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Patterns of Tumor Contrast Enhancement Predict the Prognosis of Anaplastic Gliomas with *IDH1* Mutation

Y.Y. Wang, K. Wang, S.W. Li, J.F. Wang, J. Ma, T. Jiang, and J.P. Dai



ABSTRACT

BACKGROUND AND PURPOSE: It is proposed that *isocitrate dehydrogenase 1 (IDH1)* mutation predicts the outcome in patients with high-grade glioma. In addition, contrast enhancement on preoperative MR imaging reflects tumor biologic features. Patients with anaplastic glioma with the *IDH1* mutation were evaluated by using MR imaging to determine whether tumor enhancement is a prognostic factor and can be used to predict survival.

MATERIALS AND METHODS: A cohort of 216 patients with histologically confirmed anaplastic glioma was reviewed retrospectively. Tumor contrast-enhancement patterns were classified on the basis of preoperative T1 contrast MR images. Tumor *IDH1* status was examined by using RNA sequencing. We used univariate analysis and the multivariate Cox model to evaluate the prognostic value of the *IDH1* mutation and tumor contrast-enhancement pattern for progression-free survival and overall survival.

RESULTS: In all 216 patients, *IDH1* mutation was associated with longer progression-free survival ($P = .004$, hazard ratio = 0.439) and overall survival ($P = .002$, hazard ratio = 0.406). For patients with *IDH1* mutant anaplastic glioma, the absence of contrast enhancement was associated with longer progression-free survival ($P = .038$, hazard ratio = 0.473) and overall survival ($P = .043$, hazard ratio = 0.436). Furthermore, we were able to stratify the progression-free survival and overall survival of patients with *IDH1* mutation by using the tumor contrast-enhancement patterns ($P = .022$ and 0.029, respectively; log-rank).

CONCLUSIONS: Tumor enhancement on postcontrast MR imaging is a valuable prognostic factor for patients with anaplastic glioma and *IDH1* mutation. Furthermore, the contrast-enhancement patterns could potentially be used to stratify the survival outcome of such patients.

ABBREVIATIONS: AG = anaplastic glioma; GTR = gross total resection; <GTR = residual tumor; HR = hazard ratio; *IDH1* = *isocitrate dehydrogenase 1*; KPS = Karnofsky Performance Status Scale; PFS = progression-free survival; OS = overall survival

Anaplastic gliomas (AGs), classified as World Health Organization grade III, are aggressive brain tumors. They exhibit morphometric heterogeneity on radiologic imaging, and their clinical course varies substantially so that some patients succumb

to progressive disease within weeks while others survive for a decade or more. The clinical characteristics, radiologic features, genetic changes, and extent of resection all play important roles in determining the prognosis of patients with AG.^{1–9} Interaction and synergy may exist among these factors.

The presence of contrast enhancement on MR images, which is based on pathophysiologic changes indicating the destruction of the blood-brain barrier, is considered a specific radiologic feature of high-grade gliomas. Previous studies have revealed the prognostic role of contrast enhancement in patients with AG.^{5,8,10–12} Additionally, radiologic features such as enhancement and multifocality correlate with the molecular characteristics of malignant glioma.¹³

Mutation in the *isocitrate dehydrogenase 1 (IDH1)* gene at R132 is an important molecular event and plays a significant role in gliomagenesis. This genetic change is detected in approximately 50%–70% of anaplastic astrocytomas^{1,14,15} and in 70% of anaplastic oligodendrogliomas.^{14–16} Furthermore, the presence of

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IDH1 mutations distinguishes tumors with markedly different clinical presentations, concurrent molecular genetic alterations, and overall natural history.¹ For patients with AG, the occurrence of *IDH1* mutation is associated with gross total resection (GTR) and longer overall survival (OS).¹⁷

Previous studies have demonstrated the correlation between *IDH1* status and the radiologic features of glioma, in that tumors with *IDH1* mutation are more likely to be larger, and cyst, edema, and contrast enhancement are present.¹⁸ However, investigations of the interactive and synergistic role of *IDH1* mutation and tumor contrast enhancement in predicting the survival of patients with AG are rare. By classifying tumor contrast enhancement according to the patterns observed, patients with anaplastic glioma with *IDH1* mutation were evaluated by using MR imaging to determine whether tumor enhancement is a prognostic factor and can be used to predict outcome survival stratification.

