Risk stratification in pediatric low-grade glioma and glioneuronal tumor treated with radiation therapy: An integrated clinicopathologic and molecular analysis

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#### **Abstract**

**Background.** Management of unresectable pediatric low-grade glioma and glioneuronal tumor (LGG/LGGNT) is controversial. There are no validated prognostic features to guide use of radiation therapy (RT). Our study aimed to identify negative prognostic features in patients treated with RT using clinicopathologic and molecular data and validate these findings in an external dataset.

**Methods.** Children with non-metastatic, biopsy-proven unresectable LGG/LGGNT treated with RT at a single institution between 1997 and 2017 were identified. Recursive partitioning analysis (RPA) was used to stratify patients into low- and high-risk prognostic groups based on overall survival (OS). CNS9702 data were used for validation.

**Results.** One hundred and fifty patients met inclusion criteria. Median follow-up was 11.4 years. RPA yielded low- and high-risk groups with 10-year OS of 95.6% vs. 76.4% (95% CI: 88.7% - 98.4% vs. 59.3% - 87.1%, p=0.003), respectively. These risk groups were validated using CNS9702 dataset (n=48) (4-year OS: low-risk vs. high-risk: 100% vs. 64%, p<0.001). High-risk tumors included diffuse astrocytoma or location within thalamus/midbrain. Low-risk tumors included pilocytic astrocytoma/ganglioglioma located outside of the thalamus/midbrain. In the subgroup with known *BRAF* status (n=49), risk stratification remained prognostic independent of *BRAF* alteration (V600E or fusion). Within the high-risk group, delayed RT, defined as RT after at least one line of chemotherapy, was associated with a further decrement in overall survival (p=0.021).

**Conclusion.** A high-risk subgroup of patients, defined by diffuse astrocytoma histology or midbrain/thalamus tumor location, have suboptimal long-term survival and might benefit from timely use of RT. These results require validation.

**Keywords:** pediatric low-grade glioma, radiation, risk stratification, survival

# **Key points:**

- Management of unresectable pediatric low-grade glioma is controversial.
- We identify and validate a high-risk cohort with suboptimal long-term survival.
- This high-risk cohort might benefit from timely use of radiation therapy.

## **Importance of Study:**

There are no validated prognostic features to guide use of radiation therapy in unresectable pediatric low-grade gliomas and glioneuronal tumors. Optimal management of these tumors is controversial. Using clinicopathologic and molecular data from patients treated homogenously with radiation therapy, we identified and validated low- and high-risk groups with significantly disparate 10-year overall survival. Risk stratification remained significant independent of BRAF alteration (V600E or fusion). Within the high-risk group, delayed radiation therapy, defined as radiation therapy administered after at least one line of chemotherapy, was associated with a further decrement in overall survival. High-risk tumors, defined by diffuse astrocytoma histology or thalamic/midbrain location, might benefit from timely use of radiation therapy.

## Introduction

Pediatric low-grade gliomas (LGGs) and glioneuronal tumors (LGGNTs) represent heterogeneous diseases with multiple treatment options and varied outcomes depending on clinicopathologic variables and treatment modality. Several selection biases drive the choice of therapy. For tumors that can be totally resected, no adjuvant therapy is required and outcomes are usually excellent.<sup>2,3</sup> For unresectable tumors, optimal management is controversial. Despite the efficacy of radiation therapy (RT) in locally controlling unresectable disease, concerns regarding RT-related adverse effects<sup>4</sup> in young children have resulted in a preference for initially administering chemotherapy or targeted therapy, both of which are associated with inferior progression-free survival (PFS) when compared to RT.<sup>5-9</sup> RT is recommended for older children or for those whose tumors have progressed on systemic therapy. For patients who are candidates for RT, there are no validated criteria for risk stratification based on overall survival (OS) and the timing of RT remains controversial as it is assumed that all patients can be successfully salvaged with RT. As a result, patients might experience significant morbidity and mortality due to repeated tumor progressions prior to RT in an otherwise curable disease. <sup>10</sup> This highlights the need for a data-driven approach, incorporating both clinicopathologic and molecular data, to risk-stratify this patient population, with the goal of facilitating treatment decisions and thereby optimizing outcomes for these patients.

We used a cohort of patients who were uniformly treated with RT at St. Jude Children's Research Hospital (St. Jude) with a median follow-up of greater than 10 years to identify critical clinicopathologic and molecular variables associated with OS. Patents were treated either on an institutional phase II study, RT1, (ClinicalTrials.gov identifier: NCT00187226), or according to Children's Oncology Group Study, ACNS0221 (ClinicalTrials.gov identifier:

NCT00238264). *BRAF* status was available for some patients. Our focus was on OS, in order to identify high-risk patients who could not be successfully salvaged. We hypothesized that clinicopathologic features along with molecular data could be used to risk-stratify patients based on OS. We employed recursive partitioning analysis (RPA) to divide patients with LGG/LGGNT into two prognostic groups with clinical utility and subsequently validated the model using an external prospective dataset from the Children's Cancer and Leukemia Group's (CCLG) CNS9702 trial, which included patients who were treated in a similar manner to those at St. Jude.

### **Materials and Methods**

Study Population – St. Jude Cohort

Patients with LGG/LGGNT treated with focal RT between 1997 and 2017 at St. Jude were identified. Indications for RT included radiographic progression of disease, symptomatic disease or deterioration in visual field/acuity. Patients could have been treated with chemotherapy prior to RT. Those who underwent RT between 1997 and 2010 were treated on a phase II institutional protocol, RT1, (ClinicalTrials.gov identifier: NCT00187226) and those who underwent RT after 2010 were treated according to Children's Oncology Group Study, ACNS0221 (ClinicalTrials.gov identifier: NCT00238264). Exclusion criteria are shown in the CONSORT diagram (Figure 1). A total of 150 patients were included for analysis. Tumors in relatively uncommon anatomic sites such as the spinal cord (n=3, 2 had metastatic disease) and cerebral cortex (n=8, 4 were diagnosed as LGG, not otherwise specified) were excluded. Patients who did not undergo a biopsy (n=19) or patients whose tumors were diagnosed as LGG, not otherwise specified (n=12) due to insufficient biopsy material were excluded. Additionally, patients with metastatic disease (n=9), and prior malignancy (n=1) were also excluded due to the known poor prognosis in this subgroup. Rare

histological entities represented by only a single patient, such as angiocentric glioma (n=1), neurocytoma (n=1) and gangliocytoma (n=1), were also excluded. The study was approved by the St. Jude Institutional Review Board (#Pro00009006).

