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## Investigational New Drugs for Brain Cancer

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### Abstract

**Introduction**—Despite substantial improvements in standards of care, the most common aggressive pediatric and adult high-grade gliomas (HGG) carry uniformly fatal diagnoses due to unique treatment limitations, high recurrence rates and the absence of effective treatments following recurrence. Recent advancements in our understanding of the pathophysiology, genetics and epigenetics as well as mechanisms of immune surveillance during gliomagenesis have created new knowledge to design more effective and target-directed therapies to improve patient outcomes.

**Areas covered**—In this review, the authors discuss the critical genetic, epigenetic and immunologic aberrations found in gliomas that appear rational and promising for therapeutic developments in the present and future. The current state of the latest therapeutic developments including tumor-specific targeted drug therapies, metabolic targeting, epigenetic modulation and immunotherapy are summarized and suggestions for future directions are offered. Furthermore, they highlight contemporary issues related to the clinical development, such as challenges in clinical trials and toxicities.

**Expert opinion**—The commitment to understanding the process of gliomagenesis has created a catalogue of aberrations that depict multiple mechanisms underlying this disease, many of which are suitable to therapeutic inhibition and are currently tested in clinical trials. Thus, future treatment endeavors will employ multiple treatment modalities that target disparate tumor characteristics personalized to the patient's individual tumor.

### 1. The Diverse Spectrum of Malignant Gliomas

In the United States, approximately 23,000 people are diagnosed with a malignant brain tumor each year [1], of which approximately 80% belong to the heterogeneous group of diffuse gliomas [1]. Histologically, gliomas are graded into individual classes as defined by the World Health Organization (WHO) classification of central nervous system (CNS)

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#### Declaration of Interest

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tumors based on architectural and cytological features, such as cellular atypia, mitotic activity, necrosis and vascular proliferation [2]. Clinically, we frequently distinguish between low-grade (LGG, grade II) and high-grade (HGG, grades III and IV) tumors to reflect their anticipated biologic behavior and clinical course. Over time, LGGs frequently progress to higher-grade projecting the natural course of this disease rather than a new entity. Three main glioma subtypes exist, which are based on the morphological similarities of the predominant tumor cell population; the most frequent subtypes are astrocytoma (~75%, WHO grade I–IV), followed by oligodendroglioma (~6%, WHO grade II–III) and ependymoma (~7%, WHO grade II–III), which resemble astrocytes, oligodendrocytes and ependymal cells, respectively [1].

Glioblastoma (GBM, grade IV astrocytoma) is the most malignant variant of diffuse astrocytoma in adulthood accounting for 55% of all gliomas [1]. The majority (95%) of GBM arise as *de novo* lesions (i.e. primary GBM). Less common are secondary GBM (~5%) that arise from the progression of grade II or III glioma. Secondary GBM is a genetically and clinically distinct entity that typically occurs in younger patients (mean age 45 years versus 60 years for primary GBM) [1 3].

Oligodendroglioma comprises the second most common glioma in adults [2]. The hallmark molecular abnormality that is now increasingly used to define oligodendroglioma is co-deletion of chromosomes 1p and 19q, which also predicts a better prognosis and responsiveness to radiation therapy and chemotherapy [4].

Brain cancer has superseded leukemia as the most common cause of cancer-related death in children [1,5]. As in adults, gliomas are the most common CNS neoplasms in children accounting for ~53% of all tumors although the predilection sites differ. The majority of childhood gliomas (~60%) are grade I or II lesions and of astrocytic lineage, while oligodendroglioma and ependymoma are rare [5]. Childhood HGG traditionally include GBM, anaplastic astrocytoma, anaplastic oligodendroglioma and diffuse intrinsic pontine glioma (DIPG) [5,6]. DIPG is a childhood specific brain cancer that presents most commonly between ages 6–8. Despite varying histological grades DIPGs share a universally lethal outcome [7,8]. In contrast to other HGGs, the diagnosis of DIPG has relied solely upon radiological and clinical findings, and not on histopathological features. However, a role for image-guided stereotactic biopsy has recently been re-introduced toward the goals of sample collection for biological studies and molecular characterization required for experimental personalized therapeutics [9].

Aside from radiation exposure and uncommon inherited genetic syndromes, there are few proven causes of primary brain tumors beyond the role of random spontaneous mutation associated with physiologic DNA replication. About 5% of gliomas are due to inherited germ-line mutations associated with known syndromes such as Cowden's disease (PTEN mutation), tuberous sclerosis (TSC1 9q34, TSC2 16p13), Li–Fraumeni syndrome (p53 mutation), neurofibromatosis types 1 and 2 (neurofibromin and merlin mutations) and Lynch Syndrome (miss-match DNA repair defect) [10].

## 2. Current standards of treatment and outcomes

Tumor grade, histology and their effect on prognosis are the key factors influencing therapeutic decision-making. Non-infiltrative grade I gliomas have the most favorable prognosis and chance for cure due to their indolent nature and potential for complete surgical resection with 5- and 10-year survival rates of 94% and 92%, respectively [1]. Well-differentiated oligodendroglioma (WHO grade II) and anaplastic oligodendroglioma (WHO grade III) are known to be particularly responsive to cytotoxic therapies including radiation, temozolomide (TMZ), and the combination chemotherapy regimen PCV (Procarbazine, CCNU, Vincristine). Two recent randomized Phase III trials, initiated prior to the introduction of TMZ, definitively demonstrated the benefit of aggressive combination radiation + chemotherapy for anaplastic oligodendroglioma [11,12]. Patients treated with PCV + radiation lived remarkably longer than patients treated with radiation therapy alone (~15 vs. 7 years, median survival). These studies have all but established a new standard of care that combines radiation and PCV chemotherapy for patients with anaplastic 1p19q co-deleted oligodendroglioma. An informative sub-analysis in these studies revealed no statistically significant benefit from combination PCV + radiation compared to radiation alone in the non-codeleted tumors that represent anaplastic astrocytoma. A randomized trial comparing combination radiation + TMZ with combination radiation + PCV is currently underway (CODEL study, NCT00887146). Results are anxiously awaited since anaplastic oligodendroglioma is known to be sensitive to TMZ, which is less toxic, more easily tolerated and more easily administered than PCV.

In contrast, HGGs of the astrocytic lineage are less sensitive to therapy and have a drastically lower median survival. Although complete surgical resection of these infiltrative tumors is virtually impossible, retrospective studies increasingly point to a probable survival benefit from > 70% tumor resection with clear survival benefit following gross total resection [13,14]. Surgery is followed by fractionated radiotherapy with concomitant and adjuvant treatment with the alkylating agent temozolomide (TMZ), a regimen that was established in GBM as the gold standard in 2005 [15]. In this study, the 2-year survival of GBM patients increased from ~8% to 28% [1]. Many neuro-oncologists also recommend the use of TMZ for the treatment of anaplastic astrocytomas; however, a clear benefit has not been convincingly demonstrated and is subject of an ongoing Phase III trial by the *European Organisation for Research and Treatment of Cancer* (EORTC, CATNON intergroup trial (NCT00626990)). The strongest predictive and prognostic factor for a therapeutic response and a longer survival is the epigenetic silencing of the DNA repair enzyme O-6-methylguanine-DNA methyltransferase (MGMT) [14–17]. Nonetheless, almost all GBMs ultimately relapse, and there are no treatments proven to prolong survival in patients with recurrent GBM. This is exemplified by recent disappointing results showing no significant survival benefit from the neutralizing anti-VEGF monoclonal antibody bevacizumab (Avastin®), administered either alone or in combination with existing cytotoxic regimens, which was granted accelerated approval in 2009 based on its ability to improve MRI abnormalities and reverse patient symptoms in patients with recurrent GBM [18–20].

For pediatric DIPG patients, treatment options are even more limited with no conventional chemotherapeutic agents proven effective, and radiation therapy alone representing the standard of care resulting in a survival rate of less than 10% 2 years after diagnosis [8,21].

In spite of extensive efforts to optimize existing therapies, we have likely reached the limits of what can be achieved with traditional cytotoxic regimens. Drawing from the advancements in understanding the underlying biology, new nodes of vulnerabilities have been identified that have paved the way for novel target-directed therapies, onco-metabolic and epigenetic focused strategies, as well as new immunotherapeutic approaches (Fig. 1, Tab. 1). Especially exciting have been the advances made in next-generation immunotherapies, which, in principle, can affect every single tumor cell by simultaneously or sequentially evoking specific immune responses against a theoretically unlimited number of tumor-associated antigens. In this review, we summarize the current state of these newly emerging therapeutic developments and highlight major remaining obstacles and challenges for these approaches.

### 3. Emerging Targeted Therapy for Malignant Glioma

#### 3.1 Receptor tyrosine kinase (RTK) Pathway Inhibitors

Nearly all HGGs show genetic alterations in the receptor tyrosine kinase (RTK)–PI3K–RAS pathway, although specific aberrations and frequencies greatly differ between adults and children [22,23]. RTKs are a family of cell surface receptors that include platelet-derived growth factor receptors (PDGFR- $\alpha$  and PDGFR- $\beta$ ), epidermal growth factor receptors (ErbB1/EGFR, ErbB2, ErbB3, ErbB4), vascular endothelial growth factor receptors (VEGFR1–4), fibroblast growth factor receptors (FGFR1, FGFR2) and c-MET. Oncogenic RTK hyperactivation is driven by multiple mechanisms such as RTK gene amplification and overexpression resulting in ligand-independent receptor oligomerization, receptor mutation resulting in constitutive activation and ligand over-expression. Dysregulated RTK hyperactivation drives multiple oncogenic processes such as cell proliferation, aberrant survival and therapeutic resistance, migration/invasion, and tumor cell stemness that is closely linked to tumor propagating capacity [24–26].

In adult HGGs, epidermal growth factor receptor (EGFR) EGFR is the most frequently amplified gene, with approximately one-third of GBMs also having gene rearrangements [27]. The most common is EGFR variant III (EGFRvIII), a truncated 801 base pair in-frame deletion of the wild-type EGFR, which is tumor-specific and absent in healthy tissue. Despite this frequency and that fact that EGFR inhibitors are efficacious in other molecularly defined malignancies (e.g. EGFR mutated non-small cell lung cancer) [28], multiple EGFR inhibitors (erlotinib, gefitinib, lapatinib and the chimeric EGFR monoclonal antibody cetuximab), alone or in combination, have been largely ineffective against HGG in multiple clinical trials [29–39], attributed to the absence of the kinase domain mutations required for durable therapeutic responses, insufficient CNS drug penetration or toxicity, particularly when used in combination [23,40,41]. For example, lapatinib, a dual RTK inhibitor that also targets erbB2, was investigated as a single agent for recurrent GBM and discontinued prematurely due to lack of clinical effect, owing to sub-therapeutic tissue concentrations [37]. Likewise, the combined use of lapatinib with pazopanib, a multiple

kinase inhibitor against VEGFR1–3, PDGFR $\alpha/\beta$  and c-Kit, failed to show any effectiveness in a Phase I/II study [42], while a follow-up Phase I study of lapatinib + TMZ in recurrent HGGs showed a slight effect on progression-free survival, albeit with moderate toxicity [43]. This gave rise to the current Phase II trial evaluating high-dose lapatinib in conjunction with standard chemo-radiation in patients with newly diagnosed GBM (NCT01591577).

