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### Authors

Mellinghoff, Ingo K

Chang, Susan M

Jaeckle, Kurt A

et al.

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## IDH Mutant Grade II and III Glial Neoplasms

Ingo K. Mellingshoff<sup>1</sup>, Susan M. Chang<sup>2</sup>, Kurt A. Jaeckle<sup>3</sup>, Martin van den Bent<sup>4</sup>

<sup>1</sup>Department of Neurology and Human Oncology & Pathogenesis Program, Memorial Sloan Kettering Cancer Center, New York, New York, USA

<sup>2</sup>Department of Neurological Surgery, University of California San Francisco, San Francisco, California, USA

<sup>3</sup>Department of Neurology and Hematology/Oncology, Mayo Clinic, Jacksonville, Florida, USA

<sup>4</sup>Department of Neurology, Erasmus MC Cancer Institute, Rotterdam, The Netherlands

### Abstract

Mutations in isocitrate dehydrogenase (IDH) 1 or IDH2 occur in the majority of adult low-grade gliomas and, less commonly, in cholangiocarcinoma, chondrosarcoma, acute myeloid leukemia, and other human malignancies. Cancer-associated mutations alter the function of the enzyme, resulting in production of R(-)-2-hydroxyglutarate (R-2-HG) and broad epigenetic dysregulation. Small molecule IDH inhibitors have received regulatory approval for the treatment of IDH mutant (mIDH) leukemia and are under development for the treatment of mIDH solid tumors. This article provides a current view of IDH mutant adult astrocytic and oligodendroglial tumors, including their clinical presentation and treatment, and discusses novel approaches and challenges toward improving the treatment of these tumors.

### Keywords

Isocitrate Dehydrogenase (IDH); Low-grade Glioma (LGG); Cancer Metabolism; Response Assessment in Neuro-Oncology (RANO)

### Molecular Pathogenesis of mIDH Glioma.

Cancer-associated isocitrate dehydrogenase (IDH) mutations first emerged from a comprehensive analyses of mutations in protein-coding genes in colorectal cancer <sup>1</sup>. In 2018, whole genome sequencing uncovered the presence of these mutations in glioma <sup>2</sup>, a surprising finding that was rapidly confirmed in a much larger number of tumors <sup>3 4</sup>.

The extraordinarily high prevalence of IDH mutations in adult low-grade and anaplastic glioma (see below) and that fact that these mutations clustered in key arginine residues

**Corresponding Author:** Martin van den Bent, Brain Tumor Institute Erasmus Medical Center (MC) Cancer Institute, Rotterdam, the Netherlands, m.vandenbent@erasmusmc.nl.

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within the enzyme's active site (R132 of IDH1 and R140 or R172 of IDH2), immediately pointed toward a prominent role of the mutant IDH enzyme in the pathogenesis of these tumors. Several additional observations supported this conclusion. First, IDH mutations occur at the earliest stages of low-grade glioma<sup>5</sup>. Secondly, introduction of the mIDH enzyme into cells is sufficient to induce the distinct pattern of DNA hypermethylation associated with IDH mutations in human glioma<sup>6</sup>. And, unlike many other genetic alterations found in diffuse glioma, IDH mutations remain detectable throughout the disease course, suggesting a contribution of the mutant enzyme to tumor maintenance<sup>7-9</sup>.

Metabolic studies revealed that cells expressing the mIDH enzyme produce the R(-)-enantiomer of the metabolite R(-)-2-hydroxyglutarate (R-2-HG), which accumulates in IDH-mutant human gliomas<sup>10 11 12</sup>. Accumulation of R-2-HG leads to competitive inhibition of  $\alpha$ -ketoglutarate-dependent enzymes, a large protein family that includes the ten-eleven translocation (TET) family of 5-methyl cytosine hydroxylases, the jumonji domain containing (JmjC) family of histone lysine demethylases, enzymes involved in nucleic acid metabolism, and many enzymes with still unknown functions<sup>13</sup>.

Several findings support the conclusion that the "onco-metabolite" 2-HG is the critical mediator of the oncogenic functions of mIDH. Cell-permeable esters of R-2-HG phenocopy the effects of mIDH in experimental models and ectopic expression of the dehydrogenase that counteracts the activity of the mIDH enzyme is sufficient to reverse the cellular effects of cancer-associated IDH mutants<sup>14 15 16</sup>. Nonetheless, much remains to be learned about the role of mIDH in gliomagenesis. Of note, there are currently no preclinical models that recapitulate the genetics and growth pattern of IDH-mutant low-grade glioma. It seems plausible that several factors might relieve mIDH cancer cells from their dependency on the mutant enzyme for growth and survival<sup>17</sup>.