Study Population – CNS9702 Cohort

The external validation dataset, CNS9702, was a population-based study that enrolled patients younger than 16 years of age with LGG/LGGNT across 22 treatment centers in the United Kingdom. Treatment strategy, methodology and results for the entire study population have been previously reported. The dataset we used for validation consisted of patients on CNS9702 who underwent RT as first line of therapy. The exclusion and inclusion criteria for the external validation dataset were identical to that of the St. Jude dataset (Figure 1). A total of 48 patients were included for analysis. As with the St. Jude dataset, the most common reason for exclusion was lack of a biopsy (n=12). Other reasons for exclusion included location (spinal cord: n=9, cerebral cortex: n=9), metastatic disease (n=4), the designation of LGG, not otherwise specified (n=2) and rare histologic entities represented by a single patient, such as dysembryoplastic neuroepithelial tumor (n=1).

Histopathologic Review

For both study cohorts, histopathology was centrally reviewed by a single neuropathologist (DWE). For each case, standard hematoxylin and eosin–stained histopathologic preparations were supplemented by immunohistochemistry on 5µm formalin-fixed paraffin-embedded (FFPE) tissue sections. For the St. Jude cohort, *BRAF* alterations were identified when tissue was available. A mouse monoclonal antibody (Ventana, cat. no. 7990-4855; prediluted) was used to detect *BRAF* V600E-mutant protein. <sup>12</sup> Interphase fluorescence *in situ* hybridization was used to detect chromosome 7q34 duplication, a surrogate marker for *KIAA1549-BRAF* fusion, using a probe developed in-house. <sup>13</sup> The presence of the histone *H3K27M* mutation was also detected by immunohistochemistry.

Surgery

For the St. Jude cohort, the extent of surgical resection was defined using postoperative MRI, along with clinical and operative data. Gross total resection (GTR) was defined as the removal of all tumor-related T1 enhancement and T2 FLAIR abnormality, excluding signal change thought to be post-surgical. Subtotal resection (STR) was defined as presence of residual tumor-related T1 enhancement and/or T2 FLAIR abnormalities visible on postoperative MRI. Biopsy was defined as sampling rather than resection of tumor. For the CNS9702 Cohort, primary surgical resection or biopsy was recommended for all patients except those with neurofibromatosis type 1 (NF1) and those with chiasmatic/hypothalamic tumors consistent with optic pathway glioma, as described previously. Extent of resection was based on the local operative report and on the postoperative MRI following the recommendation of the Brain Tumor Subcommittee. GTR was defined as without measurable disease. STR was defined as residual tumor of a measurable size.

Radiation Therapy

For the St. Jude cohort, RT was delivered with a 3D conformal or intensity-modulated radiation technique to a total dose of 54 GyRBE in 1.8 GyRBE fractions over 6 weeks. The gross tumor volume (GTV) included both the cystic and solid components of the tumor and was defined by T2/FLAIR hyperintensity and T1 enhancement (if present). In patients who underwent surgery before RT, the GTV was defined as the surgical bed and any residual T2/FLAIR hyperintensity or T1 enhancement. The clinical target volume (CTV) margin was 1 cm for patients treated from 1997 to 2006. The CTV margin was 0.5 cm for patients treated after 2006. For patients treated with photon therapy, a planning target volume (PTV) margin of 0.5 or 0.3 cm was used. For patients treated with pencil-beam scanning proton therapy, scenario-based optimization was used with a 3-mm positional uncertainty and a 3% range uncertainty. For tumors involving or adjacent to the optic pathway that were treated with

proton therapy, the total dose was reduced to 52.2 GyRBE to minimize risk of optic chiasm/nerve toxicity. For the CNS9702 Cohort, similar focal RT was delivered to 54 GyRBE for children >5 years and 50 GyRBE for children <5 years, as described previously. All tumors were treated with a CTV margin of 1-2 cm.

Radiographic Evaluation

For the St. Jude Cohort, all patients underwent brain MRI at baseline, every 3 months for the first 2 years, every 6 months through 5 years, and yearly thereafter. Radiographic response was categorized according to RECIST criteria, with radiographic progression being defined as >25% tumor growth or the appearance of new lesions. For the CNS9702 cohort, tumor progression was defined by local radiologic report and/or symptomatic progression, as described previously.

Outcome Measures and Variable Definitions

The primary outcome measure was OS. Secondary outcomes were progression-free survival (PFS) and cumulative incidence of secondary malignancies. Survival time was calculated from start of RT to death or last follow-up, whichever occurred first. Time to progression was calculated from start of RT to progression, death or last follow-up, whichever occurred first. All outcome measures were calculated in an identical fashion for both cohorts. Tumor size, measured prior to RT, was defined as the largest tumor diameter, and was dichotomized as ≥6cm and <6cm, consistent with adult LGG studies. <sup>16</sup> Tumors involving the optic pathways and/or hypothalamus were grouped together, consistent with published data. <sup>17</sup> Tumors involving the thalamus and/or midbrain were grouped together, as were those involving the pons and/or medulla, as both structures were often involved. Delayed RT was defined as RT administered after at least one line of systemic therapy.

### Statistical Analysis

Frequency distributions between groups were assessed with the Fisher exact test for categorical variables and Wilcoxon rank sum test for continuous variables. Survival analysis was computed using the Kaplan-Meier method and log-rank test was used to compare curves. Prognostic variables were identified via Cox proportional hazards. Variables significant on Cox univariate analysis (p<0.05) were considered for multivariate analysis. Cumulative incidence of secondary malignancies was estimated using death as a competing risk. All statistical tests were 2-sided. Statistical analyses were performed with Stata software (2014 release, StataCorp, College Station, TX).