MATERIALS AND METHODS

Patients

In total, 216 adult patients diagnosed with AG who had undergone surgical treatment at our institution from February 2007 to June 2010 were reviewed retrospectively. Patients were included on the basis of the following criteria: 1) age 18 years or older, 2) presurgical structural MR imaging scan available (T1-weighted, T2-weighted, postcontrast T1-weighted), 3) pathology-confirmed AG based on the modified World Health Organization grading system, and 4) no previous diagnosis of any brain tumor. The histopathologic diagnosis was evaluated and confirmed by 2 independent neuropathologists blinded to the patients' clinical and radiologic information. According to the Response Assessment in Neuro-Oncology criteria,¹⁹ GTR was defined as no visible contrast enhancement on postoperative MR images within 72 hours after the operation in contrast-enhanced tumors or absence of all abnormal hyperintense changes on preoperative MR images for tumors not demonstrating contrast enhancement. In this study, resections that were not GTR were considered residual tumor (<GTR). The overall follow-up duration was 85 months, which spanned March 2007 to April 2014. This study was approved by our institutional review board, and written consent was obtained from all enrolled patients.

Imaging Acquisition

MR imaging was performed on Trio 3T scanners (Siemens, Erlangen, Germany). It typically included axial T1-weighted (TR, 450 ms; TE, 15 ms; section thickness, 5 mm), T2-weighted fast spin-echo (TR, 6000 ms; TE, 140 ms; section thickness, 5 mm), and Gd-DTPA injection-enhanced (Beijing Beilu Pharmaceutical, Beijing, China; 0.1 mmol/kg) axial T1-weighted images (TR, 450 ms; TE, 15 ms; section thickness, 5 mm), with a 24-cm FOV and 256 × 256 matrix size. Postcontrast images were acquired immediately following injection of the contrast agent. The interval between contrast injection and the beginning of the contrast-enhanced T1-weighted image acquisition was maintained between 75 and 85 seconds. Postoperative MR images for determining the extent of resection were obtained within 72 hours after resection, and the radiologic parameters were maintained in accordance with the preoperative scans.

Identification of Imaging Features

Tumor contrast enhancement was assessed by 2 experienced neuroradiologists (Q.C. and X.C., who have 14 and 12 years of experience, respectively, in diagnosis using brain MR imaging) blinded to the patient clinical information. In cases in which the types of enhancement identified by the first 2 neuroradiologists were inconsistent, a third senior neuroradiologist (J.M., 25 years of experience in brain disease diagnosis) re-examined the images and determined the image to be used. "Contrast enhancement" was defined as newly emerged unequivocal increased signal intensity on the T1-weighted image following intravenous contrast administration compared with noncontrast T1 images. "Nonenhancement" was defined as no apparent hyperintensity on postcontrast T1-weighted images. Three contrast-enhancement patterns were identified on the basis of the size and morphologic features of the largest enhanced area on contrast-enhanced MR images regardless of whether it was single or multifocal: nodular, with the largest focal diameter of ≤1.5 cm; patchy, tumors with a maximum diameter of enhancement of >1.5 cm; and ringlike, cystic necrosis with peripheral enhancement (Fig 1). Multifocal tumor enhancement was defined as >1 area of tumor enhancement separated from the others on the postcontrast T1-weighted image.

DNA Sequencing for IDH1 Mutation

IDH1 mutation was determined by using DNA pyrosequencing, which we have described previously.²⁰ Briefly, a QIAamp DNA Mini Kit (QIAGEN, Hilden, Germany) was used to isolate genomic DNA from frozen tumor tissue samples. We then analyzed the genomic region spanning the wild type R132 of *IDH1* by using pyrophosphate sequencing with 5'-GCTTGTGAGTGGATGGGTAAAAC-3' and 5'-biotin-TTGCCAA-CATGACTTACTTGATC-3' primers. Duplicate polymerase chain reaction analyses were performed in 40-μL reaction volumes containing 1-μL of each 10-μmol/L forward and reverse primer, 4 μL of 10× buffer, 3.21 μL of 2.5-mmol/L deoxynucleotide triphosphates, 2.5-U HotStar Taq (Takara, Shiga, Japan), and 2 μL of 10 μmol/L DNA. The polymerase chain reaction conditions were as follows: 95°C for 3 minutes; 50 cycles of 95°C for 15 seconds, 56°C for 20 seconds, 72°C for 30 seconds; and 72°C for 5 minutes (Applied Biosystems GeneAmp PCR System 9700; Applied Biosystems, Foster City, California). Single-stranded DNA was purified from the polymerase chain reaction products and pyrosequenced with a PyroMark Q96 ID System (QIAGEN) by using a 5'-TGGATGGGTAAAACCT-3' primer and an EpiTect Bisulfite Kit (QIAGEN).