Despite such disappointing results to date, the development of new EGFR inhibitors continues to be of high interest due to its critical role in gliomagenesis, which is also underscored by the fact that the majority of agents evaluated in clinical trials for glioma patients have targeted one or multiple components of this signaling pathway. The novel agent Dacomitinib (PF-299804), an irreversible pan-HER inhibitor, was found to have promising activity in preclinical EGFR mutated glioma models [44] and is currently being evaluated in two Phase II studies for recurrent GBM patients with EGFR amplification or EGFRvIII expression (NCT01520870, NCT01112527). Afatinib (BIBW 2992) is another novel dual EGFR/Her2 inhibitor that had no meaningful single-agent activity in non-selected recurrent GBMs, but perhaps a modest response in patients with EGFRvIII mutations [45]. Although standard daily dosing was inactive, alternative dosing schedules such as pulsatile-increased dosing may be evaluated in the future, as observed in other cancer types.

Conjugated and unconjugated antibodies have also been developed to target both wildtype EGFR and EGFRvIII, including cetuximab, panitumumab, and nimotuzumab. Although none of them have been able to demonstrate a significant survival advantage in patients with recurrent GBM, several new agents are currently undergoing clinical trial evaluations: AMG 595, a first-in-human mAb for patients with EGFRvIII recurrent HGG (NCT01475006); Sym004, a recombinant antibody that specifically binds to EGFR in patients with EGFR amplified GBM (NCT02540161); and ABT-414, an antibody-drug conjugate, with a toxic payload (monomethylauristatin F) targeted to EGFR or EGFRvIII amplified GBMs (Intelligence 1; NCT02573324). ABT-414, which is currently undergoing evaluation in randomized, placebo-controlled Phase IIb/III with concomitant radiotherapy and temozolomide (TMZ), demonstrated preliminary efficacy in Phase I, however toxicities affecting the eye and liver were observed as well [46].

Small molecule tyrosine kinase inhibitors with anti-PDGFR activity, which is amplified in 10% of HGGs [47], include imatinib mesylate, the dual VEGFR/PDGFR inhibitors (sunitinib, sorafenib, pazopanib, and vatalanib), and dasatinib that targets BCR-ABL, the Src family, c-Kit, and PDGFR- $\alpha/\beta$ . PDGFR inhibitors have also largely disappointed in the treatment of HGGs despite encouraging preclinical results. For example, imatinib, which was investigated as monotherapy and in combination with hydroxyurea in newly diagnosed and recurrent GBM did not improve outcome [48–50]. Likewise, a Phase II clinical trial of pazopanib in patients with recurrent GBM failed, reporting a median progression-free survival and overall survival of 12 and 35 weeks, respectively [51]. Furthermore, dasatinib was ineffective in multiple Phase II clinical trials as a single agent and in combination with radiation, TMZ, erlotinib, lomustine, as well as crizotinib [52–57]; the results of an ongoing randomized Phase II trial adding dasatinib to bevacizumab in recurrent disease (Alliance N0872, NCT00892177) are anticipated shortly. Dose-limiting toxicities, particularly when used in combination with other cytotoxic agents (e.g. lomustine), and poor CNS

bioavailability due to the fact that dasatinib is a substrate for P-glycoprotein and breast cancer resistance protein (BCRP) have contributed to its poor efficacy [58–59]. The novel PDGFR $\alpha/\beta$  and Flt3 inhibitor crenolanib is currently in Phase II for adult HGG (NCT01229644) and in a Phase I study for recurrent pediatric HGG (NCT01393912).

Other inhibitors target c-MET or its ligand hepatocyte growth factor (HGF), which, when overexpressed promotes angiogenesis, proliferation and invasion of cancer cells and correlates with poor prognosis [60–62]. Preclinical studies have conferred sensitivity to c-Met inhibitors in glioblastomas xenograft models [60,63]. Cabozantinib, which targets Met and other kinases, failed in the treatment of primary and recurrent GBM, although patients were not pre-selected for c-Met amplification [64]. Further studies with AMG 102 (rilotumumab), a humanized monoclonal antibody against HGF, and SGX523, a direct inhibitor of c-MET, evaluated in patients with recurrent molecularly undefined GBMs were also unsuccessful in a Phase II and two Phase I trials (NCT00606879, NCT00607399) respectively [65]. Met-specific inhibitors are in clinical development, including onartuzumab, a monoclonal antibody targeting c-Met that is currently being evaluated in Phase II combination with bevacizumab in recurrent GBM (NCT01632228).

In contrast to the RTKs discussed above that are primarily expressed by glioma cells, VEGFRs are primarily expressed by glioma vascular endothelial cells and ligand-dependent activation drives tumor-associated vascular proliferation (i.e. angiogenesis) and BBB dysfunction. Bevacizumab, a mAb against the VEGF-A ligand, is the most extensively studied and successful VEGFR pathway inhibitor. Bevacizumab was FDA-approved for recurrent GBM based on objective imaging response data with no evidence of survival benefit (see Section 2, above). Two large Phase III trials have since failed to demonstrate any survival benefit for patients with newly diagnosed GBM [18,20,66]. Furthermore, the Phase III EORTC-26101 trial, investigating the therapeutic efficacy of the combination bevacizumab and lomustine versus lomustine alone in GBM patients at first recurrence, did not improve the neurological deterioration-free survival or the overall survival [67]. Apart from mAbs, various small molecule inhibitors directed against VEGFRs have been developed including cabozantinib, cediranib, pazopanib, sorafenib, sunitinib, vandetanib and vatalanib; however, clinical trials have been uniformly disappointing. Several Phase II trials of sorafenib and sunitinib in combination with other therapies in GBM have failed to meet primary endpoints [68–75]. Cediranib, which targets all VEGFRs, c-kit and PDGFR, initially showed modest potential efficacy as monotherapy in a Phase II trial in recurrent GBM patients [76]. However, these results were not substantiated in a subsequent randomized Phase III trial comparing cediranib monotherapy, lomustine monotherapy and combination cediranib and lomustine [77]. Lenvatinib (E7080), a novel multitargeted tyrosine kinase inhibitor for all VEGFRs and PDGFR, did not warrant further testing in recurrent GBM patients beyond Phase II because the therapeutic effect was very limited with no objective responses (NCT01137604) [78]. Finally, axitinib, a TKI with high affinity and specificity for the VEGF-receptors that approved for the treatment of metastatic renal cell carcinoma, did not result in any meaningful treatment response in patients with recurrent GBM who had failed surgery, radiation, and TMZ [79].



The failures of RTK pathway inhibitors in HGG have been especially frustrating in light of strong evidence supporting the molecular basis for RTK pathway targeting and mounting successes in other malignancies. Several potential explanations exist for these treatment failures in GBM. Most clinical trials have not focused sufficiently on patient cohorts with tumors enriched for target hyperactivation. Both parallel signaling pathways and bypass feedback loops can contribute to intrinsic resistance to target inhibition. Small molecule RTKs are typically hydrophilic resulting in poor drug bioavailability in the CNS due to the blood-brain barrier and insufficient target inhibition. Moreover, multiple surveys have shown substantial intratumoral cellular heterogeneity for target expression and hyperactivation, which adds another layer of complexity [80,81]. As such, future clinical trials will have to focus on stringent patient selection criteria (genotype-enriched clinical trials), customized drug design based on tumor characteristics, and drug dosing regimens rigorously proven to sufficiently inhibit target activation in patients. Combinations instead of single tyrosine kinase inhibitors may be required to sufficiently inhibit parallel bypass and feedback pathways.

### 3.2 PI3K/AKT and mTOR Pathway Inhibitors

Almost 90% of patients with GBM show at least one alteration in the phosphatidylinositol 3-kinase (PI3K) signaling, which can either result from activating mutations in PI3K itself (25%), loss of the tumor suppressor gene Phosphatase and Tensin Homolog (PTEN) (41%) or downstream activation of RTKs [47]. The downstream effectors of PI3K include AKT and mammalian target of rapamycin (mTOR) consisting of mTOR complexes mTORC1 and 2, which play key roles in cell metabolism, survival, and protein translation. Hence, the addition of pharmacological inhibitors should theoretically lead to therapeutic benefits; however, the majority of clinical trials have failed to fulfill this expectation.

Several first-generation mTOR inhibitors, including rapamycin, temsirolimus (CCI-779), and everolimus (RAD001) possess antineoplastic activity as single agents in vitro and in vivo, and have been studied in various Phase I/II trials for newly diagnosed and recurrent HGGs. Although radiographic responses were observed in a subset of HGG patients receiving everolimus or temsirolimus, no significant effect on progression-free survival and overall survival was reported for any of these inhibitors when used alone, in combination with standard chemoradiation or with bevacizumab in recurrent GBM [82–89]. Notably, the combined use of temsirolimus with TMZ and radiation therapy led to an increased infection risk, which prohibited the analysis of the efficacy of this combination [87]. Currently ongoing clinical trials focus on the evaluation of everolimus in the treatment of progressive LGGs in combination with TMZ (NCT02023905), as single agent in children with progressive LGGs (NCT01734512) and ependymoma (NCT02155920). In addition, dual mTORC1/2 inhibitor sapanisertib (INK1280) is in early clinical investigation to assess CNS penetration and response in patients with recurrent GBM (NCT02133183).

Among PI3K-targeting drugs, buparlisib (BKM120), a pan-PI3K inhibitor, has been studied in the treatment of recurrent GBM based on encouraging preclinical experiments in U87 glioma cells [90]. An initial Phase II trial achieved intratumoral drug concentrations sufficient to inhibit the PI3K pathway although efficacy was not reported. Ongoing studies

with buparlisib include a Phase I/II dose-escalation trial with concurrent standard chemoradiation in newly diagnosed GBM (NCT01473901), a Phase II trial in recurrent GBM (NCT01339052) and combination trials for recurrent GBM with bevacizumab (NCT01349660), and carboplatin or lomustine (NCT01934361). PX-886 (Oncothyreon) is a semi-synthetic derivative of wortmannin and irreversibly inhibits PI3K. In glioma cells, PX-866 dramatically inhibited proliferation, autophagy and angiogenic potential in a variety of cell lines and U87 mouse xenograft models [91]. A Phase II trial evaluating the efficacy and safety of PX-866 in patients with recurrent GBM was recently completed and results are soon to be published (NCT01259869). Other PI3K inhibitors currently undergoing clinical evaluation include XL-147 (NCT01240460).