Recursive partitioning analysis (RPA) was used to devise high-risk and low-risk prognostic groups based on OS. RPA divides patients into homogeneous groups based on chosen covariates with respect to a predetermined outcome parameter, such as OS.<sup>18</sup> We employed classification and regression tree analysis (CART) <sup>19</sup> in Stata to generate statistically significant divisions (p≤0.05), with respect to OS, based on variables that were significant on Cox multivariate analysis with the exception of tumor size. CART in Stata is specifically for failure time data and uses the martingale residuals of a Cox model to approximate chi-square values for all variables.<sup>20</sup> Tumor size was not included because measurement methods varied across patients. Patients treated prior to 2000 were more likely to have CT-based measurements and patients treated 2000 onwards were more likely to have MRI-based measurements. The minimum size for each subgroup was 10. The St. Jude dataset was used to construct a risk-stratification model. The CNS9702 dataset from the United Kingdom was used for external validation of the resulting model.

#### **Results**

Patient Characteristics - St. Jude Cohort

Median follow up was 11.4 years (Table 1). Median age at time of RT was 8 years. The following four anatomic sites were included in the analysis: optic pathway/hypothalamus, thalamus/midbrain, pons/medulla and cerebellum. The following histologic diagnoses were included in the analysis: pilocytic astrocytoma, diffuse astrocytoma and ganglioglioma. Forty percent of tumors involved the optic pathway or hypothalamus and 26% involved the thalamus or midbrain. The majority of tumors (81%) were pilocytic astrocytomas followed by diffuse astrocytoma (12%) and ganglioglioma (7%). Most patients underwent either a biopsy (53%) or STR (47%) prior to RT. Approximately a third of the patients (34%) received at least one course of chemotherapy prior to RT. The mean RT dose was 54 GyRBE and most patients were treated with photon therapy.

Prognostic Factors For Overall Survival and Progression-Free Survival – St. Jude Cohort

The 10-year OS and PFS for the entire St. Jude cohort was 90% (95% CI: 82.7 – 93.8) and
66% (95% CI: 56.6% - 73.3%), respectively. On multivariate analysis, the following
characteristics were associated with increased risk of mortality: thalamus/midbrain location
versus optic pathway/hypothalamic location (Hazard ratio [HR]: 6.02, 95% Confidence
Interval [CI]: 1.35 – 26.73; p=0.018), diffuse astrocytoma versus pilocytic astrocytoma (HR:
3.76, 95% CI: 1.18 – 11.94; p=0.025), tumor size ≥6 cm versus tumor size <6cm (HR: 32.6,
95% CI: 7.72 – 137. 7; p<0.001) (Table 2).

Inclusion of cerebral cortex tumors with a specific LGG histology (n=4) did not change the results of our original multivariate analysis (Supplemental Table 1). Given that extent of resection may be correlated with tumor size and tumor location, we performed a subgroup

analysis in patients who underwent biopsy alone and those who underwent STR. In the subgroup of patients who underwent a biopsy (n=80), thalamus/midbrain location compared to optic pathway/hypothalamus location remained associated with an increased risk of mortality (HR: 15.6, 95% CI: 2.01 - 121.74, p=0.009), and tumor size  $\geq$ 6cm compared to tumor size  $\leq$ 6cm also remained associated with an increased risk of mortality (HR: 7.38, 95% CI: 1.58 - 34.56, p=0.011). Similarly, in the subgroup of patients who underwent a STR (n=70), thalamus/midbrain location compared to optic pathway/hypothalamus location remained associated with an increased risk of mortality (HR: 5.77, 95% CI: 1.6 - 20.84, p=0.007), and tumor size  $\geq$ 6cm compared to tumor size  $\leq$ 6cm also remained associated with an increased risk of mortality (HR: 11, 100007).

On multivariate analysis, the following characteristics were associated with increased risk of progression: tumor size  $\geq$ 6 cm compared to <6cm (HR: 2.84, 95% CI: 1.1 – 7.32; p=0.031) and delayed RT (HR: 2.53, 95% CI: 1.44 – 4.44; p=0.001) (Table 3). Delayed RT was defined as RT administered after at least one line of chemotherapy. Larger tumor size was correlated with delayed RT (p=0.002).

Risk Groups Derived From RPA Independently Predict Survival – St. Jude Cohort

All patients from St. Jude (n=150) were used for RPA modeling. Only variables significant on multivariate analysis with the exception of tumor size were included as input elements for the RPA. The resulting model had two splits and three terminal nodes, which were simplified into two risk groups (Figure 2) with statistically significant differences in OS rates (Figure 3A). High-risk group consisted of tumors with diffuse astrocytoma histology or location within the thalamus/midbrain. Low-risk group consisted of tumors located outside of the thalamus/midbrain with either pilocytic astrocytoma or ganglioglioma histology. The 10-year OS for the low-risk (n=105) and high-risk group (n=45) was 95.6% vs. 76.4% (95% CI:

88.7% - 98.4% vs. 59.3% – 87.1%, p=0.003), respectively. The difference in OS between low-risk and high-risk groups increased over time with a 7% absolute difference at 5 years (5-year OS: 90.5% vs. 97%) and a 19.2% absolute difference at 10 years. Within the high-risk group, patients who underwent delayed RT had a significantly worse OS when compared to those who underwent RT as first line therapy (p=0.021, Figure 3D). In the low-risk group, timing of RT did not influence survival (p=0.061, Figure 3E). The risk groups remained independently prognostic for OS after adjusting for other variables such as tumor size and delayed RT (high-risk vs. low-risk HR: 10.37, 95% CI: 3.5 – 30.76; p<0.001) (Supplemental Table 2).

Risk Groups Retain Prognostic Significance Among Tumors with Molecular Data – St. Jude cohort

Of the 49 tumors that were tested for *BRAF* alterations, 59% harbored a *BRAF*-fusion, most commonly *KIAA1549-BRAF*, and 16% harbored a BRAF *V600E* mutation (Table 1). Patients with tumors for which molecular data were available had been treated more recently and therefore had shorter follow-up (median: 5.8 years) (p<0.01) and were treated with more advanced RT techniques (p<0.001) when compared to the entire cohort. High-risk group was the only variable predictive of OS (HR: 15.38, 95% CI: 1.36 – 173.87; p=0.027) (Supplemental Table 3). The 6-year OS for low-risk (n=42) and high-risk group (n=7) was 97.6% vs. 41.7% (95% CI: 84.3% – 99.6% vs. 1.12% - 84.2%, p=0.003), respectively (Figure 3B). Tumors harboring *BRAF* V600E were not associated with worse OS or PFS when compared to tumors harboring BRAF fusions (Supplemental Table 4). None of the tested thalamic or midbrain tumors (n=26) harbored H3 K27M.

