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Prognostic Value of Preoperative MRI Metrics for Diffuse Lower-Grade Glioma Molecular Subtypes

 P. Darvishi,  P.P. Batchala,  J.T. Patrie,  L.M. Poisson,  M.-B. Lopes,  R. Jain,  C.E. Fadul,  D. Schiff, and  S.H. Patel

ABSTRACT

BACKGROUND AND PURPOSE: Despite the improved prognostic relevance of the 2016 WHO molecular-based classification of lower-grade gliomas, variability in clinical outcome persists within existing molecular subtypes. Our aim was to determine prognostically significant metrics on preoperative MR imaging for lower-grade gliomas within currently defined molecular categories.

MATERIALS AND METHODS: We undertook a retrospective analysis of 306 patients with lower-grade gliomas accrued from an institutional data base and The Cancer Genome Atlas. Two neuroradiologists in consensus analyzed preoperative MRIs of each lower-grade glioma to determine the following: tumor size, tumor location, number of involved lobes, corpus callosum involvement, hydrocephalus, midline shift, eloquent cortex involvement, ependymal extension, margins, contrast enhancement, and necrosis. Adjusted hazard ratios determined the association between MR imaging metrics and overall survival per molecular subtype, after adjustment for patient age, patient sex, World Health Organization grade, and surgical resection status.

RESULTS: For *isocitrate dehydrogenase* (*IDH*) wild-type lower-grade gliomas, tumor size (hazard ratio, 3.82; 95% CI, 1.94–7.75; $P < .001$), number of involved lobes (hazard ratio, 1.70; 95% CI, 1.28–2.27; $P < .001$), hydrocephalus (hazard ratio, 4.43; 95% CI, 1.12–17.54; $P = .034$), midline shift (hazard ratio, 1.16; 95% CI, 1.03–1.30; $P = .013$), margins ($P = .031$), and contrast enhancement (hazard ratio, 0.34; 95% CI, 0.13–0.90; $P = .030$) were associated with overall survival. For *IDH*-mutant 1p/19q-codeleted lower-grade gliomas, tumor size (hazard ratio, 2.85; 95% CI, 1.06–7.70; $P = .039$) and ependymal extension (hazard ratio, 6.34; 95% CI, 1.07–37.59; $P = .042$) were associated with overall survival.

CONCLUSIONS: MR imaging metrics offers prognostic information for patients with lower-grade gliomas within molecularly defined classes, with the greatest prognostic value for *IDH* wild-type lower-grade gliomas.

ABBREVIATIONS: LGG = lower-grade glioma; HR = hazard ratio; *IDH* = *isocitrate dehydrogenase*; *IDH*mut-Codel = *IDH* mutation and a whole-arm deletion of chromosome arms 1p and 19q; *IDH*mut-Noncodel = *IDH*-mutant lacking 1p/19q codeletion; *IDH*wt = *IDH* wild-type; IQR = interquartile range; OS overall survival; TCIA = The Cancer Imaging Archive; *TERT* = telomerase reverse transcriptase; WHO = World Health Organization

Recognition of the biologic and prognostic significance of molecular-based characterization of diffuse lower-grade gliomas (LGGs) heralded major revisions to their classification by the World Health Organization (WHO) in 2016.^{1–3} LGGs encompass WHO grade II and III astrocytic and oligodendroglial tumors, most of which have a prognostically favorable mutation in the *isocitrate dehydrogenase* (*IDH*) gene.^{2,4} Oligodendrogliomas are defined by the presence of both an *IDH* mutation and a whole-

arm deletion of chromosome arms 1p and 19q (*IDH*mut-Codel), which confer added prognostic and therapeutic favorability.^{5,6} *IDH*-mutant astrocytomas (*IDH*mut-Noncodel) lack 1p/19q codeletion and typically have tumor protein p53 and alpha-thalassemia/mental retardation syndrome X-linked (*ATRX* chromatin remodeler, [*ATRX*]) gene mutations.^{7,8} *IDH* wild-type LGGs (*IDH*wt) are generally associated with far worse clinical outcomes and largely comprise neoplasms that are genotypically similar to primary glioblastoma.^{1,9}

Despite improved clinical applicability of the revised classification system, there remains substantial heterogeneity in clinical outcomes within existing subtypes of LGGs.^{10–12} While numerous

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From the Departments of Radiology and Medical Imaging (P.D., P.P.B., S.H.P.); Public Health Sciences (J.T.P.); Pathology, Divisions of Neuropathology and Molecular Diagnostics (M.-B.L.); and Division of Neuro-Oncology (C.E.F., D.S.), University of Virginia Health System, Charlottesville, Virginia; Department of Public Health (L.M.P.), Henry Ford Health System, Detroit, Michigan; and Departments of Radiology (R.J.) and Neurosurgery (R.J.), New York University School of Medicine, New York, New York.

