Imaging Surrogates of Infiltration Obtained Via Multiparametric Imaging Pattern Analysis Predict Subsequent Location of Recurrence of Glioblastoma

BACKGROUND: Glioblastoma is an aggressive and highly infiltrative brain cancer. Standard surgical resection is guided by enhancement on postcontrast T1-weighted (T1) magnetic resonance imaging, which is insufficient for delineating surrounding infiltrating tumor.

OBJECTIVE: To develop imaging biomarkers that delineate areas of tumor infiltration and predict early recurrence in peritumoral tissue. Such markers would enable intensive, yet targeted, surgery and radiotherapy, thereby potentially delaying recurrence and prolonging survival.

METHODS: Preoperative multiparametric magnetic resonance images (T1, T1-gadolinium, T2, T2-weighted, T2-weighted fluid-attenuated inversion recovery, diffusion tensor imaging, and dynamic susceptibility contrast-enhanced magnetic resonance images) from 31 patients were combined using machine learning methods, thereby creating predictive spatial maps of infiltrated peritumoral tissue. Cross-validation was used in the retrospective cohort to achieve generalizable biomarkers. Subsequently, the imaging signatures learned from the retrospective study were used in a replication cohort of 34 new patients. Spatial maps representing the likelihood of tumor infiltration and future early recurrence were compared with regions of recurrence on postresection follow-up studies with pathology confirmation.

RESULTS: This technique produced predictions of early recurrence with a mean area under the curve of 0.84, sensitivity of 91%, specificity of 93%, and odds ratio estimates of 9.29 (99% confidence interval: 8.95-9.65) for tissue predicted to be heavily infiltrated under the curve. Tissue predicted to be heavily infiltrated showed a statistical difference in early recurrence. Imaging predictions were validated via radiopaque markers placed in the peritumoral regions of interest on postoperative images.

CONCLUSION: Visually imperceptible imaging patterns discovered via multiparametric pattern analysis methods were found to estimate the extent of infiltration and location of future tumor recurrence, paving the way for improved targeted treatment.

KEY WORDS: Glioblastoma, Imaging pattern analysis, Infiltration, Multiparametric, Recurrence

Glioblastoma, a grade IV astrocytoma, is the most common and aggressive primary brain tumor with a median survival of 12 to 15 months after diagnosis.1 Maximal safe surgical resection combined with adjuvant radiotherapy and chemotherapy has been the cornerstone in the management of these high-grade gliomas.2-4 Advanced imaging modalities have improved the precision of radiotherapy through enhanced tumor delineation, whereas conformal...
radiotherapy and intensity-modulated radiation therapy allow for an increased radiation dose to the tumor bed and surrounding margin while preserving critical brain structures. 

Glioblastoma is well-known to extend beyond the visible borders of the enhancing tissue on magnetic resonance imaging (MRI), which is the primary target of surgical treatment. In particular, stereotactic biopsy specimens have revealed the presence of tumor in regions that were presumed to be either normal or edematous brain on the basis of MRI characteristics. Inclusion of this peritumoral region in the clinical target volume has been an area of contention when planning postoperative radiotherapy. This is in large part due to the fact that the risk of radiation toxicity increases with escalating target volume and dose. Nevertheless, dose escalation in the peritumoral region has been shown to confer a survival advantage despite a higher frequency of white matter abnormalities. Meanwhile, mounting evidence in the surgical literature suggests that the aggressive extent of resection, independent of age, is likewise associated with improved survival.

Thus, the peritumoral region remains a critical problem for both understanding and effective treatment of glioblastoma. An altered imaging characteristic of this region, commonly referred to as peritumoral edema, represents a combination of infiltrating tumor cells and vasogenic edema secondary to angiogenic and vascular permeability factors released by adjacent tumor cells, where a compromised blood-brain barrier (BBB) allows intravascular proteins and fluid to penetrate the parenchymal extracellular space. It is difficult to distinguish infiltrating neoplasm from vasogenic edema with standard imaging approaches; therefore, it is important to identify imaging biomarkers that distinguish regions of densely infiltrating glioblastoma from the biophysical response of surrounding brain tissue to inflammatory factors.