Risk Groups Are Externally Validated Using CNS9702 Cohort

A total of 48 patients from CNS9702 met inclusion criteria. Median follow up was 3.8 years from start of RT. A table of patient and tumor characteristics from CNS9702 can be found in Supplemental Table 5. Thirty-one percent of tumors involved the optic pathway/hypothalamus, 17% involved the thalamus/midbrain, 38% involved the pons/medulla and 15% involved in the cerebellum. The two histologies represented in the dataset were pilocytic astrocytoma (85%) and diffuse astrocytoma (15%). The 4-year OS for the low-risk group (n=35) was 100%, and that for the high-risk group (n=13) was 64.17% (95% CI: 30.2% – 84.3%; p<0001) (Figure 3C).

Secondary Malignancies – St. Jude Cohort

There were 13 subsequent malignancies, 11 of which occurred within the radiation field. Secondary malignancies within the RT field occurred a median of 9.05 years after the start of RT and included the following histologic diagnoses: anaplastic astrocytoma, gliosarcoma, mucoepidermoid carcinoma of the parotid gland, glioblastoma, and meningioma. The 15-year cumulative incidence of second malignancies was 7.03% (95% CI: 3.2% – 12.7%). One patient with a secondary malignancy had a known underlying genetic cancer predisposition syndrome (NF1). At last follow-up, three of the 11 patients with secondary malignancies were alive.

### **Discussion**

The current management paradigm for unresectable LGG/LGGNT supports an approach of initially administering chemotherapy or targeted therapy and reserving RT for subsequent progressions with the assumption that disease outcome will not be affected by delaying RT. While this strategy is appropriate in certain patients, others might benefit from earlier

initiation of RT. Currently, there are no validated criteria to guide use of RT. Our risk-stratification scheme addresses this significant gap in knowledge. We demonstrate that children in the high-risk group with diffuse astrocytoma or thalamic/midbrain tumors have significantly worse OS compared to the low-risk group and delaying RT in high-risk patients is associated with a further decrement in OS. This decrease in OS was not apparent until after approximately 8 years of follow-up, highlighting the necessity of reporting long-term outcomes in this patient population and the need to validate these results in a larger cohort. RT timing did not influence survival in the low-risk patients. Delayed RT was also associated with worse PFS. We validated this risk-stratification in an external dataset and showed that risk groups retained prognostic significance independent of *BRAF* status. Our findings have important implications for unresectable LGG/LGGNTs and challenge the current treatment paradigm.

It is difficult to compare our results to existing literature due to possible differences in reporting of histopathology and tumor location, as well as lack of long-term follow-up and adequate patient numbers. Interobserver agreement on histopathology can vary. 21,22

Therefore, central review by an experienced neuropathologist is critical and highlights a strength of our study. Although a large number of patients were treated on RT1, the initial report of RT1 did not analyze tumor histopathology or location with respect to OS or PFS. ACNS 0221 demonstrated inferior OS in patients with non-pilocytic phenotype with a median follow-up of 5 years; however, diffuse astrocytoma was not analyzed as a separate covariate. 23 Tumor location was also not analyzed with respect to OS or PFS. Indelicato et al. published outcomes after proton therapy in 174 patients with a median follow-up of 4 years. 24 Only age was associated with improved OS. Lower RT dose, as well as brainstem location, was associated with lower PFS on univariable analysis. However, since brainstem tumors were uniformly treated to a lower dose, collinearity of these two variables did not permit a

multivariable analysis. Furthermore, thalamic/midbrain tumors were not categorized separately. Other studies reporting outcomes after proton or photon therapy represent small and heterogeneous patient populations, limiting useful comparisons. <sup>25–27</sup> However, outcomes in children treated with chemotherapy have shown that supratentorial midline tumors <sup>28</sup> and thalamic tumors <sup>5</sup> are associated with worse outcomes. Diffuse astrocytoma has also been associated with worse PFS in the initial report of CNS9702, which included patients who were observed after surgery or treated with either chemotherapy or RT. <sup>11</sup>

None of the above studies has incorporated molecular data on BRAF alterations. There is a known association between pilocytic astrocytoma and KIAA1549-BRAF fusion, as approximately 70% of pilocytic astrocytomas harbor this fusion.<sup>29</sup> It is unclear whether the fusion is associated with improved prognosis independent of tumor type, as the two are highly correlated. BRAF V600E mutation has been reported across multiple tumor types, including pilocytic astrocytoma, (pediatric-type) diffuse astrocytoma, ganglioglioma and pleomorphic xanthoastrocytoma. 13,30,31 Whether this mutation is associated with independent prognostic significance is controversial. Lassaletta et al. reported inferior PFS in BRAF V600E-mutant tumors in a heterogeneous patient population treated with surgery alone, chemotherapy or RT.<sup>32</sup> Tumor location and histopathology were not included in the analysis as covariates, questioning the independent prognostic significance of BRAF V600E.33 Our data shed some light on this topic, as we included both molecular and clinicopathologic data in our analysis. Furthermore, all patients were treated in a consistent fashion with RT, eliminating confounding effects from treatment heterogeneity. We did not find that BRAF V600E mutation was significantly associated with worse PFS or OS when compared to BRAF fusion in the subset of patients with molecular data (n=49). More importantly, our risk stratification scheme was able to demonstrate a significant difference in OS between low- and high-risk groups in this subset of patients with molecular data, suggesting that

tumor location and histopathology likely prevails over *BRAF* alteration for LGG/LGGNT treated with RT.

