CLINICAL STUDY



Gyriform infiltration as imaging biomarker for molecular glioblastomas

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Abstract

Background Molecular glioblastomas (i.e. without the histological but with the molecular characteristics of IDH-wild-type glioblastoma) frequently lack contrast enhancement, which can wrongly lead to suspect a lower-grade glioma. Herein, we aimed to assess the diagnostic value of gyriform infiltration as an imaging marker for molecular glioblastomas.

Methods Two independent investigators reviewed the MRI scans from patients with newly diagnosed gliomas for the presence of a gyriform infiltration defined as an elective cortical hypersignal on MRI FLAIR sequence. Diagnostic test performance of this sign for the diagnosis of molecular glioblastoma were calculated.

Results A total of 426 patients were included, corresponding to 31 molecular glioblastoma, 294 IDH-wild-type glioblastoma, 50 IDH-mutant astrocytoma, and 51 IDH-mutant 1p19q-codeleted oligodendroglioma. A gyriform infiltration was observed in 16/31 (52%) molecular glioblastoma, 40/294 (14%) IDH-wild-type glioblastoma, and none of the IDH-mutant glioma. All the 56 gyriform-infiltration-positive tumors were IDH-wild-type and all but two had a *TERT* promoter mutation. The inter-rater agreement was good ($\kappa = 0.69$, p < 0.001). The sensitivity, specificity, positive predictive value and negative predictive value of the presence of a gyriform infiltration for the diagnosis of molecular glioblastoma were 52%, 90%, 29%, 96%, respectively. The median overall survival was better for gyriform-infiltration-negative patients compared to gyriform-infiltration-positive patients in the whole series and in patients with non-enhancing lesions (n=95) (25.6 vs 16.9 months, p = 0.005 and 20.2 months vs not reached, p < 0.001).

Conclusion Gyriform infiltration is a specific imaging marker of molecular glioblastomas that can help distinguishing these tumors from IDH-mutant lower-grade gliomas.

Keywords Imaging marker · T2-FLAIR · IDH-wild-type astrocytoma · TERT · Molecular glioblastoma

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Introduction

Several studies have shown that IDH-wild-type diffuse astrocytic gliomas display molecular features of glioblastoma (GBM) and, similar to IDH-wild-type GBM, are associated with a poor prognosis. Therefore, these tumors have been classified in the World Health Organization (WHO) 2021 classification [1] of central nervous system tumors as IDH-wild-type GBM [2, 3]. Thus, according to the latter classification, a diagnosis of IDH-wild-type GBM can be made in adults if the histo-molecular analysis finds an IDH-wild-type diffuse and astrocytic glioma with microvascular proliferation or necrosis, or a TERT promoter (TERTp) mutation, or an EGFR amplification, or a combined whole-chromosome-7 gain and chromosome-10 loss. However, in contrast to IDH-wild-type GBM diagnosed on histological characteristics (presence of microvascular proliferation or necrosis), contrast enhancement is frequently lacking in IDH-wild-type GBM diagnosed solely on the molecular profile (molecular GBM) and these tumors have a radiological presentation that can wrongly suggest a low grade glioma (LGG) [4–6]. Indeed, in a recent meta-analysis, 42% of grade II or III IDH-wild-type astrocytomas were found not to display contrast enhancement on magnetic resonance imaging (MRI), compared to 57% and 45% of IDH-mutant low grade astrocytomas and oligodendrogliomas, respectively [7]. Therefore, imaging biomarkers that could allow distinguishing molecular GBM from LGG would be helpful. Recently, important advances have been made in the non-invasive molecular characterization of gliomas. For example, 2-hydroxyglutarate MR spectroscopy enables to identify IDH-mutant gliomas [8] and the presence of a T2/ FLAIR mismatch sign is highly specific of IDH-mutant astrocytomas [9–11]. Additionally, molecular GBM have been shown to frequently display areas of elective FLAIR hyperintensity limited to the cortical grey matter[4]. The aim of the present study was to assess the specificity and reproducibility of this radiological feature, designated as "gyriform infiltration", as an imaging marker for the noninvasive detection of molecular GBM in an independent cohort of diffuse gliomas.

Methods

Patient selection

and June 2020 and reviewed their clinical, radiological, histological, and molecular characteristics. The list of patients was retrospectively obtained from the records of the neuropathology department. Patients were included if they were aged \geq 18 years, had a diffuse glioma according to 2016 WHO brain tumor classification, had an available TERTp mutation and IDH mutation status, and had an MRI scan performed at diagnosis (before surgery or biopsy) for radiological review with T2 FLAIR and postcontrast T1-weighted sequences. Patients with glioma limited to the brainstem or cerebellum, or with a H3K27M or an H3G34 mutation, were excluded from the analysis. We defined molecular GBM as IDH-wild-type astrocytomas without histological characteristics of GBM (presence of microvascular proliferation or necrosis) but with molecular alterations (TERTp mutation and/or EGFR amplification).

Molecular data

Data regarding *IDH1*, *IDH2*, *H3-3A/H3C2*, BRAF V600E, and TERTp mutations, *EGFR* gene amplification, ATRX expression, O6-methylguanine-DNA methyltransferase (*MGMT*) promoter methylation, and chromosomes 1p and 19q codeletion were obtained from the records of the neuropathology department. The majority of samples were molecularly characterized using a dedicated next generation sequence (NGS) panel enabling to test genetic mutations and loss or gain of chromosomal regions characteristic of gliomas [12, 13]. TERTp mutation was tested using a droplet Polymerase Chain Reaction (PCR) using commercial probes (Biorad) and confirmed using the NGS panel [14].