GBM with mutant PI3K alleles may also benefit from inhibition of AKT. Among the various agents that have entered the clinical trial Phase perifosine (KRX-0401) is the most advanced. A Phase II trial of single agent perifosine failed in recurrent GBM (NCT00590954). As data suggesting that the combined inhibition of AKT and mTOR may be more effective in combination with temsirolimus, two Phase II trials have been launched (NCT01051557, NCT02238496).

### 3.3 RAF–MEK–ERK pathway inhibitors

The presence of activating mutations in constituents of the mitogen activate protein kinase (MAPK) pathway has raised considerable interest in inhibiting this pathway in patients with HGGs and also in patients with unresectable or recurrent LGGs, which typically harbor oncogenic activation of BRAF resulting from a BRAF<sup>V600E</sup> point mutation or a KIAA1549:BRAF fusion. Evidence for the effectiveness of BRAF and MEK inhibitors or combinations thereof comes from multiple pre-clinical data in murine intracranial xenografts of pediatric gliomas and a few anecdotal clinical case reports [92,93].

There are in fact several BRAF inhibitors available, but only few have been evaluated in CNS tumors due to poor BBB penetration. Vemurafenib (PLX4032), which has demonstrated remarkable activity against metastatic BRAF<sup>V600E</sup> mutated melanoma [94], is currently undergoing Phase I/II testing in patients with BRAF<sup>V600E</sup> positive recurrent or refractory pediatric glioma (NCT01748149). Additional agents currently in clinical trial for pediatric BRAF<sup>V600E</sup> mutated glioma are the BRAF inhibitor dabrafenib (GSK436) (NCT01677741) and the two MEK1/2 inhibitors trametinib (GSK212) (NCT02034110, NCT02124772) and MEK162 (NCT02285439). Notably, the BBB penetrance has not been sufficiently studied in the majority of these agents except for dabrafenib, which appears to achieve adequate CNS drug levels [95]. Isolated case reports detailing complete clinical responses to dabrafenib in children with BRAF<sup>V600E</sup>-mutant GBMs have sparked excitement [96]; however, responses are rarely durable due to MAPK reactivation [97,98]. Therefore, multiple research groups have proposed the co-administration with other MAPK pathway inhibitors, such as the MEK inhibitor trametinib, in an effort to delay the development of resistance and to minimize the toxicities, particularly cutaneous, associated with BRAF inhibition [97].

Current BRAF inhibitors are mutation specific explaining why BRAF<sup>V600E</sup> inhibitors are ineffective against other mutations or oncogenic BRAF fusions [93,99]. This limitation can



potentially be overcome in appropriate patient subsets using MEK inhibitors that target wild-type MEK1/2 kinase that is hyperactivated downstream of oncogenic BRAF driver mutations [99]. In addition to trametinib and MEK162, sorafenib (NCT02450149) and selumetinib (AZD6244) are under clinical investigation for this purpose. The latter, selumetinib, a second-generation MEK1/2 inhibitor, is undergoing Phase II evaluation in LGGs and neurofibromatosis 1 (NF1)-associated tumors, including optic pathway gliomas as well as plexiform neurofibromas (NCT01386450, NCT01089101) that has resulted in some encouraging therapeutic responses, particularly in patients with NF1 [100].

### 3.5 Inhibitors of Onco-Metabolism

Activation of oncogenes and loss of tumor suppressors promote metabolic reprogramming to enhance nutrient uptake and energy supply in order to sustain growth and survival. In diffuse gliomas, the premier oncogenic mutations that result in metabolic reprogramming are within the genes coding for isocitrate dehydrogenases, IDH1 and IDH2. Originally detected in primary GBMs (5–10%) [23], IDH1 mutations occur in 50–80% of grade II and III astrocytoma and oligodendroglioma, as well as secondary GBM, while IDH2 mutations are less common and are mutually exclusive with mutations in IDH1 [101,102]. The most common mutation in IDH1 is arginine 132 to histidine (R132H) that results in the ability to convert  $\alpha$ -ketoglutarate ( $\alpha$ -KG) to the R(-)-2-hydroxyglutarate (2-HG), a potential onco-metabolite, with the coincident conversion of NADPH to NADP<sup>+</sup>. Current evidence suggests that 2-HG alters the epigenetic machinery in glioma, affecting histone demethylases and DNA hydroxylases, leading to chromatin modifications and gene expression dysregulation [103].

There are currently no therapies approved targeting mutant IDH; however, multiple therapeutic approaches targeting mutant IDH1/2 and the altered metabolic pathway are currently in clinical and preclinical development stages. Because IDH1 mutation is involved in the inhibition of histone lysine demethylases, DNA methyltransferase (DNMT) inhibitors have been suggested as a potential therapeutic approach [104], including decitabine and 5-azacytidine that reverse cancer-promoting hypermethylation restoring cell differentiation and tumor suppressing functions [105,106]. Specific inhibitors of IDH1 R132H are AGI-5198 and ML309, which reduced 2-HG levels and significantly decreased the growth of IDH1<sup>R132H</sup>-expressing glioma cells in vitro and human glioma xenografts [107,108]. Similar compounds of mutant-selective inhibitors of IDH1 (AG-120), IDH2 (AG-221) or both IDH1/2 (AG-881) have entered the clinical trial Phase (NCT02073994, NCT02273739, NCT02481154) (Tab. 1). While we are awaiting the results for glioma patients, an ongoing Phase I trial of AG-120 in 62 patients with advanced IDH1-mutant solid tumors reported a partial response in 1/55, stable disease in 52.7% (29/55), and progressive disease in 38.2% (21/55) [109].

Dichloroacetate (DCA) represents another strategy for targeting the oncogenic effects of IDH mutation. DCA is an inhibitor of mitochondrial pyruvate dehydrogenase kinase and thus, increases intracellular levels of pyruvate dehydrogenase, which is reduced in IDH1 mutated tumors. DCA has been shown to reduce GBM cell proliferation and clonogenicity in preclinical animal models [110]. A small case series of 5 patients treated with oral DCA

confirmed the inhibition of pyruvate dehydrogenase kinase along with p53 activation inducing apoptosis in human GBMs [111] and gave rise to a Phase II trial for primary and recurrent GBM, albeit unselected for IDH1 status, which was recently completed (NCT00540176).

In contrast to IDH-mutated tumors, IDH wild-type HGGs adapt to the increased metabolic demands of tumor growth through alternative metabolic pathways dependent on branched-chain amino acids (BCAAs), which generate nitrogen for the synthesis of neurotransmitter glutamate and macromolecule precursors for mitochondrial ATP synthesis. The key enzyme involved in this process is branched-chain amino acid transaminase 1 (BCAT1) that is restricted to a small number of tissues, including the brain. Interestingly, expression of BCAT1 is dependent on intracellular levels of  $\alpha$ -KG, which is produced by wild-type IDH1 [112]. IDH1 expression inhibition was found to impair BCAA metabolism and reduce tumor cell proliferation and invasiveness in a GBM xenograft model [112]. Gabapentin, a leucine analog that specifically inhibits BCAT1 significantly reduces glutamate concentrations in U-87MG cells *in-vitro* [112]. As BCAAs are nutritionally essential in that humans cannot synthesize them endogenously, a diet deficient in BCAAs (e.g. ketogenic diet) might provide a unique way to downregulate BCAT1 and target these neoplastic metabolic processes [113]. Currently, there are multiple clinical trials under way investigating the feasibility of ketogenic dietary therapy as an adjunct to radiation and chemotherapy in adult patients with HGGs (NCT02046187, NCT02302235, NCT01754350, NCT01865162). Historically, the success of the ketogenic diet in brain tumor patients, reported as case reports or small case series, have been mixed [114,115]. A recently reported Phase I trial examined the feasibility of a ketogenic diet in 20 patients with recurrent GBM (ERGO trial; NCT00575146). Although well tolerated, no clinical activity was found when used as single agent [116]. Regardless, these data highlight that energy metabolism reprogramming drugs or diets might be an attractive treatment option for GBM patients with mutated or wildtype IDH1.

Acetate is another nutrient of increasing interest for its possible role in HGGs and other malignancies. Acetate ligation to coenzyme A (CoA) by acetyl-CoA synthetases (ACSSs) to form acetyl-CoA is critical for the synthesis of fatty acids (lipogenesis), nucleotides, amino acids, and histone acetylation. Cancer cells increase their dependence on acetate under conditions of hypoxia and nutrient stress. *Mashimo and Comerford* et al. found that primary brain tumors utilize exogenous acetate to meet their high metabolic demands and facilitate growth [117,118]. The incorporation of exogenous acetate is achieved by acetyl-CoA synthetase 2 (ACSS2) and its expression inversely correlates with survival of patients with HGGs. Knockdown of ACSS2 expression was found to dramatically impair the incorporation of exogenously supplied acetate into lipids and histone protein and increased animal survival. These findings have been validated in patients with HGGs and brain metastasis [117]. Selective ACSS2 inhibitors are currently under development and could offer a highly promising new approach to metabolic therapy since ACSS2 is dispensable in normal cells.

### 3.6 Targeting epigenetic modulators

Epigenetic alterations, i.e. the acetylation and deacetylation of histones allowing the regulation of gene expression patterns, play a crucial role in gliomagenesis. Histone acetylation leads to the unfolding of chromatin (euchromatin) and promotes increased gene transcription while deacetylation induces chromatin condensation (heterochromatin) and mediates suppression of transcription [119,120]. Histone deacetylases (HDAC), responsible for maintaining this balance are frequently overactivated in cancers and inhibition of HDAC in glioma cell lines and GBM animal models demonstrated anti-neoplastic activity [121,122]. Nonetheless, the role of HDAC inhibitors in the treatment of adult HGG remains questionable because most of the HDAC genes are, in fact, downregulated in GBM [123], and clinical trials with the HDAC inhibitors vorinostat and panobinostat failed [124,125]. Although a Phase I/II study of vorinostat in combination with TMZ or radiotherapy + TMZ demonstrated good tolerability in recurrent and newly diagnosed GBM, respectively, it did not result in a significant survival improvement. Likewise, Phase II studies evaluating the combinations of bevacizumab and vorinostat or bevacizumab and panobinostat in recurrent GBM revealed no efficacy. In the context of these results, it is somewhat surprising that valproic acid, an anticonvulsant with relatively weaker HDAC inhibitory activity, showed modestly improved outcome in newly diagnosed GBM when used in combination with radiotherapy (Tab. 1) [126]. A possible explanation for the failure to translate the pre-clinical results is that only specific patient subgroups selected based on molecular biomarkers such as HDAC expression and/or histone acetylation patterns have the potential to benefit from HDAC inhibition. To date, clinical trial designs have not selected for such populations.