Although OS was the primary outcome measure, we also analyzed PFS. We found that tumor size and delayed RT were associated with worse PFS. Tumor size has been correlated with outcomes in prior studies and was also correlated with OS in our analysis. The association between delayed RT and worse PFS might be explained through two mechanisms. First, tumors treated with chemotherapy prior to RT tended to be larger at the time of RT, and large tumor size is associated with worse PFS. However, even after accounting for tumor size, delayed RT remained independently associated with worse PFS. Therefore, a second explanation might be that pretreated tumors have shorter PFS compared to tumors that have not been pretreated. Pretreated tumors might be inherently more aggressive and have acquired additional deleterious mutations over time. The phenomenon of temporal genomic heterogeneity has been described in the context of pediatric gliomas. 34,35

The rationale for delaying or avoiding RT in young children is to reduce the risk of late effects, such as cerebral vasculopathy, neurocognitive impairment, endocrine deficiencies and secondary malignancies. In the context of pediatric brain tumors, the reported cumulative incidence of a second malignancy is approximately 6% to 8% at 20 to 30 years, which is comparable to our result of 7% at 15 years. So, In the setting of LGG, distinction between a secondary malignancy and transformation can be challenging – approximately 3% of pediatric LGG may transform to a secondary high-grade glioma. Some patients with LGG may also harbor cancer predisposition syndromes, such as NF1, and be more susceptible to develop a subsequent malignancy. Along with young age, region and volume of brain being irradiated are significant risk factors for vasculopathy, neurocognitive effects and endocrine deficiencies. Irradiation of optic pathway gliomas, adjacent to the circle of Willis, can result

in vasculopathy,<sup>39</sup> irradiation of the hippocampus can impair memory and learning,<sup>40</sup> and irradiation of the pituitary-hypothalamic axis can result in endocrine deficiencies.<sup>41</sup>

There are several limitations to our analysis. BRAF alterations were only determined in a subset of St. Jude patients (n=49). Similarly, absence of H3 K27M mutation was confirmed in a subset of patients with thalamic/midbrain tumors for whom data were available (n=26). Additionally, the validation cohort from CNS9702 did not contain any molecular data. Although we did not find any statistically significant difference in outcome by BRAF alteration, our subset with molecular data was limited. Therefore, we cannot exclude the possibility that BRAF alterations might be associated with a difference in outcome. With additional molecular data, as well as MRI-based tumor measurements, further refinement of the risk stratification scheme might be possible. There is some difficulty in interpreting tumor size over a two-decade timespan with evolving imaging technologies from CT to highresolution MRI. Secondly, although RPA is a clinically useful tool to divide patients into prognostic groups, the results can be highly dependent on patient population and input variables. Therefore, it is reassuring that we can reproduce statistically significant differences in the low-risk and high-risk groups using an independent dataset. Finally, application of our risk stratification scheme should be limited to unresectable LGG/LGGNT treated with RT. The validity of this risk scheme in other clinical contexts is unknown.

We have identified a high-risk group of patients with diffuse astrocytoma or thalamic/midbrain tumors associated with inferior survival and validate these results using an external dataset. Risk grouping retained prognostic significance independent of BRAF alteration. In this high-risk group, delayed RT was associated with a reduction in OS, suggesting that these patients might benefit from timely RT.

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## **Figure Legends**

# Figure 1. CONSORT Diagram

Figure 2. Results of recursive partitioning analysis of St. Jude cohort

**Figure 3.** Overall survival stratified by risk group for: (A) the entire St. Jude cohort, (B) the patients in the St. Jude cohort with molecular data stratified by risk group, and (C) the external validation dataset (the CNS9702 cohort). Overall survival stratified by timing of RT in: (D) high-risk group, and (E) low-risk group. Delayed RT is defined as RT after at least one line of chemotherapy.

Table 1. Patient and Tumor Characteristics of the St. Jude Cohort

Variable	Number	Number for patients with molecular data (N = 49)	<b>P</b> α <0.01	
Median follow-up years (range)	11.4 (0.24–29.4)	5.83 (0.24–20.6)		
Median age in years (range)	8 (1.2–20)	7.9 (2.2–18.4)	0.616	
Sex (%)	,	•	0.384	
Female	71 (47)	26 (53)		
Male	79 (53)	23 (50)		
Race (%)	, ,	. ,	0.416	
White	114 (77)	35 (71)	X	
Black	28 (19)	12 (25)		
Other	8 (5)	2 (4)	* *	
NF1 (%)	- (- /	( )	0.428	
Yes	7 (5)	1 (2)		
No	143 (95)	48 (97)		
Tumor location (%)	173 (33)	70 (71)	0.029	
OPG/hypothalamus	50 (40)	20 (41)	0.029	
Thalamus/midbrain	59 (40) 30 (26)	20 (41)		
Pons/medulla	39 (26)	6 (12)		
Cerebellum	34 (23)	15 (31)		
	18 (12)	8 (16)	Λ 101	
Histology (%)	120 (01)	41 (04)	0.181	
Pilocytic astrocytoma	120 (81)	41 (84)		
Diffuse astrocytoma	19 (12)	3 (6)		
Ganglioglioma  PRAF - 14 - 14 - 10 (0/)	11 (7)	5 (10)		
BRAF alteration $(N = 49)$ (%)	0 (16)	0 (16)		
BRAF V600E	8 (16)	8 (16)		
BRAF fusion	29 (60)	29 (59)		
BRAF wildtype	12 (25)	12 (25)	0.472	
Tumor size before RT (%)	145 (07)	40 (00)	0.472	
<6 cm	145 (97)	48 (98)		
≥6 cm	5 (3)	1 (2)		
Surgical extent (%)	00 (70)		0.002	
Biopsy	80 (53)	16 (33)		
STR	70 (47)	33 (67)		
Number of surgeries before RT	101 (10)	-0 ()	0.079	
(%)	101 (69)	28 (57)		
	38 (26)	18 (37)		
2	8 (5)	3 (6)		
>2				
Chemotherapy before RT (%)	51 (QA)	20 (41)	0.271	
Yes	51 (34)	20 (41)		
No	99 (66)	29 (59)		
Radiation modality (%)	444 (= =	<b>22</b> (1 <b>-</b> )	< 0.001	
3D CRT	114 (76)	23 (47)		
IMRT	28 (19)	19 (38)		
Proton therapy	8 (5)	7 (14)		
Mean radiation dose in GyRBE (range)	54 (48–55.8)	54 (50.4–54)	0.329	

<sup>&</sup>lt;sup>a</sup>P value for frequency distributions based on Fisher exact test for categorical variables and Wilcoxon rank-sum test for continuous variables.