Multiparametric MRI has been the most informative noninvasive method for comprehensive characterization of tumor and surrounding brain tissue. Various acquisition protocols reflect different properties of tissue; however, no single imaging metric is currently sufficient to delineate areas of nonenhancing tumor infiltration. Pre- and postgadolinium T1-weighted (T1) images contain information about regional angiogenesis and the integrity of the BBB in the tumor. T2-weighted (T2) and T2-fluid attenuated inversion recovery (FLAIR) images are helpful for assessing nonenhancing tumor and edema extent because they are sensitive to water concentration. Diffusion imaging techniques such as diffusion tensor imaging (DTI) map the diffusion process of water in the brain, affected in part by tumor cell density. Dynamic susceptibility contrast-enhanced (DSC) MRI techniques reflect various aspects of perfusion in the brain, which provide quantitative measures of regional microvasculature and hemodynamics. These patterns are variably affected by infiltrating tumor cells, and independently they are not sufficiently specific enough to clearly define areas of tumor infiltration.

In the current study, we tested the hypothesis that advanced pattern analysis and machine learning methods applied to multiparametric MRI are able to quantitatively capture subtle and otherwise imperceptible imaging variations that highlight heterogeneity within edematous peritumoral tissue. Moreover, we tested the hypothesis that these methods can provide predictive spatial maps of tumor infiltration and the likelihood of early recurrence. Understanding subtle imaging characteristics of infiltrating glial tumors can lead to more aggressive, yet targeted, surgical and radiation treatment strategies that aim to maximize treatment efficacy while maintaining neurological function.

METHODS

Study Setting

Institutional review board approval was obtained for this study. Our research consisted of a retrospective cohort and a replication study cohort at the University of Pennsylvania from 2006 to 2012.

Participants

The retrospective cohort included 31 patients who had de novo glioblastoma (World Health Organization grade IV) and were treated at our institution. The inclusion criteria consisted of preoperative advanced MRI (ie, anatomic, diffusion, and perfusion imaging modalities) and subsequent recurrence. In all subjects, the clinical diagnosis of tumor recurrence was established with histopathologic analysis after repeat resection. Subjects who had a previous tumor, a previous resection, or residual tumor after surgical resection were excluded. Residual tumors were defined as any contrast-enhancing areas identified by the neuroradiologist (M.B., 14 years of experience) on the immediate postoperative MRI. The replication study included an additional 34 patients with de novo glioblastoma who met the aforementioned inclusion and exclusion criteria. All subjects in this analysis received the same treatment, ie, gross total resection of enhancing tumor followed by temozolomide and radiotherapy. The average time to recurrence was 8.04 months (median, 6.5 months).

Quantitative Variables

Preoperative MRI was performed using a 3-Tesla scanner (Tim Trio; Siemens Medical Solutions, Erlangen, Germany). T1, T2, T2-FLAIR, DTI, and DSC-MRI were captured for all the patients included in this analysis before the surgery (see Figure, Supplemental Digital Content 1, http://links.lww.com/NEU/A835). All MRIs were affinely registered intrasubject, smoothed, inhomogeneity corrected, and skull stripped as previously described. For imaging sequence details, please see Supplemental Digital Content 2, http://links.lww.com/NEU/A836.

Tumor Segmentation

To precisely segment the tumor and peritumoral edema, we used state-of-the-art GLISTR software (University of Pennsylvania, Philadelphia, Pennsylvania) to delineate tumor (enhancing and nonenhancing regions) and edematous regions for which the infiltration maps were analyzed. GLISTR software uses an image-analysis technique that incorporates probabilistic imaging and biophysical models of tumor growth and appearance to segment brain tumors and their surrounding tissues using 4 MRI modalities (T1, T1-gadolinium, T1, and T2-FLAIR).

MRI Features

Principal component analysis was used to extract all the information conveyed by the DSC-MRI, and this information was used to create the
Principal component analysis is a standard dimensionality reduction method that was used here to distill the time series of DSC-MRI down to a few parameters that capture the temporal dynamics of blood perfusion (see Figure, Supplemental Digital Content 3, http://links.lww.com/NEU/A837).

The machine-learning model combines information from 4 structural magnetic resonance images (T1, T1-gadolinium, T2, and T2-FLAIR), 4 DTI-derived images (ie, fractional anisotropy, radial diffusivity, axial diffusivity, and trace, and perfusion images (principal components 1 through 5 (PC1-PC5)). In addition, because relative cerebral blood volume is a commonly used normalized ratio of blood volume, which is calculated based on an estimate of the arterial input function, we used relative cerebral blood volume as an additional feature.