In contrast to adult HGG, chromatin-remodeling defects appear to be central to the pathogenesis of pediatric HGG. Of particular significance are mutations in histone H3.3, with 78% of DIPG carrying the amino acid substitution lysine 27 to methionine (K27M) and up to 31% of non-brainstem pediatric HGGs harboring glycine 34 to valine or arginine (G34V/R) or K27M mutations [127,128]. These mutations are thought to sequester polycomb repressive complex 2 (PRC2), which in turn represses gene expression through histone methylation resulting in broad epigenetic dysregulation [8,129–131]. It is thus not surprising that epigenetic modifier panobinostat demonstrated significant anti-tumor efficacy in a preclinical DIPG animal model, further substantiating the underlying mechanism, and has rapidly emerged as a promising therapeutic strategy for this devastating disease [132]. However, panobinostat resistant DIPG cells exist, particularly following chronic drug exposure highlighting the need to optimize regimens by utilizing combination approaches. As H3<sup>K27M</sup> DIPG cells are transcriptionally more active with reduced di- and trimethylation, one potential strategy has focused on inhibiting the Jumonji-domain demethylase JMJD3, a key enzyme responsible for the demethylation [130]. The use of the GSKJ4, an inhibitor of JMJD3 demethylase, resulted in complete growth inhibition of H3<sup>K27M</sup>-expressing DIPG cells and brainstem glioma xenografts along with increased K27 methylation and a subsequent decrease in gene expression [133], an effect that could be further enhanced by panobinostat [132,134]. With these results as a foundation, panobinostat that is FDA approved for other indications could rapidly be translated into clinic use and first clinical trials are currently being planned, as single therapy or in combination to increase clinical benefit and reduce the development of resistance.

## 4. Immunotherapy in malignant glioma

### 4.1 Vaccines in Malignant Glioma

Several vaccine studies against HGG have been completed or are underway (Tab. 1). Particularly exciting has been the development of vaccines based on GBM-specific antigens, such as EGFRvIII or IDH<sup>R132H</sup>. Rindopepimut (PEPvIII vaccine), the most extensively studied vaccine for CNS neoplasms, is targeted against the EGFRvIII peptide (H-Leu-Glu-Glu-Lys-Lys-Gln-Asn-Tyr-Val-Val-Thr-Asp-His-Cys-OH) conjugated to keyhole limpet hemocyanin (KLH). ACTIVATE (A Complementary Trial of an Immunotherapy Vaccine against Tumor Specific EGFRvIII) trial was the first Phase II study to evaluate the efficacy of rindopepimut in newly diagnosed EGFRvIII-positive GBM patients, who had undergone gross total resection and standard chemoradiation therapy [135]. 19 patients were enrolled and compared favorably with a control group that received the standard therapy with temozolomide and radiation. Not surprisingly, patients who developed immune sensitization to EGFRvIII had a longer median overall survival compared to patients that lacked this response (47.7 months vs. 22.8 months) [135]. Notably, in cases of recurrence, most tumors were found to have lost EGFRvIII expressing cells. A follow-up Phase II trial (ACT II) investigated rindopepimut/GM-CSF concurrently with standard or dose-intensive adjuvant TMZ and enrolled 22 patients with newly diagnosed EGFRvIII-positive GBM [136]. All patients were found to have an immune response to EGFRvIII, with the greatest serum response seen in the dose-intensive cohort. Most importantly, overall survival was significantly improved (23.6 months) compared to historical case-matched controls [136]. The ACT III trial was the third and largest Phase II trial that evaluated the efficacy of the peptide vaccine CDX-110 in 65 patients with newly diagnosed EGFRvIII-positive GBM following gross total resection and the successful completion of combination radiation and TMZ therapy [137]. The median overall survival was 24.6 months compared to 15.2 months for matched EGFRvIII-positive controls [137,138]. A further sub-analyzes revealed that MGMT methylated patients receiving rindopepimut were found to have a significantly longer PFS (17.5 months vs. 11.2 months) and OS (32.3 months vs. 20.9 months) [137]. Although these trials rigorously confirm that rindopepimut vaccine can be safe and suggested a possible benefit in the treatment of EGFRvIII positive GBMs, these results are unlikely to be confirmed in the ongoing Phase III trial for newly diagnosed EGFRvIII positive GBM (ACT IV, NCT01480479) [139]. However, the limited clinical responses observed in Phase III and recurrence of EGFRvIII-negative tumors highlight a potential limitation of single antigen vaccines in GBMs that display considerable intratumoral cellular heterogeneity and require additive therapies to maximize the therapeutic response and control tumor growth at recurrence.

Apart from newly diagnosed GBM, rindopepimut is currently also being evaluated in a Phase II trial in patients with EGFR-positive recurrent GBM in combination with the anti-VEGF mAb bevacizumab in an attempt to optimize the EGFRvIII specific immune response through reversing VEGF mediated immunosuppression (ReACT, NCT01498328) [140]. Preliminary results presented at the 2015 ASCO Annual Meeting demonstrated that 27% of patients reached a 6-month progression free survival (vs. 11% in contemporary controls) with a median overall survival of 12 months (vs. 8.8 months in contemporary controls)

[141]. The primary data collection of the ReAct trial was completed in April 2015 and results are anxiously awaited. Also, a Phase I trial is underway utilizing the EGFRvIII vaccine in the treatment of children with DIPG, which show EGFR expression in ~50% [142].

Another promising tumor-specific neoantigen with high uniformity and penetrance is IDH1<sup>R132H</sup>. The immunogenic epitope encircling this mutation is found on major histocompatibility complexes (MHC) class II and induces mutation-specific CD4+ T-helper-1 (TH1) responses [143]. Notably, peptide vaccination of mice that are transgenic for human MHC class I and II with IDH1<sup>R132H</sup> resulted in an effective MHC class II-restricted mutation-specific antitumor immune response [143]. Based on these results two Phase I trials were initiated for patients with IDH1<sup>R132H</sup>-positive grade III and IV gliomas (NCT02454634) and for IDH1<sup>R132H</sup>-positive grade II gliomas (RESIST, NCT02193347). Other notable contemporary clinical trials target the chaperone heat shock protein (HSP) 96-kD, which is highly expressed in GBM and when carrying a tumor antigenic peptide can be delivered to dendritic cells resulting in presentation of tumor peptides [144]. By purifying HSP-96 protein complexes from each patient's tumor, a personalized polyvalent vaccine, known as HSP protein complex-96 vaccine (HSPPC-96; Prophage), was developed that has been clinically tested in Phase I and II studies in recurrent and newly diagnosed HGG suggesting good tolerability and safety [145]. Although the vaccine resulted in a measurable systemic immune response to the patient's specific tumor antigens, the therapeutic benefit is unclear and requires further clinical studies [145]. In addition, HSPPC-96 vaccine + bevacizumab is currently undergoing Phase II testing in patients with recurrent GBM (NCT01814813).

ICT-107 is a patient derived DC vaccine, which targets six GBM associated antigens, AIM-2, TRP-2, HER2/neu, IL-13Ra2, gp100, MAGE1. The vaccine has been evaluated in newly diagnosed HLA-A1<sup>+</sup> and/or HLA-A2<sup>+</sup> GBM patients following resection and concomitant chemo-radiotherapy. Preliminary results from the Phase I study were highly exciting, demonstrating a median overall survival of 38.4 months and 5-year overall survival rate was 50% [146]. A follow-up randomized, double blind, placebo-controlled Phase II trial suggested potential therapeutic benefits, particularly in HLA-A2<sup>+</sup>/MGMT-methylated subgroup with a median progression free survival of 24 months (vs. 8.5 months in the historic controls), although the efficacy was less than what was observed in Phase I [147]. In contrast, the prognostically poorer MGMT-unmethylated subgroup did not show any therapeutic response when compared to historic controls [147]. A multi-center randomized, double blind Phase III trial has been launched to study about 400 HLA-A2<sup>+</sup> patients with newly diagnosed GBMs (NCT02546102).

In the future, vaccine therapies may be combined with other means of immune augmentation to optimize their efficacy and escape the tumor-induced immunosuppression [148–150]. Potential strategies include the use of cytokines, that have been studied in combination with anti-tumor vaccines, and site preconditioning with the tetanus/diphtheria (Td) toxoid, which significantly improved the tumor-antigen specific therapeutic response [151].

## 4.2 Immune checkpoint inhibitors

Historically, the CNS was classified as an immunologically privileged site. Recent work demonstrated the presence of a lymphatic system within the CNS and new insights in the key mechanisms of tumor escape from the immune system [152,153]. Those include the tumor's ability to produce specific inhibitors of antigen-presenting cell (APC) maturation and T-cell mediated cytotoxicity, as well as other immunosuppressive factors such PD-L1, prostaglandin E2, TGF $\beta$ , indoleamine 2,3-dioxygenase (IDO), IL-10 and STAT3 and the recruitment of immunosuppressive host cells such as regulatory T cells, or myeloid-derived suppressor cells to the tumor microenvironment [153].

Immune checkpoint inhibitors, aimed at overcoming these obstacles, are humanized antibodies, which block various cell surface receptors expressed on the tumor or the host's T-cells that negatively regulate the anti-tumor immune response, e.g. cytotoxic T-lymphocyte-associated antigen 4 (CTLA-4), the programmed death-1 (PD-1) receptor and the PD-L1 ligand (Fig. 2) [153]. Clinical trials in other cancers have already convincingly demonstrated that inhibition of CTLA4, PD1, and PDL1 can activate anti-tumor immune responses, with significant and durable therapeutic benefits. In human GBMs, PD-1 ligands are also expressed on the tumor surface, however, whether this can predict a clinical response to PD-1 blockade is questionable based on the data from other cancers [154,155]. Despite lacking evidence for clinical, preclinical studies in murine syngeneic glioma models convincingly demonstrated that blocking specific co-inhibitory receptors, such as PD1 and CTLA4, alone or in combination with chemotherapy or radiation, inhibited T-cell activation and resulted in dramatic tumor regressions and long-term survival [156–158].