### Clinical Presentation of mIDH Glioma.

IDH mutations are exceptionally common in adult low-grade and anaplastic glioma. IDH mutation has been reported in 50–81% of WHO grade II gliomas<sup>18,19</sup>, 54% of Grade III, and 15–20% of Grade IV gliomas<sup>18,20</sup>. In a series of patients with low grade glioma, 52% were mIDH, 30% mIDH 1p/19q co-deleted (codel), and 18% IDH wildtype (IDHwt)<sup>19</sup>. In a large database of 2193 mIDH gliomas, 80% were LGG and 20% Grade IV tumors<sup>20</sup>. In a correlative analysis of 106 patients with high-risk low-grade glioma treated on NRG/RTOG 9802, 41% were mIDH/non-codel; 35% mIDH/codel and 24% IDHwt. Most 1p/19q codel gliomas also have IDH R132H mutations or non-canonical mutations (e.g., R132C or IDH2)<sup>21,22</sup>. In a database study of 911 patients with mIDH high grade glioma, 47% were non-co-deleted and 53% were co-deleted<sup>23</sup>.

IDH mutated gliomas are more common in patients < 55 years of age. Grade IV mIDH tumors are more frequently encountered in older patients but also occur in young patients<sup>18 24</sup>.

IDH mutated astrocytomas usually show robust immunohistochemical staining for p53 and loss of expression of alpha-thalassemia/mental retardation X-linked (ATR-X) protein. In

contrast, mIDH/codel tumors are more often p53wt and have *TERT* promoter mutations<sup>25 26 27</sup>. In patients with mIDH tumors, the presence of *CDKN2A* homozygous deletion varies from 7–42% (median, 22% and is an unfavorable prognostic marker for progression-free and overall survival, in both low and high grade gliomas<sup>20</sup>. In the POLA database and the series from Wijnenga, *CDKN2A* homozygous deletions were not typically encountered in the WHO Grade II tumors<sup>23 28</sup>. Although most mIDH tumors have low tumor mutation burden (TMB), the presence of higher TMB was associated with a less favorable prognosis<sup>29</sup>. It should be noted, however, that these studies did not use the classical “high TMB” definition issued for checkpoint inhibitors and instead defined TMB cutoff values within their dataset to distinguish patient subgroups.

IDH R132H mutated tumors more commonly arise in the frontal lobes, whereas those with non-canonical mutations (i.e, R132C; IDH2) have a wider CNS distribution, including infratentorial or multicentric locations<sup>30 22</sup>. The ‘T2-FLAIR mismatch sign’, present in a minority of mIDH astrocytomas, is characterized by a T2 hyperintense rim on FLAIR imaging, and homogeneous T2 internal appearance (Figure 1). Gliomas with IDH mutation more frequently enhance than mIDH/codel tumors, whereas the latter (Figure 2) are more often associated with heterogeneous T2 internal appearance and calcification<sup>31 32 33</sup>. Recently, there has been interest in application of radiogenomics to distinguish mIDH tumors but thus far these efforts have shown limited sensitivity and specificity<sup>34</sup>.

IDH mutated gliomas typically have a more indolent biological behavior than IDHwt tumors. Patients often present with seizures without other focal signs or symptoms. Patients with mIDH Grade II and III tumors more commonly present with seizures than those with IDHwt tumors<sup>22</sup>. Patients with IDHwt tumors, compared to those with mIDH, appear to associate with greater cognitive and physical impairment<sup>35</sup>.

In a multivariable analysis, mIDH was one of the most significant independent variables correlating with lower risk of death, in particular among patients with WHO grade II and III tumors<sup>36</sup>. Gross total or ‘supratotal’ resection is more frequently achieved in patients with mIDH tumors compared with IDHwt tumors<sup>37,38</sup>, possibly in part due to relatively sharply demarcated borders radiographically<sup>37</sup>, and the more frequent unifocal nature and frontal location. Patients with mIDH/non-codel tumors have intermediate survival outcome between those with oligodendroglioma (mIDH/codel) and IDHwt tumors. In NRG/RTOG 9802, treatment of patients with high-risk low grade mIDH gliomas with combined modality radiotherapy plus adjuvant PCV was associated with longer survival outcome than with RT alone [PFS: mIDH/non-codel - HR 0.32, p=0.003; mIDH/codel HR 0.13, p<0.001]; OS: mIDH/non-codel – HR 0.38, p=0.13; mIDH/codel – HR 0.21, p=0.29)]<sup>39</sup>. In the Phase III CATNON trial for patient with WHO Grade III non-codeleted gliomas the OS for IDH1/2mt patients with profiles consistent with lower grade tumors has not been reached, and 5.6 years in those with higher grade features<sup>40</sup>. In a study of pediatric and young adult patients with mIDH tumors, median PFS was 4.62 years and OS 17.2 years, with shorter survival observed in the young adult cohort<sup>41</sup>.

## Current Treatment of Lower-Grade Glioma.

The treatment of lower-grade glioma is based on a multimodality approach. It is important to note that the landmark studies that provide the foundation for the current treatment approach for LGG and anaplastic glioma were designed and conducted prior to current molecular classification of glioma being established and when modern surgical or radiotherapy techniques such as Intensity Modulated Radiotherapy or proton therapy were not available. As such, there are inherent limitations in trying to extrapolate results to the IDH mutated glioma subgroup. Future studies stratifying patients into homogeneous populations will be critical to assess the benefit of novel therapies.