Infiltration Model

A multidimensional pattern classification method called support vector machines was used to create the infiltration tissue pattern classifier. To magnify the heterogeneity within peritumoral edema, 2 extremities were selected to train the model. The near extremity was defined as the area immediately adjacent to the enhancing tumor, whereas the distal edge of the edema was designated as the far extremity (see Figure, Supplemental Digital Content 4, http://links.lww.com/NEU/A838). These 2 regions served as reference examples for near-tumor and far-from-tumor tissue based on the expectation that they are likely to have relatively higher and lower infiltration, respectively. The classifier was trained on the labeled voxels using A Library for Support Vector Machines with a gaussian kernel function, weighting to balance the classes and parameters C and sigma in a standard support vector machine model based on a leave 1 subject out cross-validated grid search on the population.

In our modeling, this voxelwise map signifies spatial pseudoprobability of infiltration that was named the infiltration index. It may be noted that the infiltration index has a value between 0 and 1 representing non-infiltrated and infiltrated tissue, respectively.

Retrospective Cohort Study

Leave 1 subject out cross-validation was used in the retrospective study. Specifically, to produce the infiltration index map for 1 patient, the data from that patient were put aside, and the classifier was trained using the other 30 patients in the retrospective cohort study, which tested the remaining patient. This process was repeated 31 times, and each time a different patient was left out. This cross-validation process is critical for providing realistic estimates of how well the predictive model is likely to generalize to new patients.

Replication Study

The classifier was trained on 31 patients who participated in the retrospective cohort study. The trained classifier was used to create infiltration maps in 34 new patients who participated in the replication study.

Statistical Methods

The evaluation was performed by comparing the created infiltration map to 2 regions of interest (ROIs): recurring and nonrecurring. Recurring tissue ROIs were manually drawn by experts (H.A., L.M., M.B., R.L.W.) with the intention of selecting a small region on the preoperative MRI from which the recurrence originated (recurrence being estimated from the follow-up MRI). Preoperative mass effect, resection, and inexact registration between preoperative and follow-up recurrence scans made it nearly impossible to precisely delineate on the preoperative images the follow-up recurrence origin. Therefore, the recurrence origin was approximately placed by the raters in small regions with relatively high certainty. A 3-mm margin around the manually estimated origin-of-recurrence regions was included in the recurrence ROIs, assuming that they were likely to be heavily infiltrated, but also accounting for registration uncertainties in placing the recurrence ROIs. An additional margin between 3 and 10 mm was excluded from any labeling as recurrence or nonrecurrence because of its immediate proximity to actual recurrence, but its nonrecurrence status was deemed to render it unsuitable for evaluation, given that tumor recurrence would be considered likely to progress quickly to its surrounding, likely aggressively infiltrated, tissue. Nonrecurring ROIs were defined as all remaining peritumoral edematous regions between 5 and 20 mm around the tumor core (enhancing tumor plus nonenhancing core). The 20-mm maximum distance of evaluation was used to avoid artificially overestimating the predictive value of our analysis because distant edema is naturally unlikely to recur early; therefore, a model predicting this correctly would have limited added value over common clinical knowledge. Similarly, the 5-mm margin around the tumor was excluded in part to account for the fact that some peritumoral nonenhancing tissue is typically removed at our institution during surgery and also in part to account for registration uncertainties in mapping recurrence to preoperative scans. The 5-mm margin was also excluded to avoid overly optimistic evaluations of our predictions because the majority of recurrences occur close to the tumor, which is also where the infiltration index is high by construction, thereby rendering the added value of our predictions limited in that region.

To statistically determine the quality of the infiltration index maps, a receiver-operating characteristic curve was drawn and the area under the curve (AUC) was calculated. To draw the receiver-operating characteristic curve, sensitivity and specificity were calculated using the labeled maps for each subject and the training ROIs identified by the experts. The odds ratio was calculated to quantify how strongly the estimated preoperative infiltration maps were associated with subsequent recurrence.

| TABLE. Study Results for Retrospective (Cross-validated) and Prospective Cohorts |
|-------------------------------------------------|-------------|----------|--------|--------|---------|--------|
| OR (99% CI), % | AUC | Accuracy, % | Sensitivity, % | Specificity, % | P Value |
| Retrospective | 11.17 (10.71-11.64) | 0.80 | 87.11 | 93.55 | 88.06 | <.0001 |
| Prospective | 8.29 (8.95-9.65) | 0.84 | 91.25 | 91.18 | 93.48 | <.0001 |

OR, odds ratio; CI, confidence interval; AUC, area under the curve. Statistics are based on positive (recurrence) and negative (nonrecurrence) regions of interest.
RESULTS

Participants

Retrospective Cohort Study and Model Generation

The Table shows the results for the 31 retrospective cohort patients, and Figure 1 shows a color-coded tumor infiltration index map for a representative patient. Classification results include a mean AUC of 0.80, sensitivity of 93%, specificity of 88%, accuracy of 87%, and recurrence odds ratio estimates of 11.17 (99% CI, 10.71-11.64; \( P < .0001 \)) for tissue predicted to be relatively more infiltrated in the retrospective cohort study. All these results were obtained using cross-validation.

