Mitotic Activity in Glioblastoma Correlates with Estimated Extravascular Extracellular Space Derived from Dynamic Contrast-Enhanced MR Imaging

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ABSTRACT

BACKGROUND AND PURPOSE: A number of parameters derived from dynamic contrast-enhanced MR imaging and separate histologic features have been identified as potential prognosticators in high-grade glioma. This study evaluated the relationships between dynamic contrast-enhanced MRI–derived parameters and histologic features in glioblastoma multiforme.

MATERIALS AND METHODS: Twenty-eight patients with newly presenting glioblastoma multiforme underwent preoperative imaging (conventional imaging and T1 dynamic contrast-enhanced MRI). Parametric maps of the initial area under the contrast agent concentration curve, contrast transfer coefficient, estimate of volume of the extravascular extracellular space, and estimate of blood plasma volume were generated, and the enhancing fraction was calculated. Surgical specimens were used to assess subtype and were graded (World Health Organization classification system) and were assessed for necrosis, cell density, cellular atypia, mitotic activity, and overall vascularity scores. Quantitative assessment of endothelial surface area, vascular surface area, and a vascular profile count were made by using CD34 immunostaining. The relationships between MR imaging parameters and histopathologic features were examined.

RESULTS: High values of contrast transfer coefficient were associated with the presence of frank necrosis (P < .005). High values of the estimate of volume of the extravascular extracellular space were associated with a fibrillary histologic pattern (P < .01) and with increased mitotic activity (P < .05). No relationship was found between mitotic activity and histologic pattern, suggesting that the correlation between the estimate of volume of the extravascular extracellular space and mitotic activity was independent of the histologic pattern.

CONCLUSIONS: A correlation between the estimate of volume of the extravascular extracellular space and mitotic activity is reported. Further work is warranted to establish how dynamic contrast-enhanced MRI parameters relate to more quantitative histologic measurements, including markers of proliferation and measures of vascular endothelial growth factor expression.

ABBREVIATIONS: DCE-MRI = dynamic contrast-enhanced MRI; EnF = enhancing fraction; GBM = glioblastoma multiforme; IAUC60 = initial area under the contrast agent concentration curve; Ktrans = contrast transfer coefficient; Ve = volume of the extravascular extracellular space per unit tissue volume; VEGF = vascular endothelial growth factor; Vp = volume of the blood plasma per unit tissue volume.
are commonly derived from dynamic susceptibility contrast techniques, while the volume transfer coefficient ($K_{\text{trans}}$), fractional volume of the extravascular extracellular space ($v_e$), and fractional blood plasma volume ($v_p$) can be calculated from T1-weighted DCE-MR imaging.\textsuperscript{15} In glioma, $K_{\text{trans}}, v_e, \text{EnF, CBV, and CBF}$ have been shown to relate to histologic grade and/or subtype of tumor.\textsuperscript{16-21} In addition, $K_{\text{trans}}, \text{EnF},$ and CBV have been identified as potentially grade-independent prognosticators.\textsuperscript{22-25} The number of studies examining the relationship between DCE-MR imaging parameters and more specific histopathologic features in glioma is currently small and this research predominantly focuses on vascular metrics such as blood volume and flow; however, significant correlations have been described between CBV and microvascular density,\textsuperscript{26-29} VEGF expression,\textsuperscript{27,30} cell density,\textsuperscript{31} endothelial proliferation,\textsuperscript{31} and mitotic activity\textsuperscript{31} and between CBF and endothelial hyperplasia.\textsuperscript{32}

More recent studies have focused on the potential of $v_e$ as an imaging biomarker. In glioma, it has shown value in discriminating histologic grades,\textsuperscript{33,34} but compared with other potential candidate biomarkers of the extravascular extracellular space volume (ADC), on a voxel by voxel basis, no correlation was seen between the 2 metrics, suggesting that these parameters provide independent information about extravascular extracellular space characteristics.\textsuperscript{35} Two separate studies of patients with gliomas of various histologic grades have both reported significant correlations between both $K_{\text{trans}}$ and $v_e$ and vascular and microvascular density,\textsuperscript{33,34} but neither study comments on the relationship with cellular density or mitotic activity. A high-field, 7T MR imaging study of rat xenografts showed an image-matched significant negative correlation between $v_e$ and tumor cellularity.\textsuperscript{36}

We hypothesized the following: 1) Larger, rapidly growing tumors would show higher mitotic activity and high angiogenic activity reflected by $K_{\text{trans}}$ and EnF; and 2) more proliferative tumors would have a higher cellular density and mitotic activity associated with lower values of $v_e$.