Five immune checkpoint inhibitors are currently undergoing clinical evaluations in HGGs, both as monotherapy or in combination with other agents (eg. bevacizumab, TMZ, radiation and vaccines): nivolumab, pidilizumab, and pembrolizumab, which inhibit the interaction of the PD1 receptor with its PDL1 ligand; MEDI4736 and MPDL3280A, which neutralize the PDL1-ligand; and ipilimumab, that targets CTLA4. While preliminary data for nivolumab and nivolumab plus ipilimumab in the treatment of recurrent GBMs are cautiously encouraging with an overall survival of 60% at 9 months, two randomized open-label Phase III studies investigating the efficacy and safety of nivolumab compared to bevacizumab and nivolumab with or without ipilimumab are underway for newly diagnosed and recurrent GBM (CheckMate 143; NCT02017717), as well as nivolumab + radiation in newly diagnosed GBM (CheckMate 498; NCT02617589). In addition, the safety of ipilimumab, nivolumab, and the combination of ipilimumab and nivolumab will be evaluated in newly diagnosed GBM patients undergoing temozolomide therapy (Phase I, NCT02311920). Furthermore, pidilizumab is being tested in Phase I/II to assess safety and efficacy in children with DIPG (NCT01952769). For pembrolizumab, four clinical trials have recently opened or are ongoing: a Phase I study investigating pembrolizumab in combination with bevacizumab and hypofractionated stereotactic irradiation in recurrent HGGs (NCT02313272); a Phase I/II study investigating the safety and efficacy of pembrolizumab in combination with MRI-guided laser ablation in recurrent HGG (NCT02311582); a Phase II study of pembrolizumab with and without bevacizumab in recurrent GBM



(NCT02337491); and a Phase I/II study evaluating the safety and efficacy of pembrolizumab in combination with standard chemoradiation in newly diagnosed GBM (NCT02530502).

Despite the large number of ongoing studies, the evaluation of the immune-checkpoint inhibitors must be systematic and proceed with caution as evidenced by the recent suspension of the Phase I/II trial evaluating pembrolizumab in children with recurrent HGGs and DIPG due to severe adverse reactions (NCT02359565). This highlights one of the major challenges associated with immunotherapies: uncontrolled immune activation. Hence, there is a crucial need to identify tools to accurately assess and control the intensity of immune response to avoid potentially life-threatening conditions, such as toxic autoimmunity directed at brain antigens (allergic encephalomyelitis), elevated intracranial pressure and cerebral edema. Similar immune-mediated toxicities have also been observed with other immunotherapies, such as T-cell based immunotherapies, particularly chimeric antigen receptor T cell (CAR T-cell) therapy targeting EGFRvIII (Tab.1) [159]. Apart from toxicities, we will face unique challenges in the assessment of the radiographic response because the inflammatory responses might mimic radiological features of tumor progression with increased enhancement and edema. This has led to the development of the immunotherapy Response Assessment for Neuro-Oncology (iRANO) criteria, in which patients with imaging findings suggestive of progressive disease within 6 months of starting immunotherapy including the development of new lesions, should undergo follow-up imaging in a 3 months' time before defining the patient as non-responsive to treatment or progressive disease [160]. In addition, insights yielded from other clinical trials evaluating the immune checkpoint inhibitors in brain metastasis will hopefully provide further guidance towards addressing these unresolved questions. Finally, biomarkers that identify patients who will likely benefit from specific agents are needed. Recent data suggests that tumors with mismatch repair deficiency, which is only present in a small fraction of brain cancers leading to a higher mutational load, may be more susceptible to immune checkpoint blockade and thus, providing a potential screening tool [161]. Its role in CNS neoplasms, however, has yet to be defined although early data are encouraging [162].

#### 4.4 Oncolytic viruses

Oncolytic viruses represent a promising class of therapeutic agents that promote antineoplastic responses through a combination of two mechanisms: cytotoxicity via selective replication within neoplastic cells, resulting in direct tumor cell killing and induction of a systemic antitumor immunity. The concept of oncolytic virotherapy has a long history of nearly 100 years and is based on the observation that viruses have a preferential, although nonexclusive, tropism for tumors [163]. Virotherapy became a popular tool in cancer therapy in the 1990s, when advances in recombinant technology met an improved molecular understanding of virology and cancer genetics [163,164]. Many viruses have been investigated as agents for oncolytic immunotherapy in gliomas, and considerable work has been done to optimize viral vectors by attenuating pathogenicity and enhancing immunogenicity. As of 2015, approximately 15 viruses of six different species have advanced to clinical trials: adenoviruses, herpes simplex virus-1 (HSV-1), coxsackie virus, poliovirus, measles virus, poxvirus, Newcastle disease virus (NDV) and reovirus (Tab. 1) [163]. The majority of these oncolytic viruses have a natural tropism for cell surface proteins

that are expressed by glioma cells. For example, poliovirus uses CD155, a widely expressed receptor on GBM cells, for cell entry. Coxsackievirus can enter cells *via* intercellular adhesion molecule 1 and decay accelerating factor, and HSV-1 utilizes the herpesvirus entry mediator (HVEM) and selected nectins for cell entry. Adenovirus DNX-2401 was developed by genetically modifying adenovirus type 5 to bind integrins that are highly expressed on cancer cells. In addition, some of these viruses have also been engineered to selectively replicate exclusively in GBM-specific conditions, such as hypoxia or oncogenic mutation (e.g. p53 mutation, Retinoblastoma (Rb) mutation) to increase the tumor specificity and spare normal cells. For instance, adenovirus ONYX-015 (dl1520) is deleted in the E1B gene, which encodes a 55 kDa protein that inactivates p53 protein and thus, exclusively replicates in and lyses p53-deficient tumor cells [165,166]. Likewise, tumor-selective adenovirus Delta24 carries a deletion in the E1A region responsible for binding Rb protein and can only replicate in Rb deficient glioma cells [167].

There are more than 20 clinical trials of viral oncolytics completed or active in CNS tumors. The majority of them have used direct intratumoral injections to bypass the architectural barriers of the tumor and surrounding tissue as well as increase infectivity. Two oncolytic HSV-1 strains (G207 and 1716) have completed Phase I and II testing in adult recurrent HGGs, alone and in combination with radiation. For G207, no serious adverse events were observed and at best, 38% of patients had a clinical response, and one patient had long-term survival (>5.5 years) [168]. However, a follow-up Phase Ib study, was not able to reproduce these results [169]. A third study in combination with a single dose of 5 Gy radiation (NCT00157703) resulted in a probable clinical response and further studies are necessary to define its role in the therapy of HGGs [170]. G207 is also currently undergoing Phase I testing in children with progressive and recurrent HGGs (NCT02457845). Three Phase I clinical trials with HSV-1716 have been conducted in recurrent HGGs, demonstrating safety and paved the way for a Phase I trial in children with recurrent HGGs (NCT02031965). Furthermore G47 delta, derived from G207 with an additional deletion of nonessential  $\alpha 47$  gene to enhance replication efficacy, and M032, an HSV-1 strain equipped with a human IL-12 transgene, have entered early clinical testing for recurrent or progressive HGGs (NCT02062827).

DNX-2401 (DNATrix) is a conditionally replicative oncolytic adenovirus designed to target tumor cells that are defective in the Rb pathway. It is currently being evaluated in a Phase I trial for the treatment of recurrent HGGs (NCT00805376). Preliminary results presented at the *Society of Neuro-Oncology* (SNO) meeting in 2014 were promising showing complete and durable responses in up to 12% of the patients [171]. Interestingly, responders also demonstrated interleukin responses, particularly IL-12 elevations, re-emphasizing the concept that intratumoral administration of oncolytic viruses does not only act to directly kill tumor cells though viral replication but also induces anticancer immunity.

Another promising oncolytic is the TOCA 511 virus that belongs to the family of non-lytic retroviruses, and carries the cytosine deaminase (CD) gene to enhance the direct cancer cell killing via local conversion of systemically administered prodrug 5-fluorocytosine (5-FC) to the active 5-fluorouracil (5-FU). It is currently undergoing Phase I clinical trial in adults with newly diagnosed HGGs co-administered with standard chemoradiation

(NCT02598011), and recurrent HGGs (NCT01156584), which is soon to be completed. A Phase II/III trial just initiated patient enrollment for recurrent HGGs in November 2015 (NCT02414165).

Poliovirus, termed PVS-RIPO in its therapeutic engineered form, demonstrated potent lytic effects and dramatic tumor regressions in preclinical HTB-15 glioma xenografts [172]. PVS-RIPO has advanced to a Phase I clinical trial for patients with recurrent HGGs (NCT01491893). Although the results are not published yet, early data suggest moderate therapeutic efficacy in selected patient subsets (40% survival reported at 20 months) [173] and Phase II/III trials as well as an expansion to pediatric HGGs are currently planned. Other oncolytic viruses in early stages of clinical development include parvovirus H1 (NCT01301430), and measles virus (NCT00390299).

Although the replicating nature of oncolytic viruses poses unique and potentially life-threatening toxicities, clinical trials have shown that the majority of oncolytic therapies results in few adverse events, even in patients with brain tumors [164]. However, clinical responses in HGG clinical trials have fallen short of the promise initially generated from preclinical animal testing, in part because dosing and application regimens were chosen conservatively and the inability of current animal model systems to sufficiently replicate human viral pathogenesis. Human HGG xenograft models lack immune competency and immune-competent mouse models are frequently misleading because the viruses behave differently in rodents and humans. The BBB likely limits oncolytic virus entry into the CNS, and except for Parvovirus that naturally crosses the BBB, there have been few studies to determine the viral penetrance and distribution in the CNS. CNS bioavailability can be overcome by direct infusion of virus into the brain tumor but the viral distribution can still be limited due to patient-specific differences in tumor anatomy (e.g. cysts, necrosis). Furthermore, viral efficacy may be hampered by genetic modifications intended to limit toxicity that inadvertently affect the viral replication and spread [164]. In summary, it remains to be determined if oncolytic virotherapy can be designed in a way to manage toxicities at clinically effective doses. Perhaps more importantly, anti-tumor immune responses generated during oncolytic virus infections could be used to stimulate the systemic anti-tumor immunity and provide a strong rationale for the clinical exploration of combinations of immunoregulatory antibodies with oncolytic viruses [174,175]. Clinical trials using oncolytic viruses, such as HSV-1, in combination with immune checkpoint inhibitors (ipilimumab) are currently under way for advanced solid cancers and will provide a clinical proof of concept in humans (NCT02272855).