Statistical Analysis

We used the χ^2 test for categorical variables to compare each clinical and imaging feature between the *IDH1* mutant and wild type groups. The agreement between judgments of the enhancement patterns assessed by the 2 radiologists was evaluated by using the κ consistency test. A κ value of ≥0.81, 0.61–0.80, and ≤0.60 was considered excellent, good, and poor agreement, respectively. Additionally, log-rank analysis of Kaplan-Meier data was performed to compare the progression-free survival (PFS) and OS of the cohort. Factors that were significant ($P < .05$) in univariate anal-

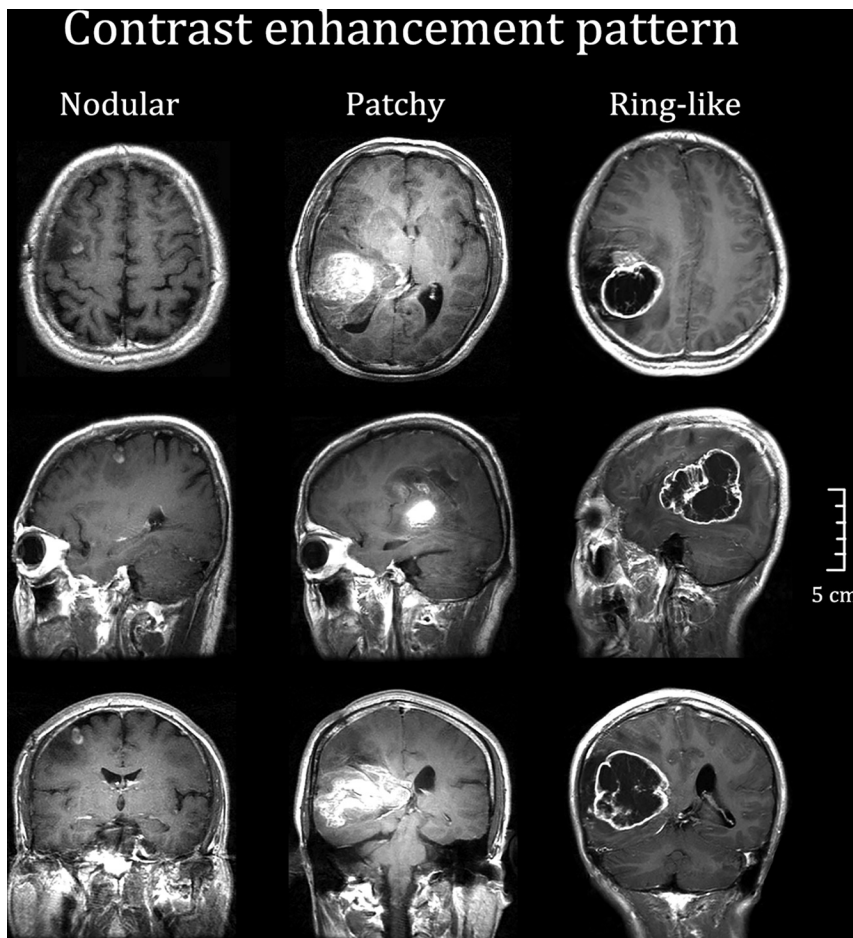


FIG 1. Tumor contrast-enhancement patterns in AG. Postcontrast T1-weighted images depict the nodular (largest focal diameter of ≤ 1.5 cm), patchy (largest focal diameter of > 1.5 cm), and ringlike (cystic necrosis with peripheral enhancement) enhancement patterns.