**MATERIALS AND METHODS**

**Patients**

Ethical approval was obtained, and all patients gave informed consent. All tumors were histologically confirmed as glioblastoma multiforme (GBM) according to the World Health Organization criteria.\textsuperscript{1} All imaging was preoperative, and patients received no treatment other than corticosteroids, which were administered for a minimum of 48 hours before imaging to allow stabilization of the effects of steroids on DCE-MR imaging measures.\textsuperscript{37} Patients were excluded from the study if they had a history of renal dysfunction or low estimated glomerular filtration rate ($<30 \text{ mL/min/1.73 m}^2$).

**MR Imaging Data Acquisition**

Imaging was performed on a 3T Achieva system (Philips Healthcare, Best, the Netherlands) by using a sensitivity encoding head coil. DCE-MR imaging was acquired in a sagittal oblique orientation to allow improved definition of the arterial input function free from flow-related artifacts. Three precontrast T1 fast-field echo (radiofrequency spoiled gradient-echo) series ($2^\circ, 5^\circ, 16^\circ$) were acquired for calculation of baseline T1 maps (TR, 3.5 ms; TE, 1.1 ms; section thickness, 4.2 mm; matrix, $128 \times 128$; FOV, $230 \times 230 \times 105 \text{ mm}$) in the same geometry. A dynamic, contrast-enhanced acquisition series (TR, 3.5 ms; TE, 1.1 ms; flip angle, 16°; section thickness, 4.2 mm; matrix, $128 \times 128$; FOV, $230 \times 230 \times 105 \text{ mm}$) consisting of 100 volumes with temporal spacing of approximately 3.4 seconds followed. A bolus dose of 0.1 mmol/kg (body weight) of gadolinium-based contrast agent (gadodiamide; Gd-DTPA-BMA; Omniscan; GE Healthcare, Piscataway, New Jersey) was injected at a rate of 3 mL/s, after acquisition of the fifth image volume. Pre- and postcontrast T1-weighted imaging sequences (TR, 9.3 ms; TE, 4.6 ms) were acquired in the same sagittal oblique geometry for definition of the volume of interest of the whole tumor.

**MR Imaging Data Analysis**

An experienced neuroradiologist (S.J.M.) manually defined VOIs for each tumor. The VOI corresponded to the enhancing tumor and all nonenhancing tissue contained within it on the postcontrast T1-weighted images. This technique of VOI definition has previously shown good interobserver agreement (intraclass correlation coefficient $>0.94$).\textsuperscript{38} Pharmacokinetic analysis was performed on all pixels within the VOI that showed significant enhancement. Parametric maps of $K_{\text{trans}}, v_e, v_p,$ and initial area under the contrast agent concentration curve (IAUC$_{60}$) were produced by using in-house software (MaDyM; Manchester Dynamic Modeling, Manchester, UK) and the extended Tofts and Kermode pharmacokinetic model.\textsuperscript{15} Automated arterial input functions were generated from an appropriately chosen section, which included the internal carotid artery.\textsuperscript{39} Summary statistics for each parameter were generated for enhancing tumor tissue.

For each tumor, EnF initial area under the contrast agent concentration curve (EnF$_{\text{IAUC60} > \text{a}}$) and thresholded EnF (EnF$_{\text{IAUC60} > 2.5}$) were calculated by dividing the enhancing volume (volume of voxels with IAUC$_{60} > 0 \text{ mmol/s}$ for EnF$_{\text{IAUC60} > \text{a}}$ and the volume of voxels with IAUC$_{60} > 2.5 \text{ mmol/s}$ for EnF$_{\text{IAUC60} > 2.5}$) by the total volume of the tumor VOI. The cutoff threshold of IAUC$_{60} > 2.5 \text{ mmol/s}$ for EnF$_{\text{IAUC60} > 2.5}$ was previously identified as an optimal threshold for allowing the distinction of high- from low-grade gliomas.\textsuperscript{40}

**Histopathologic Data Analysis**

Two experienced neuropathologists (D.d.P. and P.P.P.) performed the histopathologic analysis. Histologic specimens were assessed for the following: necrosis (presence or absence of frank and/or geographic necrosis), cell density (3-point grading score), cell atypia (3-point grading score), mitotic activity (number of mitotic figures seen per 10 high-power-field units), infiltrates (presence or absence of lymphocytes and/or macrophages), tumor vascular pattern (presence or absence of the following features: endothelial hypertrophy and/or hyperplasia, glomeruloid structures, granulation tissue, large-vessel density, thrombosis, sclerosed vessels), an overall vascular density score (3-point grading score), and histologic pattern (fibillar, gemistocytic, oligodendrocytes, sarcomatous, giant cells, and small cells) in conjunction with standard histopathologic subtyping and grading according to the World Health Organiza-
Vascular and histologic features were not mutually exclusive; therefore, 1 feature could be described in a given tumor specimen.

