, as established by RTOG 9802, a randomized phase III trial of RT alone (54 Gy) versus RT plus procarbazine, lomustine (CCNU), and vincristine (PCV) chemotherapy in patients with high-risk WHO grade 2 LGG,[4] and supported by data from ECOG-ACRIN E3305.[5]

Fig 1. EORTC 22033-26033 and the treatment landscape of high-risk low-grade glioma. *Only published in abstract form, closed because of RTOG 9802 interim analysis of RT → PCV benefit. ‡OS, extent of resection; EORTC, European Organisation For Research And Treatment of Cancer; Gy, Gray; HR, hazard ratio; IDH, isocitrate dehydrogenase; ITT, intention to treat; LGG, low-grade glioma; mFUP, median follow-up; mOS, median overall survival; mPFS, median progression-free survival; MRI, magnetic resonance imaging; mt, mutant; NOS, not otherwise specified; NR, not reported; OA, observation; OS, overall survival; PCV, procarbazine, lomustine, vincristine; PFS, progression-free survival; RT, radiotherapy; RTOG, Radiation Therapy Oncology Group; STR, subtotal resection; TMZ, temozolomide; tot, treatment.

The EORTC 22033-26033 investigators successfully reclassified 73% of enrolled patients (351 of 478) per 2021 WHO criteria, generating the largest and most robust molecularly defined high-risk WHO grade 2 LGG cohort to date. For context, the high-risk arm of RTOG 9802 enrolled 251 patients, with only 47% of tumor specimens able to be reclassified for subsequent molecular testing.[6,7] With mature outcome data, the EORTC 22033-26033 trial now bridges the intent-to-treat population remains consistent with the earlier report that there are no significant differences in progression-free survival (PFS) or overall survival (OS) by treatment sequence (RT *v* TMZ) or TMZ-7. A nuanced interpretation of this negative result is required for the field to evaluate whether ET could be deferred in favor of chemotherapy without compromising survival in an attempt to mitigate neurotoxicity and thus preserve quality of life. Importantly, the lack of progression, approximately two thirds of patients in each arm crossed over to the other modality—specifically, 66% of up-front TMZ patients received salvage treatment with RT or chemoradiotherapy. The substantial proportion of patients with crossover confounds interpretation of OS, particularly given the availability of effective salvage therapy in a disease with a long natural history such as LGG. From a practical standpoint, it is reassuring for clinicians that each modality retains efficacy as a salvage option after progression on the other. Nevertheless, the lack of survival benefit to sequential treatment delivered at time of progression. As we enter the molecular genetic era, it will be increasingly important to understand how such treatment alters the tumor ecosystem, how this influences the efficacy of subsequent lines of therapy, and what mechanisms underlie combinatorial efficacy.[8–11] This is particularly relevant to *IDH*-mutant glioma, where the most regulatory approval of vorasidenib represents the first molecularly guided approach for this disease with ongoing efforts to define the optimal integration and sequencing with RT and chemotherapy in the clinic.[12]

The molecular subgroup analyses of EORTC 22033-26033 (Fig 1) provide the most instructive findings of this report. Across both *IDH*-mutant subgroups, initial treatment modality exerted no significant effect on OS, reinforcing that molecular biology is the primary determinant of outcome in *IDH*-mutant LGG. The established chemosensitivity and radiosensitivity of oligodendroglioma (*IDH*mt/codeleted) are reflected in the favorable median OS achieved with RT and TMZ (12.3 v 14.9 years), confirming that treatment sequencing is of limited consequence in this population when modality therapy is not used. In astrocytoma (*IDH*mt/non-codeleted), the numerically longer PFS with up-front RT (3.8 *v* 2.2 years) despite equivalent OS likely reflects effective TMZ salvage after RT failure. With the *IDH*-wildtype NOS subgroup, the apparent OS advantage with TMZ-7 over RT (4.7 *v* 2.5 years) must be interpreted cautiously given the small sample size (n = 64) and biological heterogeneity.