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Please address correspondence to Sohil H. Patel, MD, Department of Radiology and Medical Imaging, University of Virginia Health System, PO Box 800170, Charlottesville, VA 22908; e-mail: shp4k@virginia.edu
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studies have reported neuroimaging features that predict currently defined glioma molecular subtypes,^{13–20} neuroimaging features might additionally contain information that allows prognostic stratification of gliomas within currently defined molecular categories. The purpose of our investigation was to determine whether neuroimaging features on preoperative anatomic MR imaging have prognostic significance for LGGs within currently defined molecular categories.

MATERIALS AND METHODS

This was a retrospective study with institutional review board approval (University of Virginia Health System, Charlottesville, Virginia) as well as Health Insurance Portability and Accountability Act compliance.

Patient Selection

Cases were accrued from a diffuse LGG data base maintained at our institution ($n = 255$) as well as from The Cancer Imaging

Archive (TCIA) ($n = 198$).²¹ Inclusion criteria were the following: 1) known *IDH* mutation status and 1p/19q-codeletion status; 2) known surgical resection status (gross total resection, subtotal resection, or biopsy); and 3) available preoperative MR imaging including (at a minimum) precontrast T1WI, contrast-enhanced T1WI, and either T2WI or FLAIR.

From the institutional dataset, 178 cases met the inclusion criteria; 43 cases were excluded for lack of *IDH* and/or 1p/19q-codeletion status, and 34 cases were excluded for lack of requisite preoperative MR imaging. From the TCIA cohort, 128 cases met the inclusion criteria; 1 case was excluded for lack of *IDH* and 1p/19q-codeletion status and 69 cases lacked requisite preoperative MR imaging. In total, our study cohort included 306 cases. Overall survival times were available for all included patients, defined as the time between the date of pathologic diagnosis to the date of death or last contact.

Neuroimaging and Analysis

Two neuroradiologists with 6 and 14 years of experience, blinded to molecular status, WHO grade, patient demographics, and clinical outcome, analyzed the preoperative MR imaging to determine the following metrics:—1) tumor size: maximum long-axis diameter (centimeters); 2) location: any glioma signal abnormality/enhancement involving the basal ganglia, thalamus, or brain stem (yes = central; no = peripheral); 3) number (n) of involved lobes: each of the following counted as 1 lobe (per hemisphere)—frontal lobe, parietal lobe, temporal lobe, insula, occipital lobe, brain stem/cerebellum; 4) corpus callosum involvement: any glioma signal abnormality/enhancement involving the corpus callosum (yes or no); 5) hydrocephalus: (yes or no); 6) midline shift: greatest degree of contralateral brain displacement (centimeters); 7) eloquent cortex involvement: any glioma signal abnormality/enhancement involving the eloquent cortex, defined per Chang et al²² (yes or no); 8) ependymal extension: any glioma signal abnormality/enhancement involving the ventricular ependyma (yes or no); 9) margins: <33%, 33%–66%, or >66% sharp/circumscribed glioma margins; 10) contrast enhancement: any glioma contrast enhancement (yes or no); and 11) necrosis: any region of glioma necrosis characterized by peripheral contrast enhancement and central nonenhancement (yes or no).

Neuropathology

Pathologic data for patients in the institutional dataset were retrieved from the electronic medical record. Molecular markers were tested in the Clinical Laboratory Improvement Amendments–certified molecular pathology laboratory at our institution. A clinically validated *IDH1* antibody was used to determine *IDH* mutation status.^{23,24} Clinically validated pyrosequencing assays (PyroMark Q24 system; <https://www.qiagen.com/us/shop/automated-solutions/pyro-mark-q24/>) were used in immunohistochemistry-negative cases in accordance with the manufacturers to determine *IDH1/IDH2* mutation status.

Table 1: Patient characteristics for each molecular subtype

	<i>IDH</i> mut-Codel	<i>IDH</i> mut-Noncodel	<i>IDH</i> wt
Total patients ^a	104 (34.0%)	129 (42.2%)	73 (23.9%)
Female sex	58 (55.8%)	61 (47.3%)	35 (47.9%)
Age ^b	48 (17–74)	35 (17–70)	57 (19–76)
WHO grade			
II	72 (69.2%)	83 (64.3%)	34 (46.6%)
III	32 (30.8%)	46 (35.7%)	39 (53.3%)

^a Data are listed as count (percentage of total cohort).

^b Data are listed as median (range). Other data are listed as count (percentage of cases within given molecular subtype).