Table 1: IDH1 mutation status of patients with AG

Characteristics	IDH1 Status			P Value ^a
	Total (n = 216)	Mutant (n = 84)	Wild Type (n = 132)	
Age (yr)				
Median (range)	44 (18–87)	43 (18–71)	45 (18–87)	
50 or older/50 or younger	77:139	23:61	54:78	.043
Sex				
Male/female	135:81	49:35	86:46	.313
KPS				
≥ 80 / < 80	181:35	78:6	103:29	.004
Contrast enhancement				
Yes/no	173:43	57:27	116:16	$< .001$
Pattern of enhancement				
Nodular/patchy/ringlike	26:62:85	9:14:34	17:48:51	.084
Extent of resection				
GTR/ $< \text{GTR}$	123:93	56:28	67:65	.021
Histopathology				
AA/AO/AOA	57:44:115	16:20:48	41:24:67	.135

Note:—AA indicates anaplastic astrocytomas; AO, anaplastic oligodendrogliomas; AOA, anaplastic oligoastrocytomas.

^a Results of the χ^2 test.

ysis were entered into multivariate survival analysis on the basis of the Cox proportional hazard ratio (HR) model. To identify the prognostic value of *IDH1* status and tumor contrast-enhancement pattern in patients according to their interactive effects, we subdivided patients into 4 subgroups according to these 2 indicators. The respective prognostic values of the tumor contrast-en-

hancement pattern of the *IDH1* mutant and wild type groups were evaluated.

RESULTS

Patient Characteristics

The clinical and radiologic data of the 216 patients with AG were reviewed (Table 1). Among these patients, 57 (26.4%) had anaplastic astrocytoma, 44 (20.4%) had anaplastic oligodendroglioma, and 115 (53.2%) had anaplastic oligoastrocytoma. Age at diagnosis, preoperative Karnofsky Performance Status Scale (KPS), and extent of resection were significantly different between patients with mutant and wild type *IDH1* ($P < .001$, χ^2 test). A total of 123 (56.9%) patients underwent GTR, and 93 (43.1%) patients had residual tumor.

Association between IDH1

Mutation and Tumor Enhancement

There was post-T1 contrast enhancement in 173 (80.1%) tumors. Patients with *IDH1* mutation were less likely to have MR imaging tumor enhancement than patients with wild type *IDH1* (67.9% versus 87.9%, $P < .001$). In addition, tumor contrast-enhancement patterns were identified in the AGs with enhancement. The κ value for the agreement of judgment of enhancement patterns between the 2 evaluators was 0.96 ($P = .012$). Enhancement was nodular in 26 (15.0%) cases, patchy in 62 (35.9%) cases, and ringlike in 85 (49.1%) cases (Table 1). However, there was no significant difference between the proportion of contrast-enhancement patterns between tumors from patients with mutant and wild type *IDH1* ($P = .084$) (Fig 2).

Association between Surgical

Resection and Tumor Enhancement

Of the tumors with contrast enhancement, those with ringlike enhancement patterns were more likely to undergo GTR than tumors without ringlike enhancement patterns, but the difference was not statistically sig-

nificant (59.7% versus 46.9%, $P = .113$). Notably, patients with mutant *IDH1* and tumors with ringlike enhancement patterns were also more likely to undergo GTR than patients with mutant *IDH1* and tumors without ringlike enhancement patterns (65.3% versus 37.5%, $P = .004$). However, in patients with wild type *IDH1*, GTR between tumors with and without

ringlike enhancement patterns was not significantly different (47.8% versus 66.7%, $P = .157$).

Progression-Free Survival

There was tumor recurrence in 165 (76.4%) patients during the follow-up period; the median PFS was 16.9 months (range, 3.1–82.8 months). Univariate analysis showed that patients with mutant *IDH1* had significantly longer PFS than patients with wild type *IDH1* ($P = .002$, log-rank). Additionally, age at diagnosis ($P < .001$), preoperative KPS ($P = .004$), and GTR ($P = .001$) were significant prognostic factors for PFS (Table 2). In multivariate Cox regression analysis, wild type *IDH1* ($P = .004$, HR = 2.277; 95% confidence interval, 1.303–3.968), preoperative KPS < 80 ($P = .015$, HR = 2.158; 95% CI, 1.179–3.471), age at diagnosis older than 50 years ($P = .018$, HR = 1.857; 95% CI, 1.111–3.106), and $< \text{GTR}$ ($P = .028$, HR = 1.598; 95% CI, 1.053–2.597) were associated with poor PFS (Table 3).