Quantitative measurement of the endothelial surface area, the vascular surface area, and the vascular profile count per square millimeter was made by using CD34 immunostaining and dedicated image-analysis software.

**Statistical Analysis**

Statistical analysis was performed by using SPSS (Version 15.0; IBM, Armonk, New York) nonparametric statistical tests. While histologic parameters produce a binary classification (necrosis, infiltrates, vascular patterns, and histologic patterns), Mann-Whitney U tests were performed to test the hypothesis that MR imaging parameter values did not differ among groups. While histologic features produce categoric scores (cell density, cell atypia, and overall vascular score), multivariate analysis of variance was used to test the hypothesis that MR imaging parameter values did not differ among groups. Spearman correlation analysis was performed to assess the relationship between quantitative histologic measures (mitotic activity, endothelial surface area, vascular surface area, and vascular profile count per square millimeter) and MR imaging parameters to identify correlations among the individual MR imaging measurements. For Mann-Whitney U and ANOVA testing, a result was considered significantly different with *P* < .01, given the number of variables assessed. For the Spearman correlation analysis, significance was <.05.

**RESULTS**

Twenty-eight untreated newly presented GBMs were included in the study (10 women; age range, 38–76 years; mean, 60 years). Histologic specimens were obtained from 12 biopsies and 16 surgical debulkings. The results of statistical analyses for comparisons of histologic and MR imaging measures are summarized in the On-line Table.

The presence of frank necrosis was associated with significantly higher values of *K*\text{trans} (*P* = .005, *Fig 1A*). Significantly higher values of *v* \text{e} were seen in the presence of fibrillary histology (estimated *P* = .007, *Fig 1B*). A positive correlation was found between *v* \text{e} and mitotic activity (*P* = .012, *r* = 0.470, *Fig 2*). No relationship was seen between mitotic activity and any of the descriptive or semiquantitative histology measures, suggesting that the relationship between *v* \text{e} and mitotic activity is independent of the relationship between *v* \text{e} and the presence of fibrillar histology. No correlation was observed between *v* \text{e} and cell density.

Cross-correlations between individual MR imaging parameters are summarized in Table 1. Positive correlations were found between *K*\text{trans} and all other MR imaging parameters. Significant correlations were present between *v* \text{e} and *K*\text{trans} (*P* < .05 and *r* = 0.450) and between *v* \text{e} and EnF. Mitotic activity did not correlate

![FIG 1. A, Boxplot of fibrillar histology and *v* \text{e} (estimated *P* = .007). Sample histologic specimens showing tumors without (B) and with (C) the presence of fibrils (small fibers measuring approximately 1 mm, black arrows, C). H&E stain ×40 magnification. D, Boxplot of frank necrosis and *K*\text{trans} (estimated *P* = .005).](image1)

![FIG 2. Scatterplot of mitotic activity versus *v* \text{e} (*P* = .012, *r* = 0.470), marker shapes depict separate scores of cell density measures.](image2)
This study identified an unexpected positive correlation between \( v_e \), a parameter thought to reflect extravascular extracellular space or cell density measures. Therefore, short sampling times will lead to relative underestimation of \( v_e \). It has also demonstrated that summary statistics presented in this and other studies reflect only perfused tissue from estimates of summary statistics, which could otherwise produce artificially low values of \( v_e \). In addition, the relatively low dynamic sampling duration (6 minutes) will affect \( v_e \) estimates to some degree. First, model fitting errors (assuming that the model is correct) will result from undersampling, but modeling studies suggest that these fitting errors are likely to be very small. Second, short sampling times will lead to relative underestimation of slow tissue exchange compartments, which would tend to reduce the impact of necrotic tissue on \( v_e \) estimation as described in several studies. These potential model fitting errors and short sampling times imply that tumors with greater necrosis would have shown higher \( v_e \) values if sampling had continued for a longer time and that the measured values of \( v_e \) in this study are more likely to reflect the extravascular extracellular space fraction in viable tissue.

The positive correlation between \( v_e \) and mitotic activity is surprising. Tumors with larger \( v_e \) values exhibited more mitotic activity, the inverse of what one might expect (and the inverse of our initial hypothesis), whereby more proliferative tumors would be more densely packed with cells. Neither mitotic activity nor \( v_e \) related to the presence of necrosis, though the lack of a relationship between \( v_e \) and necrosis may, in part, reflect the relatively low dynamic collection period (see above). These observations suggest that the size of the extravascular extracellular space in perfused enhancing tumor tissue is truly related to mitotic rate and not simply a reflection of elevated measures of \( v_e \) due to increased necrosis in rapidly proliferating tumors.