Maximal safe, surgical resection remains the initial treatment for LGG to enable an accurate diagnosis and improve clinical outcomes such as progression-free survival, overall survival and risk of malignant transformation<sup>42-44</sup>. The impact of maximal resection as first-line treatment may be more important for mIDH astrocytoma than oligodendroglioma<sup>45</sup>. Improved surgical techniques such as intraoperative MRI and electrostimulation mapping during an awake craniotomy allow for more extensive resection while minimizing neurologic injury.

Radiation therapy is an important adjunct in the management of LGG and several studies have explored the optimal timing and dosing schedule. The European Organization for Research and Treatment of Cancer (EORTC) 22845 study comparing early RT after surgery vs RT delayed until time of progression, showed no significant difference in OS (7.4 years vs 7.2 years), but patients who received early RT had improvements in seizure control and median PFS (5.3 years vs 3.4 years with delayed RT)<sup>46</sup>. Two randomized studies evaluating high dose RT versus low dose RT did not show any significant differences in PFS and OS, but long-term analysis demonstrated improved quality of life in patients treated at the lower radiation dose<sup>47-49</sup>.

The optimal use of RT and/or chemotherapy after surgery for low-grade gliomas continues to be defined. Several prognostic factors have been proposed to better identify patients at high risk for malignant transformation and may benefit from aggressive management with adjuvant chemoradiation. High-risk factors include age > 40 years, subtotal resection/biopsy only, astrocytic lineage (lack of 1p/19q codeletion), neurologic deficits prior to surgery, tumor diameter > 6 cm, tumor crossing the midline of the brain, and tumors located within or adjacent to eloquent areas of the brain<sup>50-52</sup>.

Patients without these risk factors can be considered at low risk; therefore, after gross total resection, they are usually observed closely with regularly scheduled surveillance imaging to assess for intervention at the time of progression. The EORTC brain tumor group is conducting a phase 3 study for patients with *IDH* mutated 1p/19q intact lower grade glioma following resection, without a need for immediate post-operative treatment, to establish whether early adjuvant treatment with radiotherapy and adjuvant temozolomide in this clinically favorable group of patients will improve outcome compared to active surveillance. The primary endpoint is first intervention free survival with multiple secondary

endpoints of PFS, OS, seizure control and health related quality of life. [*EORTC-1635-BTG ClinicalTrials.gov Identifier: NCT03763422*].

In an attempt to defer the adverse effects of RT, several studies have evaluated chemotherapy alone<sup>53,54</sup>. A report of the EORTC 22033–26033 study of temozolomide versus RT in high risk LGG did not demonstrate a difference in PFS, but radiotherapy tended to be superior in mIDH astrocytoma. The results regarding the effects on OS are pending<sup>55</sup>.

The survival benefit of adjuvant chemoradiotherapy for high-risk LGG was demonstrated in the Radiation Therapy Oncology Group 9802 phase III trial that randomized patients to receive RT or RT plus combination chemotherapy with PCV (procarbazine, lomustine, and vincristine). Based on the pivotal data showing an almost two-fold increase in OS for patients in the chemoradiation therapy arm compared with the RT alone arm (13.3 years vs 7.8 years), high-risk patients with low-grade gliomas should receive radiotherapy followed by adjuvant chemotherapy rather than RT alone<sup>56</sup> (Figure 3). This study was conducted prior to the molecular characterization of LGG. A post-hoc molecular analysis on a subgroup of patients from this trial<sup>39</sup> confirmed that patients with *IDH* mutated gliomas with or without 1p/19q codeletion benefited from the addition of PCV to radiotherapy, but suggested that patients with *IDH* wild-type astrocytomas may not benefit from this combination.

The CATNON trial investigated concurrent and adjuvant temozolomide in anaplastic glioma and observed only benefit of the adjuvant treatment in mIDH anaplastic astrocytoma, not in IDHwt anaplastic astrocytoma. In mIDH tumors, adjuvant temozolomide improved outcome (HR 0.48, 95% CI (0.35, 0.67);  $p < 0.0001$ ), 5 year survival increased from 62.0% (95% CI: 54.4, 68.7) to 81.6% (95% CI: 75.5, 86.4)<sup>57</sup>.

With the introduction of temozolomide as the standard of care for glioblastoma<sup>58</sup> and based on the improved safety profile compared to nitrosoureas, in clinical practice, patients are commonly treated with temozolomide. The ongoing CODEL phase III study randomizes patients with 1p/19q co-deleted WHO grade II and III gliomas to receive either RT followed by PCV or RT with concurrent and then adjuvant temozolomide to address the comparison of these 2 chemotherapy regimens. [*ClinicalTrials.gov Identifier: NCT00887146*].