Table 2: Empiric distribution summaries for the preoperative MR imaging metrics according to LGG molecular subtype

MR Imaging Metric	<i>IDH</i> mut-Codel	<i>IDH</i> mut-Noncodel	<i>IDH</i> wt
Tumor size (cm)			
Median	6.6	6.9	6.2
IQR (1st to 3rd quartile)	(5.2–8.6)	(5.3–9.3)	(4.6–8.4)
Range (minimum–maximum)	(1.9–14.2)	(1.7–15.3)	(1.1–15.0)
Location			
Central	26 (25.0%)	43 (33.3%)	29 (39.7%)
Peripheral	78 (75.0%)	86 (66.7%)	44 (60.3%)
No. of involved lobes			
Median	2.0	2.0	2.0
IQR (1st to 3rd quartile)	(1.0–3.0)	(1.0–3.0)	(1.0–3.0)
Range (minimum–maximum)	(1.0–6.0)	(1.0–9.0)	(1.0–8.0)
Corpus callosum extension	44 (42.3%)	46 (35.7%)	18 (24.2%)
Hydrocephalus	12 (11.5%)	27 (20.9%)	7 (9.6%)
Midline shift (cm)			
Median	0.00	0.30	0.00
IQR (1st to 3rd quartile)	(0.00–0.43)	(0.00–0.80)	(0.00–0.10)
Range (minimum–maximum)	(0.00–2.70)	(0.00–1.90)	(0.00–1.40)
Eloquent cortex	33 (31.7%)	45 (34.9%)	24 (32.9%)
Ependymal extension	65 (62.5%)	80 (62.0%)	58 (79.5%)
Margin			
<33% sharp/circumscribed	32 (30.8%)	31 (24.0%)	28 (38.4%)
33%–66% sharp/circumscribed	55 (52.9%)	38 (29.5%)	31 (42.5%)
>66% sharp/circumscribed	17 (16.3%)	60 (46.5%)	14 (19.2%)
Contrast enhancement	21 (31.8%)	51 (39.5%)	30 (41.1%)
Necrosis	17 (16.3%)	14 (10.9%)	18 (24.7%)

The 1p/19q-codeletion status was determined by fluorescence in situ hybridization; dual-color human probes localizing the 1p, 1q, 19p, and 19q loci were used (Vysis LSI 1p36/1q25 and LSI 19q13/

19p13 FISH Probe Kit; Abbott Laboratories, Abbott Park, IL). Histopathologic and molecular data for the TCIA cohort were retrieved from supplemental material in Ceccarelli et al.²⁵

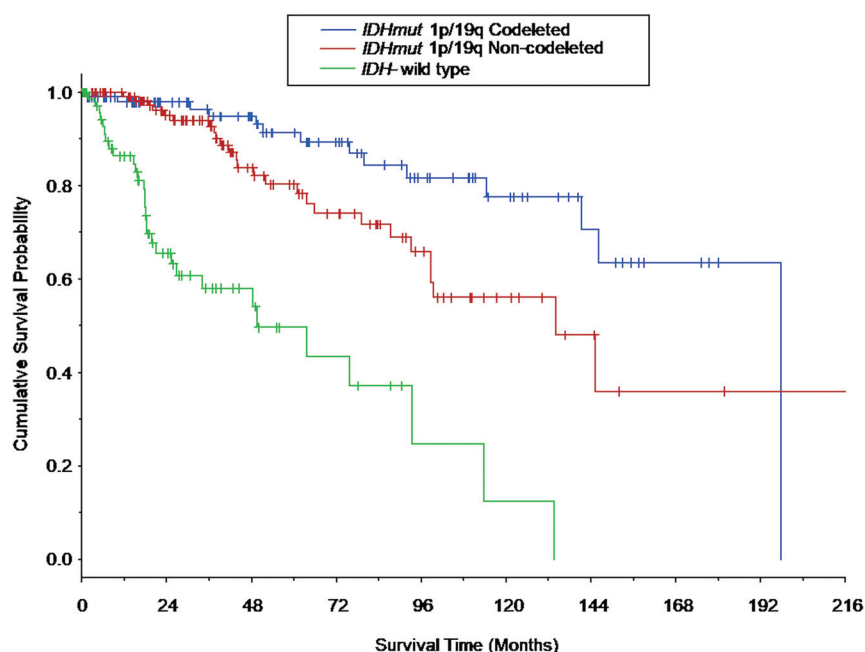


FIG 1. Kaplan-Meier survival curves based on the LGG molecular subtype. Vertical tick marks identify right-censored survival times. The survival curves differed among all 3 LGG molecular subtypes (*IDH*-mutant 1p/19q-codeleted molecular subtype versus *IDH*-mutant 1p/19q-noncodeleted molecular subtype: $P = .021$; *IDH*-mutant 1p/19q-codeleted molecular subtype versus *IDH* wild-type molecular subtype: $P < .001$; and *IDH*-mutant 1p/19q-noncodeleted molecular subtype versus *IDH*-wild type molecular subtype: $P < .001$). Survival curves for patients who composed a subset of the current patient cohort are shown in Patel et al.,⁴⁶ in 2019.