## 5. Conclusion

Unlike other solid tumors, the complexity and diversity of malignant gliomas poses unique challenges to the therapeutic approaches including intratumoral heterogeneity, divergent signaling mechanisms, insufficient tumor targeting strategies and limited drug delivery through the BBB. These obstacles have made this cancer one of the most difficult ones to treat despite an unprecedentedly high number of new investigational agents that have evolved over the last years. Though the therapeutic success has been limited so far, multiple promising strategies are still untested such as the concurrent use of multiple modalities that

target disparate tumor features, the use of local cytotoxic therapies to overcome limitations from the BBB and minimize the systemic immune-suppression, particularly when combined with immunotherapeutics, and the use of specific patient subgroups selected based on molecular biomarkers or mutations such as HDAC or RTK expression to evaluate realistically potential benefits of candidate agents. We need to be aware that novel cancer therapeutics may introduce new hurdles, as observed with immunotherapeutics and their immune-related response patterns, which pose an increasing clinical challenge for practitioners and patients. Finally, research has yet to discover a single definitive target or event that deserves the major research focus of the neuro-oncological community to successfully battle this devastating disease.

## 6. Expert Opinion

This is a time of progress and guarded optimism in Neuro-Oncology. Remarkable progress in survival has been achieved for subsets of patients with GBM and anaplastic oligodendroglioma by combining chemotherapy with radiation therapy highlighting the fact that significant advances can be made by using traditional therapeutics in creative ways. The guarded optimism while fully warranted is counterbalanced by the complexities and heterogeneity of brain cancer biology and the unique obstacles associated with diagnosing, treating, and monitoring tumors within the central nervous system. We are likely reaching the limits of what can be achieved by combining existing cytotoxic regimens and it remains frustrating that chemotherapies proven to be effective against malignant glioma remain woefully limited to the class of alkylating agents. There remains no meaningful therapy for many of our patients including but not limited to those with recurrent GBM and aggressive pediatric HGGs. Brain cancer patients have not yet benefited from the promise of personalized targeted therapy despite the fact that GBM, the most common brain malignancy, is one the most extensively characterized cancers at the genetic and epigenetic levels.

Optimism regarding the prospect for totally new therapeutic approaches is warranted and driven by research advances in multiple directions. Progress in brain cancer genomics and epigenomics has identified multiple oncogenic drivers amenable to therapeutic inhibition many of which are just entering clinical trial. However, single cell analyses of GBM are revealing an unprecedented degree of intratumoral cellular heterogeneity, findings that most certainly reflect GBM's resilience and highlight the need to more precisely define the evolutionary biology of these malignancies at the cellular and molecular levels, especially their evolution during treatment and the emergence of therapeutic resistance. The expectation is that this information will provide a roadmap to guide emerging sequential and combinatorial treatment regimens. The original focus of targeted therapy on oncogenic kinase inhibitors (e.g. EGFR, PDGFR inhibitors) has now expanded to include strategies for targeting oncometabolite production, chromatin dysregulation, and their effects on gene expression regulation. The brain is no longer misconceived as an immune-privileged site opening the prospect of harnessing the immune system to overcome brain cancer's heterogeneity and ability to evade cytotoxic and targeted therapies. This, in conjunction with advances in our understanding of the immune-suppressive strategies employed by cancer, has stimulated the development of innumerable exciting immune-based clinical trials using

vaccines and immune checkpoint inhibition. Vigilance is needed to not let current immune-suppressive standards of care (i.e. chemo-radiation) block the efficacy of immune-based therapy.

Significant challenges to the rapid and successful development of these promising approaches remain. The path from the preclinical laboratory to FDA approval is time consuming and increasingly expensive taking decades and hundreds of millions of dollars for new agents. New and more efficient clinical trial designs that minimize patient numbers and maximize the rapid detection of therapeutic signal are critical. Employing the personalized medicine approaches of tumor subtyping and patient selection for therapeutic target expression is vital to achieving these goals. Limited CNS bioavailability of systemically administered drugs remains a serious obstacle to brain tumor therapy and should no longer be relegated to the background. Toward overcoming this obstacle direct intratumoral therapies delivered by increasingly advanced forms of convection enhanced delivery, implanted biomaterials, or replication competent biological agents are becoming more sophisticated and practical. Effectively addressing current challenges will require increased financial and programmatic commitments from national funding agencies, the pharmaceutical industry and philanthropy. With this support, the growing critical mass of brain cancer scientists and clinicians, attributable in great part to the founding of The Society for Neuro-Oncology two decades ago, in conjunction with the critically important clinical trials consortia are prepared to overcome these challenges and accelerate the discovery and implementation of more effective therapies for our highly motivated brain cancer patients for whom time is of the essence.

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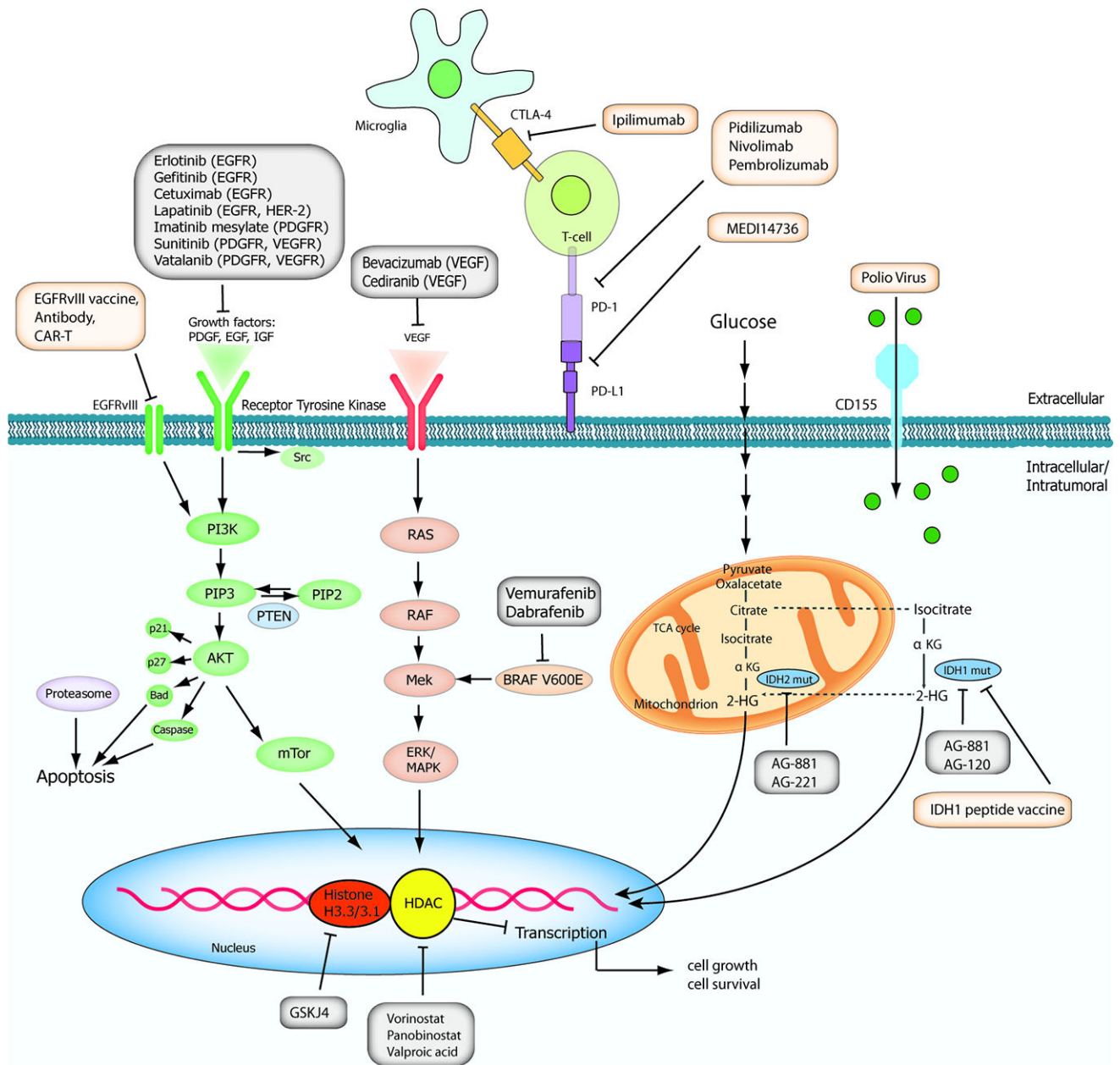
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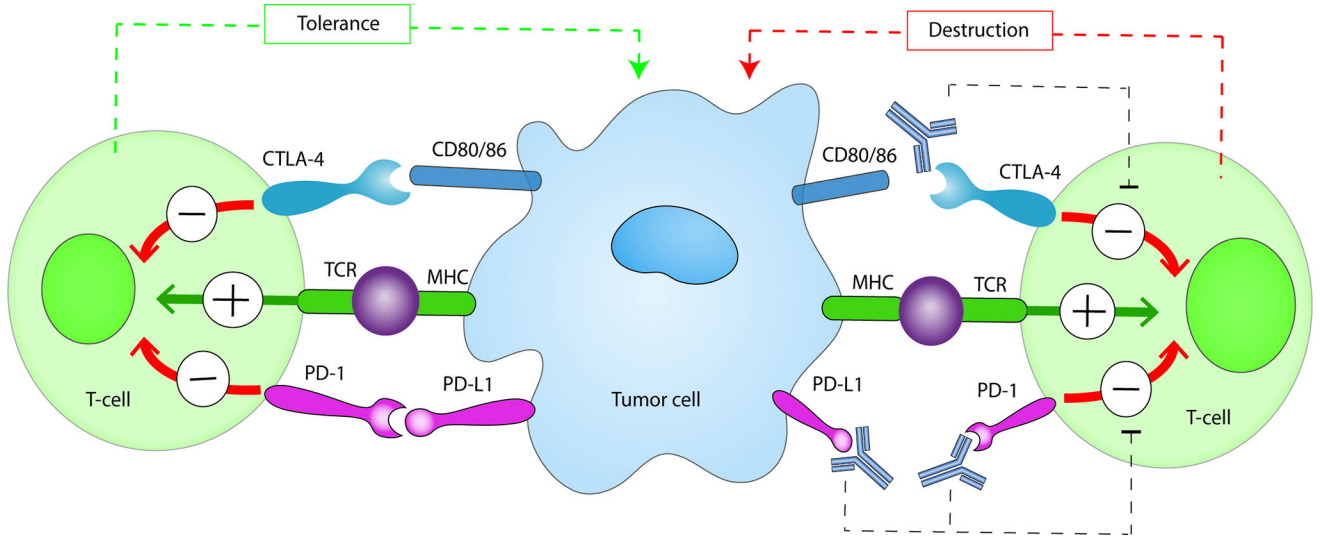
### Highlight Box

- Pediatric and adult high-grade gliomas (HGG) carry a uniformly fatal diagnosis due to high recurrence rates and the absence of effective treatments.
- Progress in brain cancer genomics and epigenomics have identified multiple oncogenic drivers targetable by therapeutic inhibition, many of which are currently undergoing evaluations in clinical trial but have yet to show improvements in survival.
- Promising new developments focusing on immunotherapies such as immune-checkpoint inhibitors or oncolytic viruses are underway to overcome the genetic drift and could lead to durable anti-tumor immune responses against a theoretically unlimited number of tumor-associated antigens.
- Future successful therapies will concurrently use multiple modalities that target disparate tumor characteristics personalized to the particular tumor of a given patient.