NF1 = neurofibromatosis-1; OPG = optic pathway glioma; RT = radiation therapy; GTR = gross total resection; STR = subtotal resection; CRT = conformal radiation therapy; IMRT = intensity-modulated radiation therapy

Table 2. Univariate and multivariate analysis of overall survival in the St. Jude cohort

Variable         HR (95% CI)         P         HR (95% CI)         P           Sex         Male         1		<u>Univariate Analysis</u>		<u>Multivariate</u>	
Sex   Male		· · · · · · · · · · · · · · · · · · ·			
Male Female       1       0.77 (0.30-1.97)       0.590          Race  <	Variable	HR (95% CI)	P	HR (95% CI)	P
Female	Sex				
Race       Caucasian       1         Black       1.23 (0.35-4.33)       0.743         Other       2.58 (0.71-9.35)       0.148         Age at RT       0.99 (0.89-1.10)       0.853         Location       To PG/hypothalamus       1         OPG/hypothalamus/midbrain       4.0 (1.40-11.35)       0.009       6.02 (1.35-26.73)       0.018         Pons/medulla       1.38 (0.33-5.81)       0.661       3.14 (0.54-18.0)       0.199         Cerebellum       No events       -       No events         Histology       Filocytic astrocytoma       1       3.76 (1.18-11.94)       0.025         Ganglioglioma       3.25 (0.70-415.0)       0.131       4.93 (1-24.1)       0.051         Tumor size before RT       <6 cm	Male	1		X	
Caucasian       1         Black       1.23 (0.35-4.33)       0.743         Other       2.58 (0.71-9.35)       0.148         Age at RT       0.99 (0.89-1.10)       0.853         Location       0PG/hypothalamus       1         Thalamus/midbrain       4.0 (1.40-11.35)       0.009       6.02 (1.35-26.73)       0.018         Pons/medulla       1.38 (0.33-5.81)       0.61       3.14 (0.54-18.0)       0.199         Cerebellum       No events       -       No events         Histology       Pilocytic astrocytoma       1       -       No events         Biffuse astrocytoma       1       -       0.001       3.76 (1.18-11.94)       0.025         Ganglioglioma       3.25 (0.70-15.0)       0.131       4.93 (1-24.1)       0.051         Tumor size before RT       < cm	Female	0.77 (0.30-1.97)	0.590		
Black Other       1.23 (0.35-4.33) 0.743 0.148         Age at RT       0.99 (0.89-1.10) 0.853         Location OPG/hypothalamus Thalamus/midbrain       4.0 (1.40-11.35) 0.009 0.009 0.02 (1.35-26.73) 0.018         Pons/medulla Pons/medulla No events       1.38 (0.33-5.81) 0.661 0.14 (0.54-18.0) 0.199 0.1	Race			* (	
Other         2.58 (0.71–9.35)         0.148           Age at RT         0.99 (0.89–1.10)         0.853           Location         OPG/hypothalamus         1           OPG/hypothalamus/midbrain         4.0 (1.40–11.35)         0.009         6.02 (1.35–26.73)         0.018           Pons/medulla         1.38 (0.33–5.81)         0.661         3.14 (0.54–18.0)         0.199           Cerebellum         No events          No events           Histology         Pilocytic astrocytoma         1          No events           Pilocytic astrocytoma         1          No events           Brass         1          No events           Ganglioglioma         3.25 (0.70–15.0)         0.131         4.93 (1–24.1)         0.051           Tumor size before RT         46 cm         1            0.001           BRAF fusion         1             No          No          No          NS          NS          NS          NS          NS          NS          NS          NS	Caucasian	1			
Age at RT       0.99 (0.89–1.10)       0.853         Location       1         OPG/hypothalamus       1         Thalamus/midbrain       4.0 (1.40–11.35)       0.009       6.02 (1.35–26.73)       0.018         Pons/medulla       1.38 (0.33–5.81)       0.661       3.14 (0.54–18.0)       0.199         Cerebellum       No events       -       No events         Histology       1       No events       -       No events         Histology       1       -       No events       -       No events         Brillian Strocytoma       1       -       -       0.001       3.76 (1.18–11.94)       0.025         Ganglioglioma       3.25 (0.70–15.0)       0.131       4.93 (1–24.1)       0.051         Tumor size before RT       46 cm       1       -       -       -       0.001         **6 cm       1       0.58 (3.67–30.52)       <0.001	Black	1.23 (0.35-4.33)	0.743		
Location         OPG/hypothalamus         1           Thalamus/midbrain         4.0 (1.40–11.35)         0.009         6.02 (1.35–26.73)         0.018           Pons/medulla         1.38 (0.33–5.81)         0.661         3.14 (0.54–18.0)         0.199           Cerebellum         No events         -         No events           Histology         Pilocytic astrocytoma         7.49 (2.91–19.23)         <0.001	Other	2.58 (0.71–9.35)	0.148		
Location         OPG/hypothalamus         1           Thalamus/midbrain         4.0 (1.40–11.35)         0.009         6.02 (1.35–26.73)         0.018           Pons/medulla         1.38 (0.33–5.81)         0.661         3.14 (0.54–18.0)         0.199           Cerebellum         No events         -         No events           Histology         Pilocytic astrocytoma         7.49 (2.91–19.23)         <0.001	Age at RT	0.99 (0.89–1.10)	0.853		
Thalamus/midbrain         4.0 (1.40−11.35)         0.009         6.02 (1.35−26.73)         0.018           Pons/medulla         1.38 (0.33−5.81)         0.661         3.14 (0.54−18.0)         0.199           Cerebellum         No events         -         No events           Histology         Pilocytic astrocytoma         1         -         No events         -         No events         0.025         Ganglioglioma         3.25 (0.70−15.0)         0.131         4.93 (1−24.1)         0.051         0.051         -         -         0.051         -	·	•			
Thalamus/midbrain         4.0 (1.40−11.35)         0.009         6.02 (1.35−26.73)         0.018           Pons/medulla         1.38 (0.33−5.81)         0.661         3.14 (0.54−18.0)         0.199           Cerebellum         No events         -         No events           Histology         Pilocytic astrocytoma         1         -         No events         -         No events         0.025         Ganglioglioma         3.25 (0.70−15.0)         0.131         4.93 (1−24.1)         0.051         0.051         -         -         0.051         -	OPG/hypothalamus	1			
Pons/medulla Cerebellum         1.38 (0.33–5.81) No events         0.661 3.14 (0.54–18.0) No events         0.199           Histology Pilocytic astrocytoma Diffuse astrocytoma         1         1         1         1         0.025         3.76 (1.18–11.94)         0.025         0.001         3.76 (1.18–11.94)         0.025         0.001         3.76 (1.18–11.94)         0.025         0.001         3.76 (1.18–11.94)         0.025         0.001         3.76 (1.18–11.94)         0.025         0.001		4.0 (1.40–11.35)	0.009	6.02 (1.35–26.73)	0.018
Cerebellum         No events          No events           Histology Pilocytic astrocytoma Diffuse astrocytoma         1          No events           Ganglioglioma         7.49 (2.91–19.23)         <0.001	Pons/medulla	· · · · · · · · · · · · · · · · · · ·	0.661		0.199