## Development of mIDH Inhibitors.

Inhibiting the aberrant activity of mutant enzymes represents an established pharmacological strategy for the treatment of human cancer, exemplified by the class of kinase inhibitors<sup>59</sup>. Cancer-associated mutant IDH enzymes represent attractive drug targets for the development of mutant-selective inhibitors because these mutations cluster in key arginine residues within the enzymes' active sites (R132 of IDH1 and R140 or R172 of IDH2) and because successful inhibition of the mutant enzyme can readily be ascertained through measurements of 2-HG in tumor biopsies<sup>10 60</sup>. In patients with acute myeloid leukemia (AML) or cholangiocarcinoma, two other human cancers with frequent IDH mutations, 2-HG can also be detected in patient serum<sup>61 62</sup>. Non-invasive imaging approaches for the detection of

2-HG in patients with glioma have been reported<sup>63 64</sup>, but their utility for clinical practice and clinical drug development remains to be defined.

Preclinical studies demonstrated that inhibition of mutant IDH enzymes retards tumor growth in experimental models of glioma, leukemia, and cholangiocarcinoma<sup>65-67</sup>.

The clinical development of inhibitors of mIDH proceeded most expeditiously for AML where, unlike in glioma, *IDH2* mutations are more common than *IDH1* mutations. Enasidenib, the first-in-class inhibitor of mIDH2, produced clinical responses in approximately 40% of patients with advanced mIDH2 AML<sup>68,69</sup>. Ivosidenib, the first-in-class inhibitor of the mIDH1 enzyme, similarly induced remissions in patients with advanced mIDH1 AML<sup>70</sup>. Both drugs have received regulatory approval for the treatment of mIDH AML.

A phase I study with ivosidenib in subjects with mIDH1 advanced solid tumors, including previously treated glioma ([ClinicalTrials.gov](https://clinicaltrials.gov/ct2/show/study/NCT02073994) identifier: [NCT02073994](https://clinicaltrials.gov/ct2/show/study/NCT02073994)), reported no dose-limiting toxicities, and the maximum tolerated dose was not reached. A dose of 500 mg once daily was selected for expansion based on the pharmacokinetic/pharmacodynamic data from all solid tumor cohorts. This trial showed early signs of clinical activity in IDH1-mutant glioma, with a reduction in tumor volume growth rates (i.e., compared with pretreatment growth rates) and tumor shrinkage in several patients<sup>71</sup>. In patients with IDH1-mutant advanced cholangiocarcinoma, ivosidenib was also well tolerated and showed preliminary evidence for antitumor activity<sup>72</sup>. The clinical benefit of targeting IDH1 mutations in advanced, mIDH1 cholangiocarcinoma was subsequently confirmed in a Phase 3 trial<sup>73</sup>.

Vorasidenib (AG-881) is a first-in-class, dual inhibitor of mIDH1 and mIDH2 that was developed for improved penetration across the blood-brain barrier<sup>74</sup>. In a phase I study ([ClinicalTrials.gov](https://clinicaltrials.gov/ct2/show/study/NCT02481154) identifier: [NCT02481154](https://clinicaltrials.gov/ct2/show/study/NCT02481154)), Vorasidenib showed a favorable safety profile at doses <100 mg QD in previously treated patients with non-enhancing glioma. Many patients remained on treatment after several years of continuous treatment and tumor shrinkage was observed in multiple patients with non-enhancing glioma<sup>75</sup>. In a follow-up perioperative phase I study in patients with non-enhancing glioma ([ClinicalTrials.gov](https://clinicaltrials.gov/ct2/show/study/NCT03343197), [NCT03343197](https://clinicaltrials.gov/ct2/show/study/NCT03343197)), vorasidenib 50 mg QD resulted in >90% reduction in intratumoral 2-HG concentrations compared with untreated controls, indicating near complete inhibition of the enzyme<sup>60</sup>.

Since a watch-and-wait approach following surgery remains a treatment option for patients with low-risk LGG, there is an opportunity to explore the activity of mIDH inhibitors during the active observation period. Vorasidenib (50 mg QD) is now being tested versus placebo in the ongoing, randomized, phase III INDIGO study ([ClinicalTrials.gov](https://clinicaltrials.gov/ct2/show/study/NCT04164901), [NCT04164901](https://clinicaltrials.gov/ct2/show/study/NCT04164901)) which enrolls patients with grade II non-enhancing mIDH glioma treated with surgery only.

Several other inhibitors targeting the mIDH enzymes are in earlier stages of clinical development for mIDH human cancers, including glioma.



## Other Therapeutic Approaches for mIDH Glioma.

Clinical and preclinical studies conducted since the first discovery of IDH mutations in cancer have provided deeper insights into the pathogenesis of IDH mutated human cancer and uncovered alternative and potentially complimentary approaches to exploit the effects of IDH mutations on cellular metabolism, epigenetic regulation and immune function:

### Targeting Tumor Metabolism:

D-2-HG directly and indirectly influences multiple and diverse metabolic intracellular events, but the myriad of interactions, the specific and most critical oncogenic driving events have yet to be elucidated. IDH mutated glioma cells are prone to oxidative stress and which is associated with increase reactive oxygen species (ROS) <sup>76</sup>. Strategies involving activation of antioxidant pathways, including glutathione synthesis inhibition, have been proposed. Metabolic reprogramming characterized by increased oxidative metabolism in the Krebs cycle, with suppression of reductive glutamine metabolism are hallmarks of IDH mutation. This increased rate of reductive glutaminolysis in preclinical models <sup>77</sup> provides the rationale for evaluating the safety and efficacy of the oral glutaminase inhibitor CB-839 in combination with radiation and temozolomide in mIDH glioma ([NCT03528642](#)).