Replication

After cross-validation, we created a single model using the MRI data of 31 retrospective subjects. This model was utilized to
generate the infiltration index maps for 34 new subjects who participated in the replication study. The Table shows the results for this cohort, and Figure 2 shows color-coded tumor infiltration index maps for a representative subject. This method produced predictions with a mean AUC of 0.84, sensitivity of 91%, and specificity of 93%, and recurrence odds ratio estimates of 9.29 (99% confidence interval: 8.95-9.65; \( P < .0001 \)) for tissue predicted to be relatively more infiltrated in the replication study. Figure 3 shows the receiver-operating characteristic curve for these subjects.

Imaging Features in Recurrence and Nonrecurrence

Five PCs captured more than 99% of the variance in the perfusion signal, which quantified all subtleties of these time curves. Each PC conveys different characteristics of the perfusion signal (see Figure, Supplemental Digital Content 5, http://links.lww.com/NEU/A839). MRI signals contained significant discriminating information, which has been organized in Figure 4 into 3 rows in accordance with respective interpretation discussed in the following section (see Figure, Supplemental Digital Content 6, http://links.lww.com/NEU/A840). Figure 4 also

**FIGURE 2.** Replication study results. **Left,** an estimated map for tumor infiltration from preoperative magnetic resonance imaging analysis; yellow arrow points to a region estimated to be relatively more infiltrated. **Right,** the corresponding magnetic resonance images after tumor resection and subsequent recurrence (red arrow) for the same patient. Recurrence occurred in the vicinity of peritumoral tissue originally estimated to be highly infiltrated. The first row represents a 3-dimensional rendering of the images. The second, third, and fourth rows show T1-weighted magnetic resonance images with contrast fused infiltration map, T1-weighted magnetic resonance images with contrast, and T2-weighted fluid attenuated inversion recovery (FLAIR), respectively. Gad, gadolinium.
shows the distribution of our infiltration index in recurring and nonrecurring tissue, indicating markedly greater separation compared with any individual MR signal. The infiltration index is calculated by integrating the top imaging features for both tissue groups to determine how distinctive (separable) the classifier is for each group.

**DISCUSSION**

Current surgical treatment of glioblastoma is guided largely by enhancement on the postgadolinium T1 MRI, and it is mainly this region that is the focus of resection. Although this practice is known to leave the majority of the infiltrating tumor unresected, this is inevitable in the absence of spatially specific knowledge of the infiltration pattern of the tumor. Likewise, the clinical target volume of radiation therapy includes the resection bed plus a variable margin that commonly receives a reduced and largely spatially uniform radiation dose. Although this practice is aimed to address the infiltrative component of glioblastoma, recurrence in this disease is currently certain and almost always occurs within this peritumoral edematous tissue. Nonetheless, aggressive use of both treatment modalities has been associated with prolonged survival. Surgical treatment confers a greater survival benefit as the extent of resection increases, whereas dose escalation within the peritumoral region similarly leads to longer survival compared with more traditional radiotherapy plans. It stands to reason that the efficacy and safety of these therapies could be significantly improved if the spatial extent of the infiltrative component were known and could be specifically targeted.

**Key Results and Interpretation**

The current study addresses a critical and unresolved need in the field of glioblastoma therapy; specifically, by using advanced analytical methodologies, we can quantify subtle, yet important, heterogeneity within the peritumoral region and provide an estimate of the spatial extent and pattern of tumor infiltration. Although we used the current standard-of-care MRI data for our analysis, this information is not apparent when the imaging data are interpreted using conventional means and measurements, but this can be teased out by integrating multiparametric images with an optimized predictive algorithm. These estimated infiltration patterns represent a promising tool for characterizing the spatial pattern of glioblastoma infiltration and pave the way for aggressive, yet targeted, treatment of the infiltrated peritumoral region.

The second row summarizes statistics from features derived from DTI, which reflect lower mean diffusivity and increased fractional anisotropy in regions of higher infiltration. This finding is consistent with the hypothesis that these regions harbor a higher ratio of malignant cells to fluid content. The T1 signal is also higher in those regions, which would be consistent with lower fluid content. Finally, slightly higher T1-gadolinium implies relatively more compromised BBB in tissue that later recurred, also consistent with infiltrating tumor characteristics.