Statistical Analysis

Data Summarization. Categorical variables are summarized by frequencies and percentages, and continuous scaled data are summarized by the median, the interquartile range (IQR), and the range of the empiric distribution.

Survival Analyses. Cox proportional hazards regression was used to examine the prognostic utility of the preoperative MR imaging metrics to predict survival within the currently defined molecular categories of LGGs. The survival analyses were conducted per molecular category (ie, *IDH*mut-Codeled, *IDH*mut-NonCodeled, and *IDH*wt) and per MR imaging metrics. For each MR imaging metric, a multivariate Cox proportional hazards regression analysis was conducted in which the MR imaging metric of interest served as the primary predictor variable and patient age, patient sex, WHO grade, and surgical resection status served as concomitant adjustment variables. The follow-up times of survivors were treated as right-censored survival times in the Cox model. With

Table 3: Associations between preoperative MR imaging metrics and overall survival per LGG molecular subtype, after adjustment for patient age, sex, tumor grade, and surgical resection status^a

Relative Comparison (Non-Reference:Reference) ^b	<i>IDH</i> mut-Codeled	<i>IDH</i> mut-Noncodeled	<i>IDH</i> wt
Tumor size (3rd to 1st quartile)	2.85 (1.06–7.70) ($P = .039$)	2.90 (1.54–5.48) ($P < .001$)	3.82 (1.94–7.75) ($P < .001$)
Location (central:peripheral)	0.59 (0.16–2.18) ($P = .430$)	0.50 (0.22–1.17) ($P = .101$)	0.78 (0.36–1.70) ($P = .534$)
No. of involved lobes ($X + 1:X$)	1.37 (0.84–2.23) ($P = .211$)	1.25 (0.99–1.58) ($P = .065$)	1.70 (1.28–2.27) ($P = .001$)
Corpus callosum extension (yes:no)	2.60 (0.69–9.84) ($P = .160$)	1.42 (0.59–3.41) ($P = .433$)	1.89 (0.78–4.58) ($P = .160$)
Hydrocephalus (yes:no)	2.57 (0.42–15.81) ($P = .308$)	0.63 (0.19–2.11) ($P = .457$)	4.43 (1.12–17.54) ($P = .034$)
Midline shift (3rd to 1st quartile)	1.32 (0.77–2.26) ($P = .320$)	1.14 (0.54–2.41) ($P = .730$)	1.16 (1.03–1.30) ($P = .013$)
Eloquent cortex (yes:no)	1.00 (0.29–3.43) ($P = .740$)	1.91 (0.77–4.78) ($P = .165$)	2.01 (0.90–4.49) ($P = .087$)
Ependymal extension (yes:no)	6.34 (1.07–37.59) ($P = .042$)	1.72 (0.64–4.64) ($P = .289$)	1.51 (0.52–4.35) ($P = .447$)
Margins (global test P value) ^c	($P = .775$)	($P = .190$)	($P = .031$)
33%–66%:<33% ^d	1.57 (0.42–5.97) ($P = .505$)	0.51 (0.19–1.33) ($P = .166$)	0.39 (0.17–0.90) ($P = .027$)
>66%:<33% ^d	0.98 (0.09–10.85) ($P = .985$)	0.39 (0.13–1.19) ($P = .098$)	0.24 (0.06–1.05) ($P = .057$)
>66%:33%–66% ^d	0.62 (0.06–6.40) ($P = .688$)	0.77 (0.24–2.38) ($P = .646$)	0.63 (0.14–2.87) ($P = .554$)
Contrast enhancement (yes:no)	3.11 (0.71–13.68) ($P = .132$)	1.67 (0.62–4.51) ($P = .313$)	0.34 (0.13–0.90) ($P = .030$)
Necrosis (yes:no)	1.84 (0.32–10.78) ($P = .498$)	0.28 (0.03–2.42) ($P = .247$)	1.93 (0.82–4.58) ($P = .134$)

^a Data are listed as adjusted hazard ratios (95% confidence interval) and corresponding P values.

^b Relative comparison (nonreference:reference) identifies the nonreference predictor variable value/level and the reference predictor variable value/level at which the adjusted instantaneous risk of death ratio (ie, hazard ratio) was evaluated. For example (3rd:1st quartile) represents a comparison of the adjusted instantaneous risk of death (ie, hazard) between 2 patients, 1 patient whose predictor variable value is at the 3rd quartile of the predictor variable empiric distribution (nonreference) and 1 patient whose predictor variable value is at the 1st quartile of the predictor variable empiric distribution (reference). A relative comparison denoted as ($X + 1:1$) represents a comparison of the adjusted instantaneous risk of death (ie, hazard) between 2 patients, 1 patient whose predictor variable value is $X + 1$ units (nonreference) and 1 patient whose predictor variable value is X units (reference). A relative comparison, denoted as (yes:no), represents a comparison of the adjusted instantaneous risk of death (ie, hazard) between 2 patients, 1 patient who has the factor of interest (nonreference = yes) and 1 patient who does not have the factor of interest (reference = no). Note, if the adjusted hazard ratio is >1 (<1), it indicates that the estimate for the instantaneous risk of death is greater (less) for the patient who has the nonreference predictor value/level.