### PARP Inhibitors:

PARP1 (and other PARPs) play critical roles in the repair of DNA single-strand breaks (SSBs) through several mechanisms that include base excision repair, nucleotide excision repair, and other DNA damage response pathways <sup>78</sup>. PARP inhibition leads to persistence of unrepaired SSBs and cytotoxic PARP-DNA complexes, which leads to the formation of potentially lethal DNA double-strand breaks (DSBs) <sup>79</sup>. Cells with deficient homologous recombination, the main compensatory mechanism to manage the increased DSB stress imposed by PARP inhibition, are unable to efficiently repair these DSB and subsequently enter mitotic catastrophe and apoptosis <sup>80</sup>. Recent evidence suggests that 2-HG produced by mIDH enzymes causes homologous recombination (HR) processes to become dysfunctional <sup>81</sup>, with evidence of activation of compensatory PARP-driven base excision repair mechanisms <sup>82,83</sup>. This has raised interest in exploring PARP inhibitors such as olaparib ([NCT03212274](#)) or BGB-290 ([NCT03749187](#)) for the treatment of mIDH gliomas.

### DNA Demethylation Agents:

Another major consequence of IDH mutations and 2-HG accumulation is the inhibition of various components of the epigenetic machinery including histone and DNA demethylases (DNMTs). This leads to aberrations in numerous biological processes that result in the glioma-CpG island methylator phenotype (G-CIMP), characterized by genome-wide DNA hypermethylation <sup>84</sup>. Restoring epigenetic programming via DNA demethylation is a current research strategy in IDH mutant glioma. The two prototypal DNA-demethylating agents, decitabine (DAC, trade name Dacogen, Eisai) and 5-azacitidine (AZA, trade name Vidaza, Celgene), are FDA-approved for treating patients with myelodysplastic syndrome. These drugs are cytidine analogs that incorporate into the DNA in the case of both agents, and RNA in the case of azacitidine, and form an irreversible covalent bond with DNMTs triggering the ubiquitin-dependent degradation of the enzymes. Both DAC and AZA have



short half-lives and poor *in vivo* stability due to their rapid deamination by the ubiquitously expressed cytidine deaminase (CDA). Overcoming this barrier for sustained and effective therapy of DNA methylation inhibition is being addressed in a clinical study of ASTX727, which consists of DAC and E7727 (cedazuridine), a novel CDA, in IDH mutant glioma (NCT03922555). The value of these strategies has yet to be demonstrated clinically.

### Immunotherapy Approaches:

Recent studies showed that the most common form of the mIDH enzyme in glioma (IDH1R132H) is presented on human MHC class II and induces mutation-specific CD4+ antitumor T cell responses<sup>85</sup>. The opportunity for peptide-based vaccination strategies using mutation-specific peptides has been evaluated in early clinical trials demonstrating safety and immunogenicity<sup>86</sup>. In addition, two pilot studies in LGG patients are exploring the neoadjuvant administration of vaccines with immune modulatory adjuncts. Several studies suggest that mIDH and the 2-HG oncometabolite may play critical roles in shaping the immunological landscape of the tumor microenvironment. *IDH*-mutation in glioma appears to be associated with impaired T-cell recruitment and T-cell receptor signaling, decreased tumor-infiltrating lymphocytes, and reduced programmed death-ligand 1 (PD-L1) expression<sup>87-93</sup>. These observations have prompted initial clinical trials involving checkpoint inhibition with avelumab (NCT02968940) and pembrolizumab (NCT02658279). These studies illustrate the interest in modulating the immune response using check point inhibitors in IDH mutated tumors as a single agent, but also in combination with peptide vaccination strategies. However, the role and efficacy of immunomodulatory therapies in treatment of mIDH glioma remain open questions.

### Evaluation of Treatment Response and Tumor Growth in mIDH Glioma.

Treatment efficacy in oncology is traditionally assessed with survival endpoints, in particular overall survival. This is assumed to present the most reliable endpoint reflecting ultimate patient benefit. For some tumors, the use of progression-free survival is a well-established surrogate for overall survival, and, in individual patients, response or absence of progression to a particular treatment is taken as evidence of benefit to that treatment.

This general approach to assess efficacy is less straightforward in mIDH lower grade glioma for several reasons. First, many of these patients are for many years clinically asymptomatic apart from usually well controlled seizures, and maintaining that status is clinically relevant. Secondly, these tumors are slowly but continuously growing entities if left untreated<sup>94</sup>. That implies that the tumor will progress after surgery, from the first day after surgery, and the date of progression is arbitrarily based on the cut-off that is taken to define progression on imaging. Thirdly, response assessment in glioma is usually based on changes in enhancement which is an indirect and a-specific measure of tumor growth and not applicable to non-enhancing tumors. Changes in non-enhancing tumor volume are often limited and may appear late, even after the end of lengthy chemotherapy cycles<sup>95</sup>. Lastly, radiotherapy and surgical effects may induce areas with increased signal intensities on T2 weighted and FLAIR MR images that are similar to radiographic changes associated with tumor progression, challenging the distinction between these two opposite conditions.