**Figure 1. Schematic overview of current targeted therapies in HGGs**

Aberrant oncogenic RTK pathways including the PI3K-AKT (green) and RAS (pink) oncogenic pathways are targeted with a variety of small molecule inhibitors (grey boxes) and monoclonal antibodies. Oncometabolites produced by IDH1/2-mutated glioma cells can be intracellularly targeted by small molecule inhibitors and a IDH1 mutant specific vaccine, which is currently being tested HGG patients. Epigenetic modifiers targeting on HDAC and histone 3.3/3.1 are shown as well. Notably, various immunotherapeutic strategies that are currently undergoing evaluations in clinical trials are pictured including tumor vaccines, immune checkpoint inhibitors, CAR-T cell therapies, and oncolytic viruses (yellow box). Abbreviations: mut: mutant.



**Figure 2. Immune checkpoint interactions on T cells and cancer**

Antigens released from tumor cells are taken up by antigen-presenting cells. The T-cell receptor (TCR) interacts with a presented antigen to activate the immune system. PD-1, which is expressed on T-cells, has an inhibitory role (red arrow) through its interaction with PD-L1 on the tumor cells resulting in tolerance and inhibition of tumor destruction. Similarly, CTLA-4, which is also expressed on T-cells, mediates inhibitory signals after binding with its ligands CD80 or CD86 dampening the immune response and preventing activation. Blocking both interactions, with monoclonal antibodies to PD-1, PD-L1 or CTLA-4 (for example, nivolumab, pembrolizumab, ipilimumab) precludes inhibition allowing for T-cell activation (green arrow) with subsequent tumor cell lysis.

**Table 1**

Overview of selected completed and ongoing clinical trials for malignant gliomas in adults and children.

Drug	Target	Treatment Arms	Trial Phase	Tumor Type	Clinical Trial Identification Number	Status/Comments/Reference
<i>RTK, RAS, PI3-kinase pathway targeting drugs</i>						
Erlotinib	EGFR	Erlotinib	II	rec HGG	NCT00387894	Terminated due to low accrual [31]
		Erlotinib+TMZ+RDX	II	initial GBM	NCT00187486	Completed [33]
		Erlotinib+TMZ+Vorinostat	I/II	rec GBM	NCT01110876	Terminated due to toxicities
		<b>Erlotinib + Bev</b>	II	<b>initial GBM</b>	<b>NCT00720356</b>	<b>Active</b>
Gefitinib	EGFR	Erlotinib + temsirolimus	I/II	rec HGG	NCT00112736	Completed [41]
		Gefitinib	II	initial GBM	NCT00014170	Completed [34]
		Gefitinib+RDX	I/II	rec GBM	NCT00025675	Completed [30]
		Gefitinib+TMZ	I	initial GBM	NCT00052208	Completed [36]
Lapatinib	EGFR	Lapatinib	I/II	rec HGG	NCT00027625	Completed [35]
		Lapatinib	I/II	rec GBM	NCT00099060	Completed [37]
		Lapatinib+TMZ	I/II	ped rec HGG	NCT00095940	Completed [38]
		<b>Lapatinib+TMZ+RDX</b>	II	rec HGG	NCT00107003	Completed
Cetuximab	EGFR	Lapatinib+pazopanib	II	rec HGG	<b>NCT01591577</b>	<b>Active</b>
		Cetuximab+TMZ+RDX	II	rec HGG	NCT00350727	Completed [42]
		<b>IA Cetuximab+Bev</b>	I/II	initial GBM (MGMT -) <b>rec HGG, DIPG</b>	NCT01044225	Terminated [39]
Dacomitinib	EGFR	<b>Dacomitinib</b>	II	<b>rec GBM</b>	<b>NCT01520870</b>	<b>Active, not recruiting</b>
Afatinib	EGFR	Afatinib+/-TMZ	II	<b>rec GBM</b>	<b>NCT01112527</b>	<b>Active, not recruiting</b>
Sym004	EGFR	<b>Sym004</b>	II	rec HGG	NCT00727506	Completed [45]
AMG 595	EGFRvIII	<b>AMG 595</b>	I	<b>rec GBM</b>	<b>NCT02540161</b>	<b>Active</b>
ABT-414	EGFRvIII	<b>ABT-414+TMZ+RDX</b>	Ib/III	rec HGG	<b>NCT01475006</b>	<b>Active, not recruiting</b>
		<b>ABT-414+TMZ</b>	II	<b>initial GBM</b>	<b>NCT02573324 (Intelligence I)</b>	<b>Active</b>
Imatinib	PDGFR c-KIT BCR-ABL	Imatinib	I/II	rec HGG	NCT00010049	Completed [49]
		Imatinib+TMZ	I	HGG	NCT00354068	Completed [176]
		Imatinib+HU	I/II	rec HGG	NCT00290771	Terminated [48]
Sunitinib	PDGFR VEGFR	Sunitinib	II	rec GBM	NCT00606008	Completed [73]
		Sunitinib+RDX	II	ped rec HGG	NCT01462695	Completed
		Sunitinib+cilengitide	I	initial HGG	NCT01100177	Completed [74]
Sorafenib	PDGFR VEGFR	Sunitinib	I	initial GBM	NCT01122888	Terminated
		Sorafenib+TMZ	II	rec HGG	NCT00597493	Completed [68]
		Sorafenib+TMZ+RDX	I/II	initial GBM	NCT00734526	Withdrawn [75]
		Sorafenib+Bev	II	rec GBM	NCT00621686	Completed [72]
Pazopanib,	PDGFR VEGFR	<b>Sorafenib+everolimus</b>	I/II	<b>rec GBM</b>	<b>NCT01434602</b>	<b>Active, not recruiting</b>
		Pazopanib	II	rec GBM	NCT00459381	Completed [51]
		Pazopanib+Lapatinib	II	rec GBM	NCT00350727	Completed [42]
Vatalanib	PDGFR VEGFR	<b>Pazopanib+TMZ</b>	I/II	<b>initial GBM</b>	<b>NCT02331498</b>	<b>Active</b>
		<b>Pazopanib+topotecan</b>	II	<b>rec GBM</b>	<b>NCT01931098</b>	<b>Active</b>
Lenvatinib	PDGFR VEGFR	Vatalanib+TMZ+RDX	I	initial GMB	NCT00385853	Completed [177]
		Lenvatinib	II	rec HGG	NCT01137604	Terminated [78]
Dasatinib	PDGFR c-KIT BCR-ABL Src	Lenvatinib+E7050	I/II	rec HGG	NCT01433991	Completed
		<b>Dasatinib</b>	II	<b>rec GBM</b>	<b>NCT00423735</b>	<b>Active</b> [52]
		<b>Dasatinib+TMZ+RDX</b>	II	<b>initial GBM</b>	NCT00895960	Terminated
		Dasatinib+CCNU	I/II	rec GBM	<b>NCT00869401</b>	<b>Active</b> [57]
Crenolanib	PDGFR Flt3	<b>Dasatinib+Bev</b>	I/II	<b>rec GBM</b>	(Alliance N0877) NCT00948389 <b>NCT00892177</b>	Terminated [53]
			I/II		(Alliance N0872)	<b>Active, not recruiting</b>
		<b>Crenolanib</b>	II	rec HGG+LGG <b>ped rec</b>	NCT01229644 <b>NCT01393912</b>	Terminated <b>Active, not recruiting</b>



			II	HGG, DIPG rec GBM	NCT02626364	Active
AMG 102	c-MET	AMG AMG 102+Bev	II II	rec HGG rec HGG	NCT00427440 NCT01113398	Completed [65] Completed
SGX523	c-MET	SGX523	I I	rec HGG rec HGG	NCT00606879 NCT00607399	Terminated Terminated
Onartuzumab	c-MET	Onartuzumab+Bev	II	rec GBM	NCT01632228	Completed
Cabozantinib	VEGFR c-MET c-KIT	Cabozantinib Cabozantinib+TMZ+RDX	II I	rec GBM Initial GBM	NCT00704288 NCT00960492	Completed [64] Completed [178]
Bevacizumab	VEGF	Bev+/-RDX Subc. Bev Bev+BV-111 (GLOBE) Bev+CCN	II II III II	rec GBM rec GBM rec GBM rec GBM	NCT01730950 NCT02157103 NCT02511405 NCT01067469	Active Active Active Active, not recruiting
Cediranib	VEGFR	Cediranib+TMZ+RDX	II	initial GBM	NCT01062425	Active, not recruiting
Vandetanib	VEGFR	Vandetanib+RDX Vandetanib+dasatinib Vandetanib+TMZ+RDX Vandetanib+carboplatin	I I I/II II	rec HGG DIPG initial GBM rec HGG	NCT00822887 NCT00996723 NCT00441142 NCT00995007	Completed [179] Completed [180] Completed [181] Active
Axitinib	VEGFR	Axitinib	II	rec HGG	NCT01562197	Completed [79]
Temsirolimus (TS)	mTOR	TS+Perifosine TS+TMZ+RDX	I/II II I	rec HGG rec HGG initial HGG	NCT01051557 NCT02238496 NCT00316849	Active, not recruiting Active Completed [87]
Everolimus	mTOR	Everolimus Everolimus+TMZ	II II II	rec LGG ped rec LGG prog LGG	NCT00823459 NCT01734512 NCT02023905	Active Active Active
Sapanisertib	mTOR	Sapanisertib Sapanisertib+Bev	0 I	rec HGG rec HGG	NCT02133183 NCT02142803	Active Active
Buparlisib (B)	PI3K	B+carboplatin/CCNU B+Bev	I/II I/II	rec GBM rec GBM	NCT01934361 NCT01349660	Active Active
PX-866	PI3K	PX-866	II	rec GBM	NCT01259869	Completed
XL-147	PI3K	XL-147	I	rec GBM	NCT01240460	Completed
Perifosine	AKT	Perifosine Perifosine+TS	II I/II II	rec HGG rec HGG rec HGG	NCT00590954 NCT01051557 NCT02238496	Active, not recruiting Active, not recruiting Active
Vemurafenib	BRAF <sup>V600E</sup>	Vemurafenib	0	rec BRAF V600E-mutant gliomas	NCT01748149	Active
Dabrafenib	BRAF <sup>V600E</sup>	Dabrafenib Dabrafenib+Trametinib	I/IIa II	rec BRAF V600E-mutant gliomas	NCT01677741 NCT02034110	Active Active
Trametinib	BRAF	Trametinib+Dabrafenib	I II	ped glioma HGG, LGG	NCT02124772 NCT02034110	Active Active
MEK 162	BRAF	MEK 162	I/II	ped HGG, LGG	NCT02285439	Active
Selumetinib	BRAF	Selumetinib	I/II	ped rec LGG	NCT01089101 NCT01386450	Active Active
<i>Onco-metabolism targeting drugs</i>						
AG-881	IDH1/2	AG-881	I	rec IDH1/2 mut glioma	NCT02481154	Active
AG-120	IDH1	AG-120	I	rec IDH1 mut glioma	NCT02073994	Active
AG-221	IDH2	AG-221	I/II	rec IDH2 mut glioma	NCT02273739	Active
DCA	Anti-metabolite	DCA	II	rec HGG Initial HGG	NCT00540176	Completed
<i>Epigenetic modulators</i>						
Vorinostat	HDACi	Vorinostat+TMZ Vorinostat+FSRT Vorinostat+RDX Vorinostat+TS+RDX	I I I I/II I	rec HGG rec LGG/HGG rec glioma DIPG DIPG	NCT00268385 NCT01076530 NCT01378481 NCT01189266 NCT02420613	Completed [182] Terminated Active, not recruiting Active Active