^c Global test P value is the P value for testing the null hypothesis that the instantaneous risk of death is same for all “margin” categories.

^d Percentage of LGGs displaying sharp/circumscribed margin.

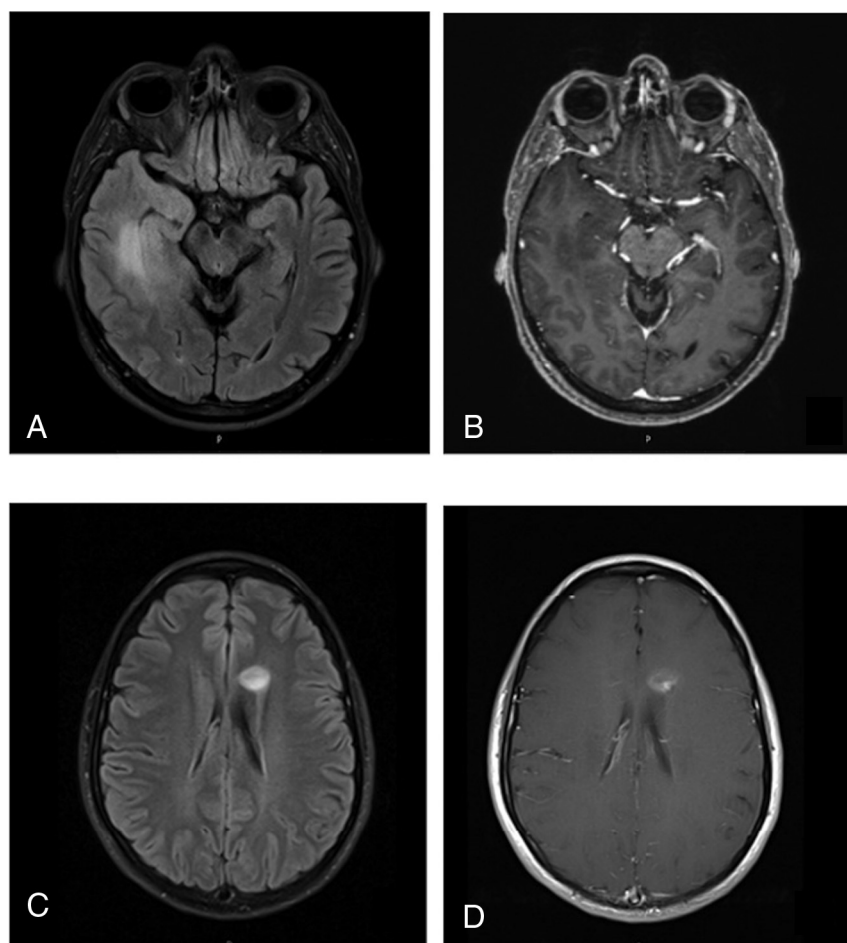


FIG 2. Representative cases of *IDHwt* LGGs. A 51-year-old man with an *IDH* wild-type diffuse astrocytoma. FLAIR (A) shows a 6-cm mass in the right temporal lobe with ill-defined margins. Contrast-enhanced T1WI (B) shows no contrast enhancement of the mass. This patient is deceased, with a survival time of 262 days. Further molecular testing in this case was positive for *TERT* promoter (–124 C > T) mutation. A 26-year-old woman with an *IDH* wild-type diffuse astrocytoma. FLAIR (C) and contrast-enhanced T1WI (D) show a 1.6-cm left frontal lobe mass with fairly well-circumscribed margins and contrast enhancement. This patient was alive at last follow-up, with a survival time of 2757 days. Further molecular testing in this case was positive for the *BRAF* V600E mutation.

regard to hypothesis testing, the Wald χ^2 test was used to test the null hypothesis that the instantaneous risk of death (ie, hazard) is not associated with the values/categories of the MR imaging metric of interest after adjustment for patient age, patient sex, WHO grade, and surgical resection status. A $P \leq .05$ decision rule was used as the null hypothesis rejection criterion for the tests of association.

Tumor Grade Analyses. MR imaging metrics versus WHO grade association were examined via multivariate logistic regression, per LGG molecular subtype. The dependent variable of the multivariate logistic regression model was an indicator variable (Y) that distinguished WHO grade III tumors ($Y = 1$) from WHO grade II tumors ($Y = 0$), and the MR imaging metrics served as the predictor variables. Regarding hypothesis testing, the type III Wald χ^2 test was used to test the null hypothesis that the adjusted odds for a tumor being WHO grade III are not associated with

the MR imaging metric. A $P \leq .05$ decision rule was used as the null hypothesis rejection criterion.