To address the many differences between high and low grade gliomas specific RANO criteria for outcome and assessment of low grade glioma have been proposed which incorporate measures for seizure and cognition assessment<sup>96</sup>. RANO response criteria to treatment for unenhancing low grade glioma are built on the classical Macdonald's criteria, with a 50% reduction of unenhancing area qualifying for response and 25% increase for progression. With modern computer technology, it has become feasible to assess (changes in) tumor volume and relate that to outcome in a semi-automated manner<sup>45</sup>. This has resulted in new ways of evaluating outcome, like the assessment of change in volumetric growth rate during treatment<sup>97</sup>. It will take however a review of large prospective datasets to validate such endpoints, preferably of homogeneously treated patients.

For everyday clinical practice, looking at change in tumor size will remain the standard approach for many years to come. With that in mind, it is important to realise the confusion that may arise of enhancing pseudoprogression after radiotherapy, and white matter changes after extensive surgery and after radiotherapy which both may give the false impression of tumor progression<sup>98</sup>.

## Future Directions in mIDH Glioma.

Despite the progress made in the understanding of the prognostic significance and altered cellular events in mIDH gliomas, many unanswered questions remain regarding the specific oncogenic mechanisms resulting from this alteration. In order to improve the outcome of these patients, it will be important to delineate the most significant oncogenic driving mechanisms in mIDH glioma and translate key findings to targeted and combined treatment/modality strategies.

In the clinical arena, there are several unanswered questions with respect to the best therapeutic compounds beyond standard radiation and chemotherapy, the role of combination strategies, and optimal timing of therapy and sequencing of treatments. Some of these questions are articulated below:

### (1.) What are relevant endpoints in the evaluation of novel agents for mIDH glioma?

A major issue in low grade glioma is the assessment of survival from a functional perspective. Cognition is often impaired in glioma patients, and therapy (radiotherapy, surgery) may contribute to that. A particularly notorious delayed effect of radiotherapy is delayed leuko-encephalopathy associated with decreased memory function and attention span<sup>99,100</sup>. It is unclear to what extent radiotherapy delivered with modern techniques induces this side-effect. Given the relatively favorable prognosis of most lower grade glioma patients, the challenge is not only long survival, but also survival with a good quality of functioning in the post-treatment period. Surviving without cognitive deficits is of vital interest to patients, and some ongoing studies have this as a primary endpoint but data from these trials and a critical evaluation of this type of endpoint is still lacking. An impossible to answer question is what level of change in a cognitive test equals some loss in overall survival. Seizures are in general better controlled after extensive surgery, radiotherapy and chemotherapy. Although recurrent seizures do not necessarily indicate tumor progression, being seizure free obviously is a matter of importance for patients and

does reflect a relevant clinical endpoint<sup>101</sup>. Lastly, with the addition of chemotherapy to radiotherapy, survival from the start of these treatments in low grade mIDH glioma and anaplastic oligodendroglioma is beyond 14 years<sup>56</sup>. From a practical perspective, the development of alternative endpoints that reflect patient benefit are urgently needed and are being explored in the CODEL and POLA studies. Such endpoints could be radiological, provided a validated relationship with ultimate clinical patient benefit is demonstrated, or cognitive functioning, assuming that this might best reflect the patient's well-being.

**(2.) Is there a role for direct inhibition of IDH and, if so, at what stage of the disease?**

While the mIDH enzyme likely plays a critical role in the initiation of mIDH low-grade gliomas, its contribution to the relentless growth of fully developed diffuse gliomas remains to be proven. The current experience with the mIDH inhibitors ivosidenib and vorasidenib suggests that this contribution might be greatest at the earlier disease stage. In contrast, ivosidenib and vorasidenib showed no clear antitumor activity in patients with enhancing tumors. The lack of single-agent antitumor efficacy for ivosidenib or vorasidenib in patients with enhancing gliomas may be due to the presence of additional genetic alterations in these tumors that can bypass the need for the mIDH enzyme for tumor maintenance.

**(3.) Can treatment be delayed for a specific subgroup of patients with mIDH glioma?**

At this time, there are several additional markers, such as 1/p/19q codeletion status, CDKN2A/B, TP53, ATRX, TERT mutation, TMB, methylation status of the MGMT gene promoter, and clinical factors including conventional histologic grade, performance status and extent of resection, which can influence the prognosis of IDH MT patients, and new genomic alterations continue to be discovered. It is conceivable that in the future a subgroup of better prognosis patients may be identified in which radiation therapy, or radio chemotherapy, can be delayed; conversely, poor prognostic patient groups may be identified who should receive earlier and aggressive treatment. It is likely that such factors will be identified and utilized as grouping or stratification factors, or eligibility for entry in future clinical trials. The other possibility is that patients with more favorable prognosis derive greater benefit from early aggressive treatment, and that patients with more unfavorable prognosis fare worse regardless of what is tried. Answering these important questions will require carefully collected and molecularly annotated datasets.