Panobinostat	HDACi	Panobinostat <b>Panobinostat+RDX</b>	II <b>I</b>	rec HGG <b>rec HGG</b>	NCT00848523 <b>NCT01324635</b>	Terminated due to low accrual <b>Active, not recruiting</b>
Valproic Acid	HDACi	Vaproic acid+TMZ+RDX	II	Rec HGG	NCT00302159	Active, not recruiting [126]
<i>Glioma Vaccines</i>						
EGFRvIII peptide vaccine	EGFRvIII	Vaccine+ GM-CSF	II	initial GBM	NCT00643097 (ACTIVATe)	Completed [135]
			II	initial GBM	(ACT II)	Completed [136]
		<b>Vaccine+GM-CSF+Bev</b>	II	initial GBM	NCT00458601 (ACT III)	Completed [137]
			III	<b>initial GBM</b>	<b>NCT01480479 (ACT IV)</b>	<b>Active [139]</b>
			II	<b>rec GBM</b>	<b>NCT01498328 (ReACT)</b>	<b>Active [141]</b>
Vaccine +GM-CSF	I	DIPG	NCT01058850	Terminated		
IDH1 Peptide Vaccine	IDH1 mut	<b>IDH1 peptide vaccine</b>	<b>I</b> <b>I</b>	<b>IDH1 mut HGG</b> <b>IHD1mut LGG</b>	<b>NCT02454634</b> <b>NCT02193347</b>	<b>Active</b> <b>Active</b>
ICT-107	6 tumor-associated antigens	ICT-107	II <b>III</b>	initial GBM <b>initial GBM</b>	NCT01280552 <b>NCT02546102</b>	Completed [147] <b>Active</b>
Glioma Vaccine IMA950	11 tumor-associated peptides	IMA950	I	rec HGG	NCT01403285	Terminated due to low accrual
HSPPC-96 Vaccine	HSP-96	Vaccine+TMZ	II	initial HGG	NCT00905060	Completed [269]
		<b>Vaccine + Bev</b>	II <b>II</b>	rec HGG <b>rec HGG</b>	NCT00293423 <b>NCT01814813</b>	Completed [145] <b>Active</b>
Tumor lysate vaccine	Tumor Antigens	Vaccine alone	I	rec HGG	NCT00068510	Completed [183]
Tumor lysate vaccine	Tumor Antigens	Vaccine+TMZ	I	rec HGG	NCT00323115	Completed [183]
Tumor lysate vaccine	Tumor Antigens	<b>Vaccine alone</b>	<b>I</b>	<b>rec HGG</b>	<b>NCT01204684</b>	<b>Active, not recruiting</b>
DCVax-L	Tumor Antigens	DCVax-L	<b>III</b>	<b>rec HGG</b>	<b>NCT00045968</b>	<b>Active, not recruiting [183]</b>
CMV vaccine (Pep-CMV)	Tumor Antigens	Vaccine+TMZ	I	rec HGG	NCT01854099	Withdrawn
<i>Immune-checkpoint inhibitors</i>						
Pidilizumab	anti-PD1	<b>Pidilizumab</b>	I/II	<b>rec GBM, DIPG</b>	<b>NCT01952769</b>	<b>Active</b>
Nivolumab (N)/ Ipilimumab (I)	anti-PD1	<b>N</b> <b>N+I</b> <b>Bev</b>	<b>III</b>	<b>rec GBM</b>	<b>NCT02017717</b>	<b>Active</b>
	anti-CTLA4	<b>N+I+TMZ</b>	<b>I</b>	<b>intial GBM</b>	<b>NCT02311920</b>	<b>Active</b>
Nivolumab (N)	anti-PD1	<b>N+RDX</b>	<b>III</b>	<b>initial GBM</b>	<b>NCT02617589</b>	<b>Active</b>
		<b>N (neoadjuvant)</b>	<b>II</b>	<b>initial GBM</b>	<b>NCT02550249</b>	<b>Active</b>
		<b>N+DC vaccine</b>	<b>I</b>	<b>rec HGG</b>	<b>NCT02529072</b>	<b>Active</b>
MEDI4736	anti-PDL1	<b>MEDI4736</b> <b>MEDI4736+RDX</b> <b>MEDI4736+Bev</b>	<b>II</b>	<b>initial and rec GBM</b>	<b>NCT02336165</b>	<b>Active</b>
Pembrolizumab (P)	anti-PD1	<b>P + Bev</b> <b>P</b>	<b>II</b> <b>-</b>	<b>rec GBM</b> <b>ped rec HGG, DIPG</b>	<b>NCT02337491</b> <b>NCT02359565</b>	<b>Active</b> <b>Suspended</b>
		<b>P+TMZ+RDX</b>	<b>I/II</b>	<b>initial GBM</b>	<b>NCT02530502</b>	<b>Active</b>
		<b>P+MRI guided laser ablation</b>	<b>I/II</b>	<b>rec GBM</b>	<b>NCT02311582</b>	<b>Active</b>
		<b>P+HFSRT+Bev</b>	<b>I</b>	<b>rec GBM</b>	<b>NCT02313272</b>	<b>Active</b>
<i>CAR T-cell therapies</i>						
anti-EGFR CAR T	EGFR	<b>anti-EGFR CAR-T</b>	<b>I</b>	<b>rec GBM</b>	<b>NCT02331693</b>	<b>Active</b>
anti-EGFRvIII CAR	EGFRvIII	<b>anti-EGFRvIII CAR-T</b>	<b>I</b>	<b>rec GBM</b>	<b>NCT01454596</b>	<b>Active</b>

Genetically modified T-cells		<b>T-cells</b>	<b>I</b>	<b>rec GBM</b>	<b>NCT02208362</b>	<b>Active</b>
HER2-specific CAR	HER2	<b>HER2-specific T cells</b>	<b>I</b>	<b>rec GBM</b>	<b>NCT02442297</b>	<b>Active</b>
<i>Oncolytic Viruses</i>						
Ad5-Delta24RGD	Rb	Virus	I/II	rec GBM	NCT01582516	Completed
DNX2401	Rb	Virus <b>Virus+TMZ</b> <b>Virus+Interferon gamma</b>	<b>I</b> <b>I</b> <b>I</b>	rec GBM <b>rec GBM</b> <b>rec GBM</b>	NCT00805376 <b>NCT01956734</b> <b>NCT02197169</b>	Completed [171] <b>Active</b> <b>Not recruiting yet</b>
HSV1G207		Virus Virus+RDX <b>Virus +RDX</b>	I/II <b>I</b> <b>I</b>	rec GBM rec GBM <b>ped rec HGG</b>	NCT00028158 NCT00157703 <b>NCT02457845</b>	Completed[168] Completed [170] <b>Not recruiting yet</b>
HSV-1716		<b>Virus</b>	<b>I</b>	<b>ped rec HGG</b>	<b>NCT02031965</b>	<b>Active</b>
M032-HSV-1		<b>Virus</b>	<b>I</b>	<b>rec HGG</b>	<b>NCT02062827</b>	<b>Active</b>
Reovirus	RAS	Virus	I/II	rec GBM	NCT00528684	Completed [184]
Poliovirus (PVS-RIPO)		<b>Virus</b>	<b>I</b>	<b>rec GBM</b>	<b>NCT01491893</b>	<b>Active [173]</b>
CEA-expressing measles virus		<b>Virus</b>	<b>I</b>	<b>rec GBM</b>	<b>NCT00390299</b>	<b>Active</b>
Toca 511		<b>Virus+FC</b> <b>Virus+FC+TMZ+RDX</b> <b>Virus+FC</b>	<b>II/III</b> <b>I</b> <b>I</b>	<b>rec HGG</b> <b>initial GBM</b> <b>rec HGG</b>	<b>NCT02414165</b> <b>NCT02598011</b> <b>NCT01156584</b>	<b>Not recruiting yet</b> <b>Active</b> <b>Active</b>
Parvovirus H1		Virus	I	rec HGG	NCT01301430	Completed
Ad5CMV-p53		Virus	I	rec HGG	NCT00004041	Completed
Ad-hCMV-TK and Ad-hCMV-Flt3L		<b>Virus</b>	<b>I</b>	<b>rec HGG</b>	<b>NCT01811992</b>	<b>Active</b>
AdV-tk		<b>Virus+valcylovir</b>	<b>II</b>	<b>initial HGG</b>	<b>NCT00589875</b>	<b>Active, but not recruiting</b>

A search was carried out on <http://www.clinicaltrials.gov>. Currently active trials are highlighted in bold.

Abbreviations: Bev: bevacizumab; CEA: carcinoembryonic antigen; DIPG: diffuse intrinsic pontine glioma; FSRT: fractionated stereotactic radiation therapy; GBM: glioblastomas; HGG: high-grade glioma; HDACi: HDAC inhibitor; HU: hydroxyurea; IA: intra-arterial; mut: mutant; Ped: pediatric; rec: recurrent; Prog: progressive; RDX: radiation; subc: subcutaneous; TMZ: temozolomide.