Overall Survival

At the time of analysis, 69 patients (whose follow-up data were available) were still alive; the median follow-up period was 22.9 months (range, 3.3–86.4 months). In univariate analysis, *IDH1* status ($P = .004$), age at diagnosis ($P = .007$), preoperative KPS ($P = .002$), and extent of resection ($P < .001$) were prognostic factors of OS (Table 2). These 4 factors remained significant in the multivariate Cox proportional hazards analysis: Wild type *IDH1*

($P = .002$, HR = 2.463; 95% CI, 1.389–4.386), age at diagnosis 50 years and older ($P = .016$, HR = 1.431; 95% CI, 1.342–2.434), preoperative KPS < 80 ($P = .026$, HR = 1.836; 95% CI, 1.087–3.402), and $< \text{GTR}$ ($P = .023$, HR = 1.488; 95% CI, 1.210–2.432) were poor prognostic factors for OS (Table 3).

Prognostic Role of Tumor Contrast-Enhancement Pattern for Patients with Mutant *IDH1*

In the mutant *IDH1* group, patients with contrast-enhanced tumors had significantly shorter PFS (median PFS, 15.9 months; range, 2.6–76.8 months) than those with nonenhanced tumors (median PFS, 26.3 months; range, 2.7–80.3 months) ($P = .038$, log-rank) (Fig 3). Furthermore, the tumor contrast-enhancement patterns allowed us to stratify the PFS of patients with mutant *IDH1*. Patients with nodular enhancement patterns had significantly longer PFS (median, 23.8 months) than those with patchy (median, 17.6 months) ($P = .042$, log-rank) or ringlike enhancement patterns (median, 14.7 months) ($P = .010$, log-rank). There were no significant differences between the PFS of patients with patchy or ringlike enhancement patterns ($P = .273$, log-rank) (Fig 4). In comparison, the tumor contrast enhancement patterns had no prognostic value in patients with wild type *IDH1* ($P = .896$, log-rank).

Patients were subdivided into 4 groups according to *IDH1* status and tumor contrast enhancement. Notably, among the 4 groups, patients with mutant *IDH1* and nonenhanced tumor had significantly longer OS than patients in the other groups ($P = .043$, Fig 3). Tumor contrast-enhancement pattern played a prognostic role in the OS of patients with mutant *IDH1*. In the mutant *IDH1* group, patients with nodular enhancement patterns had significantly longer OS (median, 31.8 months) than those with patchy (median, 27.1 months) ($P = .025$, log-rank) and ringlike enhancement patterns (median, 20.3 months) ($P = .012$, log-rank). There were no significant differences between the OS of patients with patchy enhancement patterns and those with ringlike enhancement patterns ($P = .441$, log-rank) (Fig 4). In comparison, tumor contrast enhancement patterns were not a prognostic factor in patients with wild type *IDH1* ($P = .842$, log-rank).

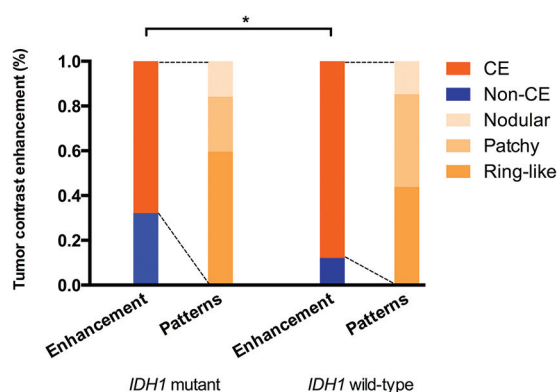


FIG 2. Constitution of tumor contrast enhancements between AG accompanied by mutant or wild type *IDH1*. The difference in contrast-enhancement rate (asterisk) between tumors from patients with mutant and wild type *IDH1* was significant ($P < .001$). There was no significant difference in enhancement-pattern distribution between tumors from patients with mutant and wild type *IDH1* ($P = .135$). CE indicates contrast enhancement.