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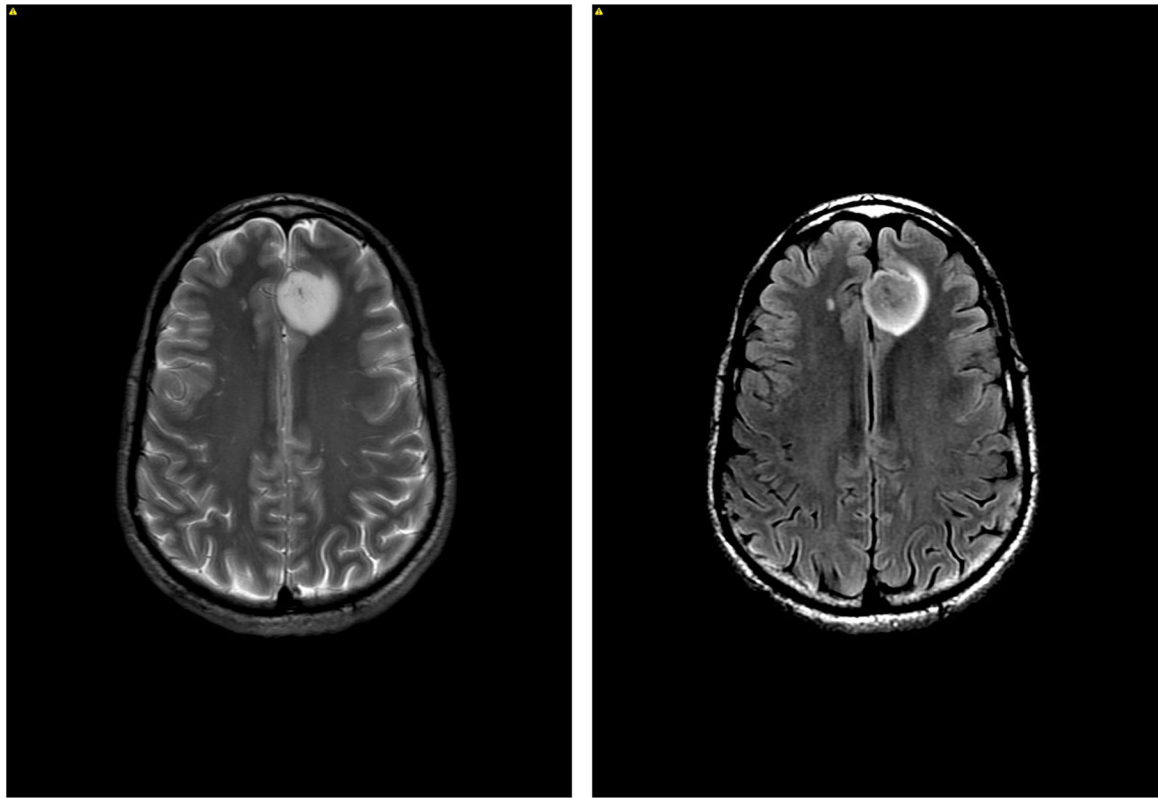
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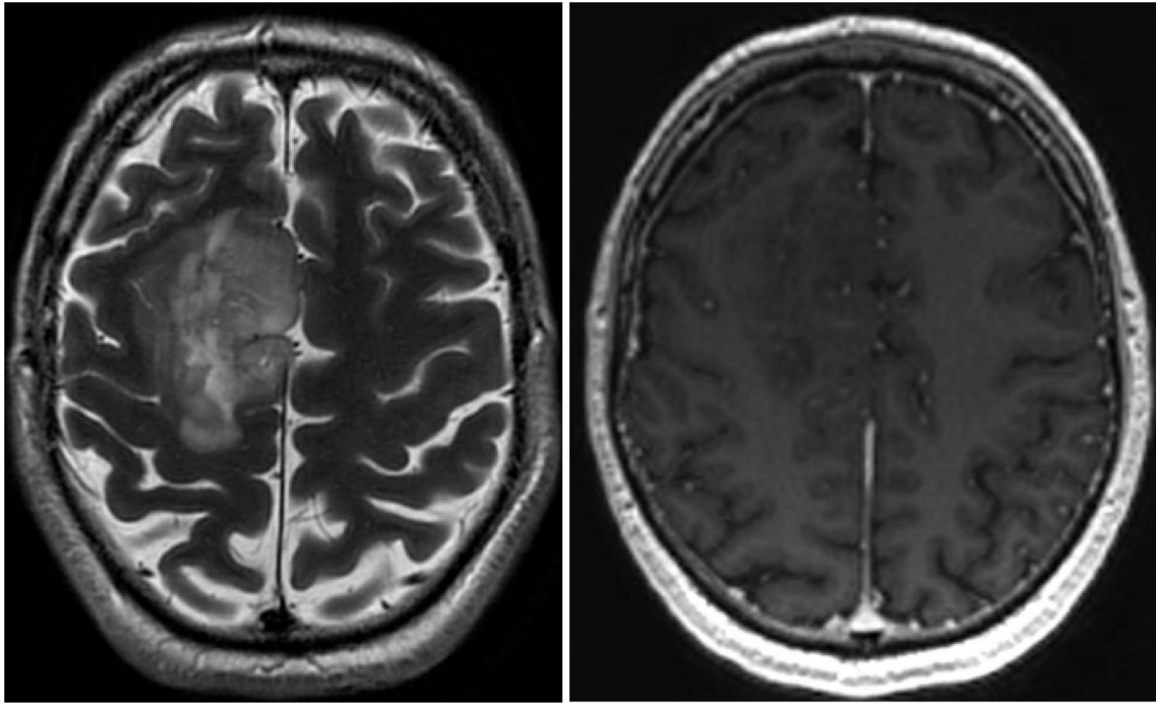