This suggestion is initially counterintuitive. In normally developing tissues, mitotic activity is higher in areas of low cell packing due to the inhibition of proliferation in response to cell-to-cell contact, a process known as contact inhibition of proliferation in developing normal tissue. No similar relationship has been described in malignant tissues, and loss of contact inhibition of mitosis is one of the hallmarks of the cancer cell. This finding leads to the hypothesis, stated in the introduction, that rapidly prolif-
erting tumors will continue to proliferate, leading to increased cell density and a decreased size of the extravascular extracellular space and consequently of \( v_e \). We have no evidence to explain why the observed relationship should exist. One possible explanation may be that tumors with high cellular density still have impaired responses to tumoral growth factors despite loss of contact inhibition of proliferation. Another possible explanation is that tumors with short mitotic cycles are characterized by a reduced time for cellular maturation, resulting in smaller cells and reduced cell packing. Whatever the underlying biologic mechanism, this finding appears particularly interesting and requires further study.

The ability to obtain an MR imaging–based biomarker of mitotic activity and/or tumor cell proliferation is highly desirable. The findings presented here may be tumor-specific or reflect an unrecognized phenotype, but the possibility that \( v_e \) may be a potential marker of mitotic activity merits further evaluation.

The data also demonstrated a positive relationship between the presence of tumor cell fibrils within the histologic specimens and higher \( v_e \), though the numbers were very small (\( n = 4 \)). Fibrillar cell processes are cell extensions containing cytoplasm, which are surrounded by cell membranes. These are visible through a microscope and allow tumor astrocytes to be recognized as “fibrillar.” A less cell-dense arrangement due to such cell extensions and/or intercellular edema facilitates its recognition via light microscopy. Thus, the ability to identify fibrillar cell processes suggests that the tumor cells are more loosely packed or that there is localized extracellular edema. No relationship was identified between fibrillar histology and mitotic activity, indicating that the relationships between \( v_e \) and these measures are independent.

Most previous work evaluating the relationship between specific histologic features and DCE-MR imaging has focused predominantly on CBV derived from DSC techniques with significant relationships seen among CBV and microvascular density, VEGF expression, cell density, endothelial proliferation, and mitotic activity. No such relationships were identified between the latter 3 measures and \( v_p \) in this study. A study by Lüdemann et al. compared a variety of MR imaging techniques with \( H_2O^{15} \)-PET for measuring perfusion and found that while both DSC and T1-weighted DCE-MR imaging techniques correlated with the criterion standard \( H_2O^{15} \)-PET measure, only borderline correlation was seen between the DSC techniques and the T1-weighted technique, whereby DSC-derived blood volumes were generally lower than those derived from the T1-weighted DCE-MR imaging technique. This finding may account for the failure of \( v_p \) to relate to any of the histologic measures in this study.

The major limitation of the present work is the lack of stereotactic image–matched histologic specimens; therefore, correlation between the histology and the DCE-MR imaging measurements at a local level cannot be made. GBMs are notoriously heterogeneous tumors, and histologic analysis of small tumor specimens may lead to undergrading of a tumor if the sample is not a true reflection of the tumor as a whole. At the time of the study, image-matched histologic samples were not obtained and there was no concurrent postoperative imaging performed, which could have helped to identify the site of the histologic sample. All histologic specimens in this study did confirm the diagnosis of GBM and therefore are considered representative samples. Previous nonimage–registered studies comparing \( v_e \) with histopathologic vascular measures have been reported, but these have been performed with a selection of 3–4 small ROIs, taking the maximal value of \( v_e \) and \( Ktrans \), which is unlikely to provide a representation of the tumor as a whole. In our current study, for the DCE-MR imaging measures \( Ktrans \), \( v_e \), and \( v_p \), median values from whole-tumor VOIs were used. Studies using whole-tumor VOIs have previously identified significant differences in DCE-MR imaging parameters among tumor grades and have been shown to convey important potential prognostic information. Thus, a comparison between small histologic samples and DCE-MR imaging parameters from whole-tumor VOIs seems reasonable. Further work is required to confirm that the correlations identified in the current study hold true for image-matched stereotactic samples. In addition, some histologic features such as Ki-67 and VEGF expression, which have previously shown correlation with DSC-MR imaging–derived CBV, were not available.

**CONCLUSIONS**

The DCE-MR imaging–derived measure \( v_e \) has been identified as a potential correlate of mitotic activity in GBM. While this is an interesting result, our understanding of the biologic mechanisms responsible for this possible relationship is limited. Further work with the correlation of \( v_e \) to more precise measures of cell density, additional markers of cellular and vascular proliferation, and measures of VEGF expression is warranted.

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