Prognostic Role of Extent of Resection for Patients with Mutant *IDH1* and Contrast Enhancement

In the mutant *IDH1* and contrast-enhancement groups, patients with GTR had significantly longer PFS (median, 19.2 months; range, 2.1–75.8 months) than those with $< \text{GTR}$ (median, 13.0 months; range, 3.0–73.7 months) ($P = .018$, log-rank). Similarly, for patients with mutant *IDH1* and contrast-enhanced tumors, GTR also predicted significantly longer OS than $< \text{GTR}$ ($P = .030$,

Table 2: Univariate analysis of survival outcomes for patients with AG

Characteristic	PFS			OS		
	P Value	HR	95% CI	P Value	HR	95% CI
Age 50 yr or older	$< .001$	1.813	1.318–2.494	.007	1.729	1.134–2.421
Sex (male)	.215	0.873	0.695–1.064	.415	0.914	0.612–1.283
Preoperative KPS < 80	.004	2.603	1.154–3.548	.002	2.872	1.270–3.341
Enhancement	.570	1.200	0.641–2.247	.625	1.187	0.596–2.365
Enhancement pattern	.150	1.107	0.964–1.271	.247	1.902	0.941–1.266
$< \text{GTR}/\text{GTR}$.001	1.734	1.237–2.429	$< .001$	1.926	1.346–2.758
Histopathology	.086	0.855	0.715–1.022	.054	0.789	0.654–1.002
<i>IDH1</i> wild type	.002	2.364	1.362–4.098	.004	2.688	1.231–4.717

log-rank). Nevertheless, GTR did not have prognostic power for PFS and OS in the wild type *IDH1* and contrast-enhancement groups (PFS, $P = .224$; OS, $P = .141$, respectively; log-rank).

DISCUSSION

We combined clinical, radiologic, and specific genetic characteristics to investigate the prognostic factors for a large cohort of patients with AG, demonstrating that *IDH1* mutation was an independent prognostic factor for patients with AG. Furthermore, the tumor contrast-enhancement pattern identified from post-contrast MR imaging was associated with the survival outcomes of patients with mutant *IDH1*. To our knowledge, this is the first

investigation of the prognostic role of combined *IDH1* mutation and tumor contrast-enhancement pattern for predicting survival in patients with AG.

Previous studies have demonstrated the impact of *IDH1* mutation on the clinical prognosis of patients with malignant glioma.¹ In patients with AG, it has been demonstrated that *IDH1* mutation is a good prognostic marker and potential stratification factor for anaplastic astrocytoma and anaplastic oligodendroglioma.^{20,21} Consistent with these findings, the present study showed that patients with mutant *IDH1* had significantly longer PFS and OS than those with wild type *IDH1*. The better prognosis of patients with mutant *IDH1* may be partly attributed to the effect of *IDH1* interaction with other clinical characteristics. First, *IDH1* mutation is frequent in diffuse low-grade gliomas but rare in primary glioblastomas, and the survival outcome of patients with low-grade glioma is generally better than that of patients with glioblastoma.^{1,14,15,22} This variance of *IDH1* mutation incidence between low- and high-grade gliomas may contribute to the association of *IDH1* mutation with good prognosis. In addition, *IDH1* mutation is more common in younger than elder patients.²³ Because age is a widely reported significant prognostic factor,^{2,4,6,18,24} the prognosis of patients with mutant *IDH1* might be good.

In this study, the frequency of *IDH1* mutation between the 2

Table 3: Multivariate analysis of survival outcomes

Predictor	P Value ^a	HR	95% CI
PFS			
Age 50 yr or older	.018	1.857	1.111–3.106
Preoperative KPS < 80	.015	2.158	1.179–3.471
<GTR	.028	1.598	1.053–2.597
<i>IDH1</i> wild type	.004	2.277	1.303–3.968
OS			
Age 50 yr or older	.016	1.431	1.342–2.434
Preoperative KPS < 80	.026	1.836	1.087–3.402
<GTR	.023	1.488	1.210–2.432
<i>IDH1</i> wild-type	.002	2.463	1.389–4.386

^a Cox proportional hazard regression analyses. A P value of .05 denoted significance.

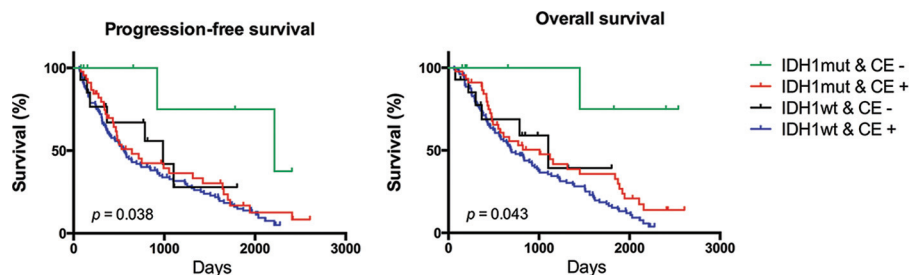


FIG 3. Kaplan-Meier plots of the series of 216 patients with AG showing the association between the PFS and OS according to combined *IDH1* status and tumor contrast enhancement. Mutant *IDH1* with no contrast enhancement predicts better survival (PFS, $P = .038$; OS, $P = .043$). mut indicates mutant; wt, wild type.