RESULTS

The study population included 306 patients, with 154 women (50.3%) and 152 men (49.7%). Patient characteristics per LGG molecular subtype are shown in Table 1, and preoperative MR imaging metric empiric distribution summaries per LGG molecular subtype are shown in Table 2.

Survival curves for overall survival (OS) are shown in Fig 1 for the LGG molecular subtypes. OS differed among all the LGG molecular subtypes ($P < .022$ for all comparisons), with patients with *IDHwt* LGGs having the shortest median OS (49.5 months; 95% CI lower bound, 25.5 months), followed by the patients with *IDHmut-Noncodel* (134.6 months; 95% CI lower bound, 98.7 months), and patients with *IDHmut-Codel* (196.6 months, 95% CI lower bound, 141.4 months).

The associations between preoperative MR imaging metrics and OS per LGG molecular subtype are expressed as hazard ratios (HRs) in Table 3, after adjustment for patient age, patient sex, tumor grade, and surgical resection status. For *IDHwt* LGGs, greater tumor size (HR, 3.82; 95% CI, 1.94–7.75; $P < .001$), greater number of involved lobes (HR, 1.70; 95% CI, 1.28–2.27; $P < .001$), hydrocephalus (HR, 4.43; 95% CI, 1.12–17.54; $P = .034$), greater degree of midline shift (HR, 1.16; 95% CI, 1.03–1.30; $P = .013$), and less circumscribed margins ($P = .031$) were associated with shorter OS,

whereas the presence of contrast enhancement (HR, 0.34; 95% CI, 0.13–0.90; $P = .030$) was associated with longer OS (Fig 2). For *IDHmut-Noncodel* LGGs, greater tumor size (HR, 2.90; 95% CI, 1.54–5.48; $P = .001$) was associated with shorter OS. For *IDHmut-Codel* LGGs, greater tumor size (HR, 2.85; 95% CI, 1.06–7.70; $P = .039$) and ependymal extension (HR, 6.34; 95% CI, 1.07–37.59; $P = .042$) were associated with shorter OS.

A secondary analysis determined unique associations between MR imaging metrics and tumor grade (II or III) per LGG molecular subtype (Table 4). Contrast enhancement was associated with grade III for both *IDHwt* LGGs (OR, 16.75; 95% CI, 3.47–80.86; $P < .001$) and *IDHmut-Noncodel* LGGs (OR, 6.08; 95% CI, 2.12–17.41; $P = .001$), but not for *IDHmut-Codel* LGGs. For *IDHwt* LGGs, ependymal extension was associated with grade II (OR, 0.09; 95% CI, 0.01–0.59; $P = .012$), and for *IDHmut-Noncodel* LGGs, central

Table 4: Associations between preoperative MR imaging metrics and WHO grade III per molecular subtype^a

Relative Comparison (Non-Reference:Reference) ^b	IDHmut-Codel	IDHmut-Noncodel	IDHwt
Tumor size (3rd to 1st quartile)	0.97 (0.18–5.26) (<i>P</i> = .975)	1.24 (0.34–4.57) (<i>P</i> = .742)	1.60 (0.31–8.17) (<i>P</i> = .573)
Location (central:peripheral)	1.55 (0.22–10.49) (<i>P</i> = .654)	0.21 (0.05–0.80) (<i>P</i> = .035)	0.42 (0.10–1.84) (<i>P</i> = .252)
No. of involved lobes (X + 1:X)	1.45 (0.64–3.30) (<i>P</i> = .370)	1.12 (0.65–1.94) (<i>P</i> = .673)	0.96 (0.48–1.92) (<i>P</i> = .906)
Corpus callosum extension (yes:no)	1.88 (0.39–9.05) (<i>P</i> = .430)	2.51 (0.72–8.76) (<i>P</i> = .149)	1.46 (0.25–8.35) (<i>P</i> = .672)
Hydrocephalus (yes:no)	8.55 (0.33–223.32) (<i>P</i> = .197)	2.07 (0.50–8.54) (<i>P</i> = .313)	0.26 (0.01–5.09) (<i>P</i> = .372)
Midline shift (3rd to 1st quartile)	0.88 (0.37–2.10) (<i>P</i> = .774)	0.55 (0.16–1.85) (<i>P</i> = .331)	1.31 (0.91–1.89) (<i>P</i> = .148)
Eloquent cortex (yes:no)	1.71 (0.49–5.99) (<i>P</i> = .402)	1.87 (0.69–5.09) (<i>P</i> = .220)	0.35 (0.08–1.54) (<i>P</i> = .165)
Ependymal extension (yes:no)	1.68 (0.28–10.07) (<i>P</i> = .570)	0.65 (0.15–2.86) (<i>P</i> = .571)	0.09 (0.01–0.59) (<i>P</i> = .012)
Margins (global test) ^c	<i>P</i> = .970	<i>P</i> = .360	<i>P</i> = .221
33%–66%:<33% ^d	1.20 (0.28–5.23) (<i>P</i> = .804)	1.47 (0.37–5.79) (<i>P</i> = .583)	1.62 (0.24–10.84) (<i>P</i> = .619)
>66%:<33% ^d	1.16 (0.15–9.17) (<i>P</i> = .888)	0.64 (0.12–3.56) (<i>P</i> = .614)	0.27 (0.02–3.35) (<i>P</i> = .305)
>66%:33%–66% ^d	0.96 (0.17–5.56) (<i>P</i> = .967)	0.44 (0.13–1.42) (<i>P</i> = .170)	0.16 (0.02–1.28) (<i>P</i> = .084)
Contrast enhancement (yes:no)	1.73 (0.47–6.43) (<i>P</i> = .411)	6.08 (2.12–17.41) (<i>P</i> = .001)	16.75 (3.47–80.86) (<i>P</i> < .001)
Necrosis (yes:no)	9.68 (0.83–112.49) (<i>P</i> = .070)	0.61 (0.08–4.67) (<i>P</i> = .632)	0.67 (0.13–3.36) (<i>P</i> = .630)