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**Figure 1. “T2-FLAIR mismatch” sign.**

Shown is a MRI Brain of a 35 year-old man with a histologically-confirmed mIDH 1p/19q non-codel WHO Grade II glioma. Left: T2-weighted sequence showing a hyperintense, relatively homogeneous area; Right: FLAIR sequence with hyperintense rim surrounding a relatively hypointense central area.



**Figure 2. Low grade oligodendroglioma.**  
Shown are Brain MRI T2 (left image) and T1 after intravenous contrast (right image), showing inhomogeneous lesion with cortical involvement and no contrast uptake.

### Progression Free Survival

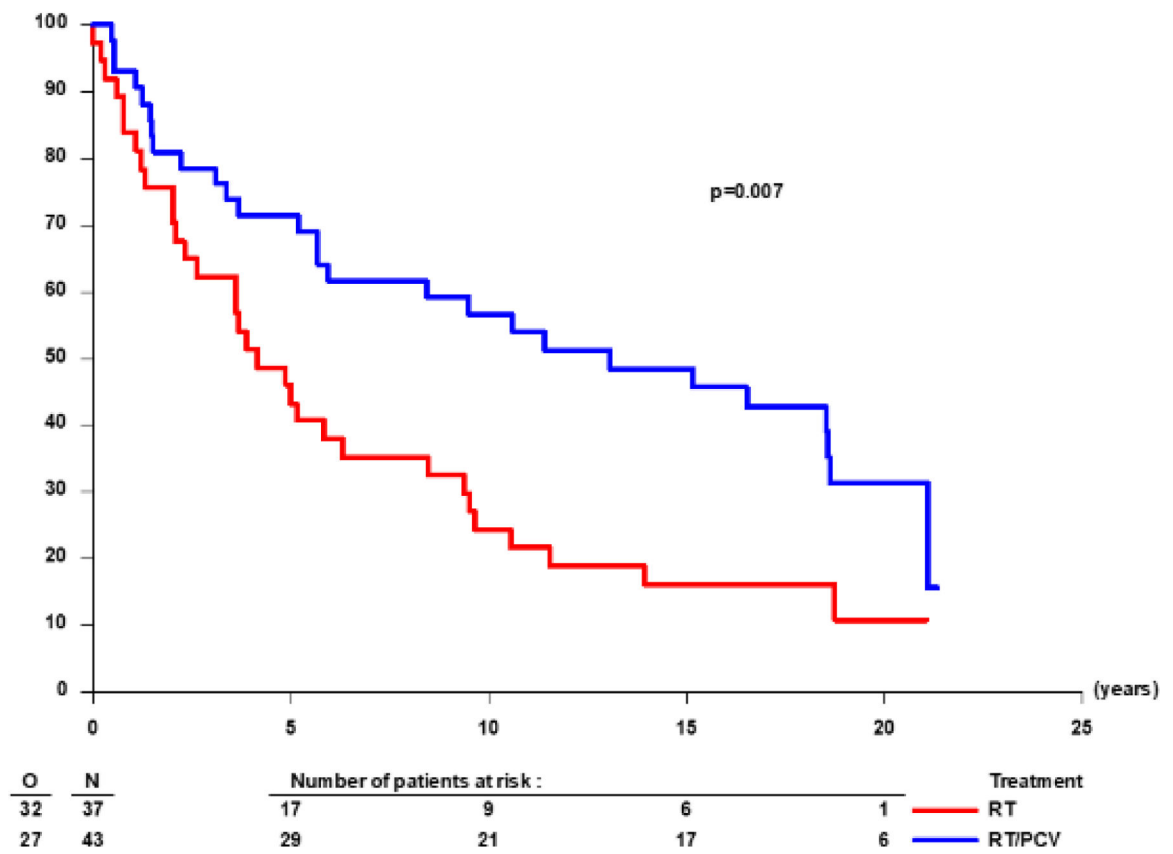


Figure 3. Long-term follow-up of Progression-free survival (PFS) of RT/PCV versus RT alone in 1p/19q codon glioma patients (n=80) in EORTC 26951.