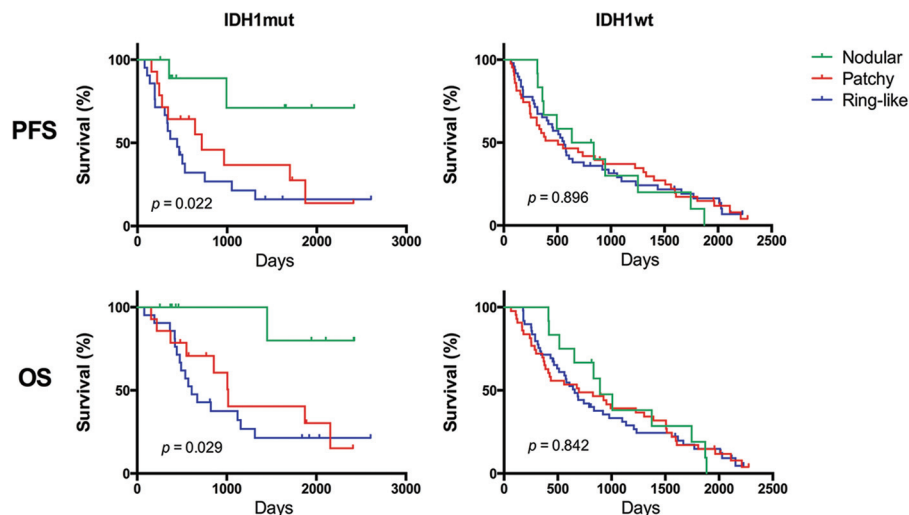


FIG 4. Kaplan-Meier plots showing that the tumor contrast enhancement pattern enabled stratification of the PFS and OS of patients with mutant *IDH1* (PFS, $P = .022$; OS, $P = .029$). Meanwhile, Kaplan-Meier plots show that the tumor contrast-enhancement pattern did not enable stratification of the PFS and OS of patients with wild type *IDH1* (PFS, $P = .896$; OS, $P = .842$).

age groups (50 years and older and younger than 50 years of age at diagnosis) was marginally significantly different ($P = .043$); these results agreed with the previous ones. Furthermore, this and a previous study²⁵ found that GTR was more likely to be achieved in AG with mutant *IDH1* than in AG with wild type *IDH1*. There was a higher rate of ringlike enhancement patterns in patients with mutant *IDH1*; in addition, these patients were more likely to undergo GTR than patients whose tumors did not have ringlike enhancement patterns, which could also have contributed to the difference in survival outcome between the 2 subgroups.

On the other hand, previous studies have suggested that the good prognosis of patients with mutant *IDH1* is primarily due to the less aggressive biologic behavior of tumors with mutant *IDH1* compared with tumors with wild type *IDH1*.^{17,19,26} Because *IDH1* mutation is considered an early genetic event in tumorigenesis and may drive other genetic changes in tumor cells, tumors accompanied by *IDH1* mutation may consequently have different genetic characteristics compared with tumors unaccompanied by the mutation, which may lead to their varied biologic features. The intrinsic difference in the tumor biologic features may explain why the *IDH1* mutation, though associated with other clinical characteristics, was an independent prognostic factor for patients with AG. Other than *IDH1* mutation, however, the histopathologic subtypes in the present study did not have prognostic value, indicating that the variety of tumor components could not predict survival for patients with AG.