Pilocytic astrocytoma       1         Diffuse astrocytoma       7.49 (2.91–19.23)       <0.001	Cerebellum			No events	
Pilocytic astrocytoma       1         Diffuse astrocytoma       7.49 (2.91–19.23)       <0.001	Histology	. 0			
Diffuse astrocytoma       7.49 (2.91–19.23)       <0.001       3.76 (1.18–11.94)       0.025         Ganglioglioma       3.25 (0.70–15.0)       0.131       4.93 (1–24.1)       0.051         Tumor size before RT        <6 cm	<b>0.</b>	1	<b>&gt;</b>		
Ganglioglioma       3.25 (0.70–15.0)       0.131       4.93 (1–24.1)       0.051         Tumor size before RT       46 cm       1       1       1       4       4       0.001       32.6 (7.72–137.7)       <0.001	· · ·	7.49 (2.91–19.23)	< 0.001	3.76 (1.18–11.94)	0.025
Tumor size before RT         <6 cm	——————————————————————————————————————		0.131		0.051
≥6 cm       10.58 (3.67–30.52)       <0.001       32.6 (7.72–137.7)       <0.001         BRAF alteration (N = 49)       BRAF fusion       1         BRAF WT       No events       8.83 (0.79–98.71)       0.077         Chemotherapy before RT         No       1       1         Yes       2.68 (1.11–6.48)       0.028        NS         RT dose (GyRBE)       1 (0.99–1)       0.804        NS         NF1       No       1				,	
BRAF alteration (N = 49)         BRAF fusion       1         BRAF WT       No events         BRAF V600E       8.83 (0.79–98.71)       0.077         Chemotherapy before RT         No       1       1         Yes       2.68 (1.11–6.48)       0.028        NS         RT dose (GyRBE)       1 (0.99–1)       0.804         NF1       No       1       1       1         Yes       1.07 (0.14–8.13)       0.945       0.945         Surgery       Biopsy       1       2       0.93 (0.31–2.88)       0.912 </td <td>&lt;6 cm</td> <td>1</td> <td></td> <td></td> <td></td>	<6 cm	1			
BRAF alteration (N = 49)         BRAF fusion       1         BRAF WT       No events         BRAF V600E       8.83 (0.79–98.71)       0.077         Chemotherapy before RT         No       1       1         Yes       2.68 (1.11–6.48)       0.028        NS         RT dose (GyRBE)       1 (0.99–1)       0.804         NF1       No       1       1       1         Yes       1.07 (0.14–8.13)       0.945       0.945         Surgery       Biopsy       1       2       0.93 (0.31–2.88)       0.912 </td <td>≥6 cm</td> <td>10.58 (3.67–30.52)</td> <td>&lt; 0.001</td> <td>32.6 (7.72–137.7)</td> <td>&lt; 0.001</td>	≥6 cm	10.58 (3.67–30.52)	< 0.001	32.6 (7.72–137.7)	< 0.001
BRAF fusion         BRAF WT       No events         BRAF V600E       8.83 (0.79–98.71)       0.077         Chemotherapy before RT         No       1       1         Yes       2.68 (1.11–6.48)       0.028        NS         RT dose (GyRBE)       1 (0.99–1)       0.804         NF1       No       1       2       0.93 (0.31–2.88)       0.912 <td></td> <td>30</td> <td></td> <td>· · · · · · · · · · · · · · · · · · ·</td> <td></td>		30		· · · · · · · · · · · · · · · · · · ·	
BRAF WT BRAF V600E       No events 8.83 (0.79–98.71)       0.077         Chemotherapy before RT No 1 Yes 2.68 (1.11–6.48)       0.028 NS         RT dose (GyRBE)       1 (0.99–1)       0.804         NF1 No 1	` ,	1			
BRAF V600E       8.83 (0.79–98.71)       0.077         Chemotherapy before RT       1       1         No       1       1         RT dose (GyRBE)       1 (0.99–1)       0.804         NF1       No       1         Yes       1.07 (0.14–8.13)       0.945         Surgery       Biopsy       1         STR       0.94 (0.31–2.88)       0.912         No surgery       0.86 (0.11–6.61)       0.885         Number of surgeries before RT       1       1         1       1       0.93 (0.31–2.88)       0.912		No events			
Chemotherapy before RT         No       1         Yes       2.68 (1.11–6.48)       0.028        NS         RT dose (GyRBE)       1 (0.99–1)       0.804         NF1       0.804       0.804         No       1       0.945         Surgery       0.945       0.945         Surgery       0.94 (0.31–2.88)       0.912         No surgery       0.86 (0.11–6.61)       0.885         Number of surgeries before RT       1       1         1       1       0.93 (0.31–2.88)       0.912			0.077		
No     1       Yes     2.68 (1.11-6.48)     0.028      NS       RT dose (GyRBE)     1 (0.99-1)     0.804       NF1     No     1       Yes     1.07 (0.14-8.13)     0.945       Surgery     Surgery       Biopsy     1       STR     0.94 (0.31-2.88)     0.912       No surgery     0.86 (0.11-6.61)     0.885       Number of surgeries before RT     1       1     1       2     0.93 (0.31-2.88)     0.912		,			
Yes       2.68 (1.11-6.48)       0.028        NS         RT dose (GyRBE)       1 (0.99-1)       0.804         NF1            No       1           Yes       1.07 (0.14-8.13)       0.945         Surgery            Biopsy       1           STR       0.94 (0.31-2.88)       0.912          No surgery       0.86 (0.11-6.61)       0.885          Number of surgeries before RT       1           1       1           2       0.93 (0.31-2.88)       0.912		1			
RT dose (GyRBE)       1 (0.99-1)       0.804         NF1           No       1          Yes       1.07 (0.14-8.13)       0.945         Surgery           Biopsy       1          STR       0.94 (0.31-2.88)       0.912         No surgery       0.86 (0.11-6.61)       0.885         Number of surgeries before RT       1       1         2       0.93 (0.31-2.88)       0.912		2.68 (1.11–6.48)	0.028		NS
NF1 No 1 Yes 1.07 (0.14–8.13) 0.945  Surgery Biopsy 1 STR 0.94 (0.31–2.88) No surgery 0.86 (0.11–6.61) 0.885  Number of surgeries before RT 1 1 2 0.93 (0.31–2.88) 0.912	RT dose (GvRBE)				
No       1         Yes       1.07 (0.14–8.13)       0.945         Surgery       1         Biopsy       1         STR       0.94 (0.31–2.88)       0.912         No surgery       0.86 (0.11–6.61)       0.885         Number of surgeries before RT       1       1         2       0.93 (0.31–2.88)       0.912		(			
Yes     1.07 (0.14-8.13)     0.945       Surgery     1       Biopsy     1       STR     0.94 (0.31-2.88)     0.912       No surgery     0.86 (0.11-6.61)     0.885       Number of surgeries before RT     1       1     1     0.93 (0.31-2.88)     0.912		1			
Surgery         Biopsy       1         STR       0.94 (0.31–2.88)       0.912         No surgery       0.86 (0.11–6.61)       0.885         Number of surgeries before RT       1       1         2       0.93 (0.31–2.88)       0.912			0.945		
Biopsy       1         STR       0.94 (0.31–2.88)       0.912         No surgery       0.86 (0.11–6.61)       0.885         Number of surgeries before RT       1       1         2       0.93 (0.31–2.88)       0.912		,			
STR       0.94 (0.31–2.88)       0.912         No surgery       0.86 (0.11–6.61)       0.885         Number of surgeries before RT       1       1         1       1       0.93 (0.31–2.88)       0.912		1			
No surgery       0.86 (0.11–6.61)       0.885         Number of surgeries before RT       1       1         1       1       0.93 (0.31–2.88)       0.912		0.94 (0.31–2.88)	0.912		
Number of surgeries before RT  1					
1 2 0.93 (0.31–2.88) 0.912		· · · · · /			
2 0.93 (0.31–2.88) 0.912	,	1			
			0.912		
	>2	0.86 (0.11–6.61)	0.885		