^a Note that for the multivariate logistic regression analysis, WHO grade III tumors were assigned the value 1 and WHO grade II tumors were assigned the value 0. Data are listed as adjusted odds ratios, (95% confidence interval) and corresponding *P* value determined by a Wald type III χ^2 test.

^b Relative comparison (nonreference:reference) identifies the nonreference predictor variable value/level and the reference predictor variable value/level at which the adjusted odds ratio was evaluated. For example (3rd:1st Quartile) represents a comparison of the adjusted odds for tumor grade III between 2 patients, 1 patient whose predictor variable value is at the 3rd quartile of the predictor variable empiric distribution (nonreference) and 1 patient whose predictor variable value is at the 1st quartile of the predictor variable empiric distribution (reference). A relative comparison denoted as (X + 1:1) represents a comparison of the adjusted odds for tumor grade III between 2 patients, 1 patient whose predictor variable value equals X + 1 (nonreference) and 1 patient whose predictor variable value equals X (reference). A relative comparison denoted as (yes:no) represents a comparison of the adjusted odds for a tumor grade III between 2 patients, 1 patient who has the factor of interest (nonreference = yes) and 1 patient who does not have the factor of interest (reference = no). Note, if the adjusted odds ratio is >1 (<1), it indicates that the estimate for the adjusted odds for a tumor grade III is greater (less) for the patient who has the nonreference predictor value/level.

^c Global test *P* value is the *P* value for testing the null hypothesis that the adjusted odds ratio is same for all “margin” categories.

^d Percentage of LGG displaying sharp/circumscribed margin.

location was associated with grade II (OR, 0.21; 95% CI, 0.05–0.80; *P* = .035).

DISCUSSION

Despite the markedly improved prognostic relevance of the 2016 WHO classification of LGGs, clinical outcome variability persists for LGGs within existing molecular subtypes.^{26–29} For each LGG molecular subtype, we separately explored the association between preoperative MR imaging metrics and patient OS after adjustment for WHO grade, surgical resection status, patient age, and patient sex. We further explored the relationship between MR imaging metrics and tumor grade separately for each LGG subtype. Among our results, we found that preoperative MR imaging features have more prognostic value for IDHwt LGGs than for IDH-mutant LGGs. Tumor size is significantly associated with OS in all LGG molecular subtypes, and MR imaging associations with tumor grade vary among LGG molecular subtypes.

Our finding that preoperative MR imaging has the greatest prognostic value for the IDHwt subtype is potentially explained by the increasingly well-recognized biologic and clinical heterogeneity of IDHwt LGGs, in particular based on molecular alterations such as telomerase reverse transcriptase (*TERT*) promoter mutation (Fig 2A, -B), epidermal growth factor receptor (*EGFR*) gene amplification, and chromosome 7/10 alterations.^{12,26,27,30} These molecular alterations have not yet been formally incorporated into the WHO classification scheme, and it is conceivable that some of our neuroimaging metrics correlated with these prognostically significant molecular alterations among our IDHwt cohort. We further speculate that our apparently counterintuitive finding of a positive correlation between contrast enhancement and survival time for the IDHwt cohort could be explained by the potential inclusion of *B-raf proto-oncogene*