Traditionally, age >40 years has been incorporated as a criterion to define clinical high risk in LGG, used in both the RyanB scoring system[13] and the RTOG high-risk LGG criteria that defined eligibility for RTOG 9802. However, whether older age retains prognostic relevance in the molecular era or merely serves as a crude surrogate for other molecular features is an open question. When excluding this *IDH*-wildtype subset, age loses prognostic significance among *IDH*-mutant populations, with older patients demonstrating superior outcomes. However, if the historical basis of age-based risk stratification reflects the predominance of *IDH*-wildtype tumors rather than a true biological impact, then the RTOG high-risk definition and the trial populations defined by these criteria likely enrolled biologically and prognostically heterogeneous cohorts obscuring the treatment effect and outcome signals. Incidentally discovered molecularly annotated data sets is warranted and may facilitate development of novel definitions and criteria for high-risk *IDH*-mutant LGG for use in future trials.

A key hypothesis for testing up-front TMZ was that it may better preserve neurocognition and health-related quality of life (HRQoL) relative to RT. These hypotheses were largely not attained to these patient-centric considerations, which are of importance in LGG, a disease predominantly affecting younger adults with a relatively favorable prognosis. Prior reports from this trial found no significant differences in HRQoL, memory, or global cognitive function within the first 3 years between arms and also noted that early HRQoL declines were more often associated with tumor progression rather than with volume of treatment exposure.[14,15] More than 80% of patients received 1-dimensional conformal RT, whereas more modern techniques and modalities such as intensity-modulated RT, volumetric-modulated arc therapy, and proton RT are increasingly used in contemporary practice to mitigate neurotoxicity.[16] The available evidence from EORTC 22033-26033 therefore does not substantiate selection of up-front TMZ over RT on the basis of neurotoxicity concerns. Nevertheless, late cognitive effects remain a legitimate concern. A longitudinal study of LGG patients treated with RT demonstrated declines in attention and executive function at a median follow-up of 12 years—deficits that were not apparent at earlier time points and underscore that RT technique and treatment volumes reflect clinical practice in the 1990s.[17] With a median follow-up of 13 years, EORTC 22033-26033 is ideally positioned to evaluate long-term HRQoL and neurocognitive outcomes by modality; however, attrition of follow-up questionnaires at later time points limits these efforts. Early side effects were low in both arms, 1.7% with TMZ and 4.2% with RT, including high-grade hematologic toxicity (eg, lymphopenia, neutropenia) and cognitive fatigue. Late toxicities (eg, radiation-induced neurocognitive decline, cognition, memory) inadequately tied to MTD. Although preservation of neurocognition was a primary justification for up-front TMZ versus RT, the reporting of comprehensive long-term neurocognition remains an undercharacterized aspect of this mature data set.

The long-term update of EORTC 22033-26033 confirms that molecular subtype, rather than treatment sequence, is the chief determinant of clinical outcomes. The comparable interim overall benchmarks of both monotherapy arms serve as a compelling reminder that combined modality therapy remains the standard against which future approaches should be evaluated for high-risk *IDH*-mutant LGG. As the field further refines molecular risk stratification and advances toward biology-guided treatment paradigms, the insights offered by EORTC 22033-26033 data set will continue to shape how we select, sequence, and individualize treatment for patients with *IDH*-mutant LGG.

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Helen A. Shih

Employment: Mass General Brigham, Elliot Hospital, Manchester, NH (I)

Honoraria: UpToDate (less than $10,000 USD in any calendar year), MedLink Neurology (less than $10,000 USD in a single calendar year)

Consulting or Advisory Role: Advanced Accelerator Applications/Novartis (less than $10,000 USD in any calendar year), GERSHO (less than $10,000 USD in a single calendar year), Nurocor (less than $10,000 USD in a single calendar year)

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