NF1 = neurofibromatosis-1; OPG = optic pathway glioma; RT = radiation therapy; GTR = gross total resection; STR = subtotal resection; CRT = conformal radiation therapy; IMRT = intensity-modulated radiation therapy; NS= not significant

Table 3. Variables significant on univariate analysis of progression-free survival in the St. Jude cohort

	<u>Univariate Analysis</u> <u>Analysis</u>		<u>Multivariate</u>	
$\mathbf{Variable}^{\beta}$	HR (95% CI)	P	HR (95% CI)	P
Age at RT	0.91 (0.85–0.98)	0.014	-	NS
Tumor size before RT				
≥6 cm	1			
<6 cm	3.89 (1.54–9.86)	0.004	2.84 (1.1–7.32)	0.031
Delayed RT <sup>α</sup>			)	
No	1			
Yes	2.73 (1.57–4.74)	< 0.001	2.53 (1.44–4.44)	0.001

<sup>&</sup>lt;sup>B</sup>Only variables that were significant on univariate analysis are listed.

RT = radiation therapy; NS= not significant

<sup>&</sup>lt;sup>a</sup>Delayed RT is defined as RT after at least one line of chemotherapy.

Figure 1

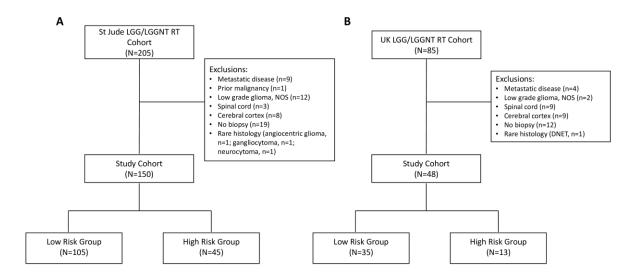




Figure 2

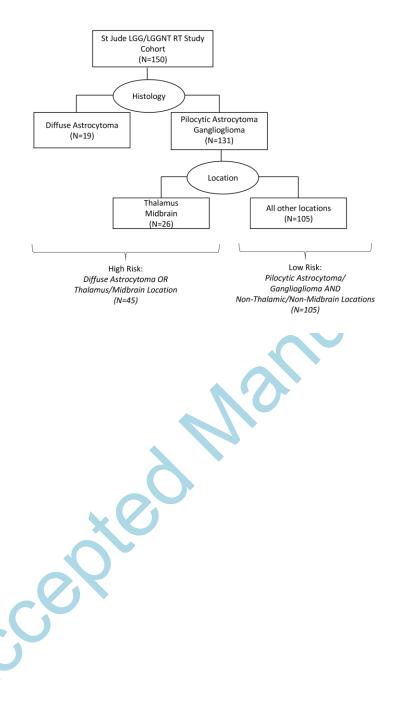


Figure-3