The third set of imaging features relates aspects of tissue vascularization, perfusion, and permeability of blood vessels. Glioblastoma remains the most angiogenic human glioma, therefore exhibiting extensive neovascularization. When brain tumors exceed a critical volume, the resultant ischemia triggers the secretion of angiogenic factors. These factors, such as vascular endothelial growth factor, promote vascular proliferation leading to the formation and maintenance of tumor vessels. These new, immature vessels tend to be tortuous and leaky. PC2 is inversely related to the magnitude of signal drop in relation to the baseline (see Figure, Supplemental Digital Content 5, http://links.lww.com/NEU/A839). Our results indicate a relatively lower PC2 in areas of high infiltration, indicative of a higher degree of BBB compromise and leaky neovascularulation. Finally, the relatively higher values of PC3 in these regions were particularly interesting as they suggested a relative time delay in the contrast agent reaching the highly infiltrated tissue, likely due to higher flow resistance, tortuosity, and other characteristics of tumor vasculature, which is in contrast to healthy brain vasculature known to display extremely low blood flow resistance. In addition...
to this time delay, the infiltrated tissue was found to have slower and incomplete signal recovery, both indicative of immature, leaky vessels. Remarkably, even though each of these MRI signals displayed very subtle differences between recurrence and nonrecurrence, which were not useful for prediction individually, the integration of these complementary signals via machine learning methods provided predictive indices of infiltration and recurrence that can potentially guide personalized treatment.

**Limitations**

An important limitation of this study is the lack of a histopathologic ground truth from the infiltrated peritumoral areas. Although it is not practical, or sometimes even possible, to finely sample the peritumoral region in routine clinical practice, in our future work, we plan to perform targeted biopsies in several of these regions and relate the imaging-based predictions to pathology-proven concentration of malignant cells. This model also predicts areas that have a high probability of recurrence, where, in
fact, no recurrence was observed on follow-up imaging. Due to the aggressive and infiltrative nature of glioblastoma, it is quite likely that these areas also harbor a higher concentration of malignant cells that will recur in the future, given enough time. Future work will elaborate how these areas are biologically different.

Finally, it is important to emphasize that the near and far peritumoral tissue ROIs used to train our model does not imply any certainty that there is more infiltration near to as opposed to far from the tumor. These regions should be viewed as part of the algorithmic procedure used to derive the model that highlights the heterogeneity within peritumoral edema. Although there are certainly other ways to derive predictive models, our choice was not arbitrary, but it was based on ample literature that generally shows a denser malignant cell concentration near the enhancing solid tumor component.\(^\text{8,9}\)

**CONCLUSION**

The results of this study demonstrate that new ways of analyzing multiparametric MRI data can elucidate patterns of tumor infiltration within the peritumoral region that predict subsequent tumor recurrence, a task that is virtually impossible based on visual interpretation of standard MRIs. The proposed methodology could offer significant advantages to clinicians treating patients with newly diagnosed glioblastoma. Neurosurgeons could potentially perform supratotal resections when anatomically possible, addressing highly infiltrated peritumoral tissue at higher risk of recurrence. Likewise, radiation oncologists could escalate the radiation dose to high-risk areas. Although our modeling was built specifically for evaluating the infiltrative component of glioblastoma, our methodology is quite general and could be applied to a host of image analysis problems.

**Disclosures**

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**REFERENCES**

Supplemental digital content is available for this article. Direct URL citations appear in the printed text and are provided in the HTML and PDF versions of this article on the journal’s Web site (www.neurosurgery-online.com).

COMMENT

The multimodality, multiparametric approach using machine learning algorithms has been increasingly used to develop prognostic and predictive imaging biomarkers in neuro-oncology.1-6 The study by Akbari et al has taken this a step further by devising a spatially resolved predictive model of tumor recurrence. The clinical implication of this biomarker is significant; it can potentially be used to define over- or undertreated tumor at a voxelwise resolution. Furthermore, the magnetic resonance imaging (MRI) acquisition techniques from which the model is derived are those routinely performed for preoperative evaluation at most cancer centers, although the generalizability of the model needs to be confirmed by multicenter imaging data. This work also requires prospective validations to assess the model’s predictive value when patients with long-term non-recurrent tumor are included.

Raymond Y. Huang
Alexandra J. Golby
Boston, Massachusetts