(*BRAF*)-mutant IDHwt gliomas in our cohort (Fig 2C, -D). *BRAF*-mutant diffuse gliomas commonly demonstrate contrast enhancement and are associated with favorable clinical outcomes.^{31–33}

On the other hand, we found relatively few significant associations with overall survival for the IDH-mutant LGG subtypes. These results might reflect that the IDH-mutant LGG subtypes represent more uniform genomic entities compared with the IDHwt subtype. Nonetheless, for both IDHmut-Noncodel and IDHmut-Codel subtypes, larger tumor size was associated with worse OS. Moreover, we found that ependymal extension is associated with worse OS for the IDHmut-Codel subtype. It is known that subventricular zone extension by glioblastoma portends a worse prognosis, possibly due to recruitment of migratory progenitor cells in this location.³⁴ It is unclear whether a similar process contributes to a more aggressive clinical course among the IDHmut-Codel LGGs, and further investigation into the impact of ependymal extension in the IDHmut-Codel subtype could build on our results.

To our knowledge, the literature investigating prognostically relevant imaging metrics in molecularly defined LGG subtypes is sparse. Perhaps the most applicable study is by Wu et al,³⁵ in 2019, who found that tumor size and elevated relative cerebral blood volume on preoperative imaging correlate with a more aggressive subtype of IDHmut-Noncodel gliomas. While our results support these findings in regard to tumor size, we observed a similar relationship for IDHmut-Codel and IDHwt subgroups as well. Small sample size and lack of adjustment for WHO grade by Wu et al are limitations that may have contributed to differences observed between our studies. Suchorska et al,³⁶ in 2019, found that contrast enhancement is a viable prognostic metric for IDH-mutant subgroups, a finding that was not reproduced in our study. Finally, the apparent diffusion

coefficient has been reported as a potential prognostic marker for *IDHwt* LGGs.³⁷

Multiple prior studies have investigated imaging findings that correlate with WHO grade.^{38–42} However, to the best of our knowledge, we are the first to report the relationship between neuroimaging metrics and WHO grade separately for each LGG molecular subtype. Our results indicate that contrast enhancement predicts WHO grade III for the *IDHmut*-Noncodel and *IDHwt* LGGs, but not for *IDHmut*-Codel LGGs. A previous study of oligodendrogliomas by White et al,⁴³ in 2005, also reported no association between contrast enhancement and WHO grade; however, 1p/19q testing was not undertaken in their study, and their cohort was small ($n=24$). Our results strengthen this conclusion and may serve to modify the well-established dictum that contrast enhancement correlates with tumor grade in adult diffuse gliomas.^{44,45}

While efforts to identify preoperative neuroimaging predictors of LGG molecular status are valuable, the clinical utility is frequently limited because nearly all such cases undergo biopsy or resection for a definitive pathologic diagnosis. Our investigation aimed to uncover simple neuroimaging metrics that could relay prognostic information for LGGs beyond what can be inferred from their molecular and histologic characterization. An added strength of our study design was the adjustment of our analysis for both tumor grade and surgical resection status. The extent of surgical resection in particular is well-recognized for its impact on the overall survival of a patient with LGG.^{46,47}

Nonetheless, our study has limitations. This is a retrospective study and thus inherently limited by design in its ability to infer relationships beyond association. Further prospective multi-institutional investigations would be necessary to confirm our results. Another potential limitation is the binary method of analysis for several metrics (eg, yes or no contrast enhancement). While this method was chosen to reduce ambiguity in the data acquisition and maintain a simple set of imaging metrics, a more graded approach to measurement may allow a more nuanced and potentially revealing analysis. Similarly, we measured “tumor size” as a single long-axis diameter, and volumetric measurements based on 3D MR imaging acquisitions would be preferable. Furthermore, we investigated a limited set of MR imaging pulse sequences (pre-contrast T1WI, contrast-enhanced T1WI, and either T2WI or FLAIR), and additional study into the prognostic value of more advanced techniques (eg, perfusion-weighted imaging, diffusion tensor imaging, susceptibility-weighted imaging) might be fruitful. We did not adjust our analysis for nonsurgical treatment (eg, chemotherapy and radiation therapy), which presumably varied across our cohort. Finally, future investigations evaluating links between MR imaging metrics and molecular alterations pertinent to *IDHwt* LGGs, such as *BRAF* or *TERT* mutations, would be highly valuable.

CONCLUSIONS

Preoperative MR imaging metrics offer prognostic information for patients with LGG within molecularly defined classes. Multiple neuroimaging features had unique prognostic significance for the *IDHwt* subtype, including hydrocephalus, midline

shift, margin features, and contrast enhancement, likely reflecting the known biologic and clinical heterogeneity of *IDHwt* LGGs. Contrast enhancement was associated with WHO grade III among *IDHwt* and *IDHmut*-Noncodel LGGs, but not *IDHmut*-Codel LGGs.

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