It has been reported that the radiologic features of glioma are associated with *IDH1* mutation. A recent study showed that *IDH1* mutation status in glioblastoma can be predicted from the radiologic features derived from MR images.¹⁸ The study identified 4 subjective tumor characteristics observable on MR images: tumor size, contrast enhancement, and the presence or absence of cyst and satellite lesions, which were associated with *IDH1* status; these characteristics predicted the presence of *IDH1* mutation with 94% accuracy (by receiver operating characteristic analysis). Another study demonstrated that *IDH1*-mutated gliomas were predominantly located in a single lobe and were more likely to have a unilateral growth pattern, sharp tumor margin, homogeneous signal intensity, and less contrast enhancement on MR imaging.²⁷ In the present study, we also found that tumor contrast enhancement was associated with *IDH1* status. Tumors accompanied by mutant *IDH1* were less likely to show contrast enhancement on MR images compared with tumors without *IDH1* mutation. Most interesting, although there was no significant difference in the contrast-enhancement patterns between tumors accompanied by mutant or wild type *IDH1*, multivariate Cox analysis identified the tumor contrast-enhancement pattern as an independent prognostic factor in patients with mutant *IDH1*. This result implies that the tumor contrast-enhancement pattern may be a particularly important factor reflecting the biologic features of tumors in the presence of *IDH1* mutation.

The prognostic value of *IDH1* status and tumor contrast enhancement was determined in patients with AG, considering their interactive effects. Notably, among the 4 classifications based on the 2 indicators, patients with mutant *IDH1* and nonenhanced tumor had significantly longer PFS and OS (Fig 3) than patients who did not; there was no difference in survival time among the

other 3 groups. This indicates that *IDH1* mutation and the absence of contrast enhancement may have synergistic effects in reflecting tumor malignancy and predicting survival outcome.

Most interesting, tumor contrast-enhancement patterns were identified as a prognostic marker that could be used to stratify PFS and OS only for patients with mutant *IDH1*, but not for patients with wild type *IDH1*. Specifically, patients with mutant *IDH1* with nodular enhancement patterns had longer PFS and OS than those with mutant *IDH1* with patchy or ringlike enhancement patterns ($P = .022$). Tumor contrast enhancement reflects the degree of destruction of the blood-brain barrier, which is induced by tumor cell invasion. Therefore, the tumor contrast-enhancement pattern may be strongly associated with the biologic features of a tumor. A small area of tumor enhancement (nodular pattern) possibly indicates lower grade malignancy compared with a relatively large area of tumor enhancement (patchy or ringlike pattern), which explains the present findings. Why the tumor contrast-enhancement pattern plays a more important role in predicting the survival of patients with AG with mutant *IDH1* compared with those with wild-type *IDH1* remains to be investigated.

In addition, the extent of resection in the enhanced tumors could be used to stratify PFS and OS for patients with mutant *IDH1*—that is, patients with GTR had longer PFS and OS than patients with <GTR; but this finding was not true for patients with wild type *IDH1*. As discussed above, there was a higher rate of ringlike enhancement pattern in the tumors of patients with mutant *IDH1*. Compared with tumors with nodular or patchy enhancement patterns, tumors with ringlike enhancement patterns had a relatively clear border on postcontrast T1-weighted images, which might facilitate surgical resection of the tumor bulk. On the other hand, *IDH1* mutation is common in diffuse low-grade gliomas and is more common in younger patients; these findings indicate less aggressive behavior and contribute to effective resection of most of the tumor in patients with mutant *IDH1*.

Our study has some limitations. First, we retrospectively enrolled patients from a single institution; therefore, the prognostic role of tumor contrast-enhancement patterns and *IDH1* mutation requires confirmation by a prospective multicenter investigation. Second, due to the limited number of patients and the similar distribution of *IDH1* mutation in the 3 anaplastic glioma subtypes, we did not separate patients by pathology for further discussion. Third, due to the relatively suboptimal timing of the postoperative scans, the potential presence of granulation tissue is a confounder in the MR imaging. The interval from contrast agent administration to image acquisition could also have influenced the level of enhancement. Although the study was carefully controlled, a slight discrepancy in the interval between contrast injection and scanning may still have been present among individuals. Future studies should investigate the association between the radiologic characteristics and survival of patients with tumors with other gene mutations.

CONCLUSIONS

We retrospectively reviewed 216 patients with AG and identified *IDH1* mutation as a significant prognostic factor. In this study, we found that the tumor contrast-enhancement patterns were asso-

ciated with the survival outcome of patients with mutant *IDH1*. Our results imply that there may be a synergistic effect between radiologic morphology and the genetic features of a tumor in determining prognosis, and this effect should be considered in future investigations.

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