Imaging protocol

Elective gyriform infiltration was defined as an area of FLAIR hyperintensity limited to the cortical grey matter, without involvement of the underlying white matter and without contrast enhancement.

Brain MRI were acquired at 1.5 T or 3.0 T and were independently reviewed by two investigators (E.M. and F.D.) for the presence of a gyriform infiltration. The two investigators were blinded to the clinical history of the patient and the molecular characteristics of the lesion during this review. If there were discordant reviews between the two investigators, both assessed the MRI sequences a second time (agreement by consensus).

Statistical analysis

Sensitivity (Se), specificity (Sp), positive predictive value (PPV), and negative predictive value (NPV) were calculated using the final score after adjudication. An interrater agreement analysis was performed to determine the reproducibility between the two reviewers using Cohen's kappa statistic (κ). A κ value ≤ 0.2 indicates slight agreement, 0.21–0.4 fair agreement, 0.41–0.6 moderate agreement, and > 0.6 substantial agreement[15]. Comparisons of categorical variables were performed using the Fisher's exact test, and comparisons of quantitative variables were performed using the Student's t-test. The probability of survival was estimated using the Kaplan–Meier method from the date of the histological diagnosis to the date of last follow-up or death, and differences between curves were assessed using the Log-rank test. All calculations were performed using SPSS software package version 28.0 (SPSS Inc, IBM Corp, Armonk, New York), and *p* values < 0.05 (two-sided) were considered significant.

The design of the study was approved by the institutional review board and conducted according to the European ethical guidelines (MR004 $n^{\circ}20_{5178}$).

Results

Tumor characteristics

We retrospectively identified 426 patients who met the inclusion criteria (flowchart in Fig. 1). The tumor population

consisted in 31 molecular GBM, 294 IDH-wild-type GBM, 50 grade II or grade III IDH-mutant astrocytomas, and 51 grade II or grade III IDH-mutant and 1p19q codeleted oligodendrogliomas (Table 1). Molecular GBM presented as an IDH-wild-type grade II astrocytoma in 10/31 (32%) patients and as an IDH-wild-type anaplastic astrocytoma in 21/31 (68%) patients, and were classified as molecular GBM because of the presence of a TERTp mutation (30/31, 97%) or/and an *EGFR* amplification (16/31, 52%). A T2-FLAIR mismatch was identified in 11/95 (12%) patients presenting non-enhancing tumors. All of these patients had an IDH-mutant astrocytoma (11/50; 22%).

Detection of gyriform infiltration

Gyriform infiltration was observed in 56 (13%) patients (Table 2). Representative examples are shown in Fig. 2. Blinded MRI analysis found a substantial inter-rater agreement for gyriform infiltration identification with a κ of 0.69 (p < 0.001; 95% confidence interval [0.61; 0.77]). Both reviewers identified gyriform infiltration as present in 43 (10%) cases and as absent in 351 (82%) cases; 31 (8%) cases were discordant, among these, a gyriform infiltration was identified by reviewer 1 only in 20 cases and by reviewer 2 only in 11 cases. After second assessment, no discordant

656 patients (\geq 18 years old) had a new diagnosis of diffuse glioma between September 2017 and June 2020.



Table 1 Characteristics of the study population subdivided in four histomolecular groups

	IDHmt Astro	OD	Molecular GBM	GBM IDHwt
n	50	51	31	294
Median (range) age, yr	38.0 (18.0-75.9)	47.8 (25.1-85.8)	59.7 (27.9-82.4)	65.9 (21.3-90.2)
Sex, n (%)	· · · ·	· · · ·	· · · ·	
Male	23 (46%)	31 (61%)	21 (68%)	169 (58%)
Female	27 (54%)	20 (39%)	10 (32%)	125 (42%)
KPS at diagnosis, n (%)				
≥70%	49 (98%)	51 (100%)	26 (84%)	221 (75%)
<70%	1 (2%)	0 (0%)	5 (16%)	73 (25%)
Clinical presentation, n (%)				
Seizure	29 (58%)	33 (65%)	21 (68%)	88 (30%)
Focal deficit	8 (16%)	3 (6%)	7 (23%)	98 (33%)
Cognitive deficit	1 (2%)	6 (12%)	1 (3%)	58 (20%)
IHS	9 (18%)	5 (10%)	1 (3%)	43 (15%)
Casual	3 (6%)	4 (8%)	1 (3%)	7 (2%)
Type of surgery, n (%)				
Biopsy	21 (42%)	21 (41%)	29 (94%)	206 (70%)
Partial or complete resection	29 (58%)	30 (59%)	2 (6%)	88 (30%)
Radiological characteristics, n (%)				
Location				
Frontal	34 (68%)	41 (80%)	13 (42%)	138 (47%)
Parietal	15 (30%)	9 (18%)	6 (19%)	100 (34%)
Occipital	0 (0%)	0 (0%)	2 (7%)	31 (10%)
Temporal	18 (36%)	16 (31%)	19 (61%)	138 (47%)
Insula	11 (22%)	10 (20%)	14 (45%)	82 (28%)
Corpus callosum	13 (26%)	9 (18%)	5 (16%)	61 (21%)
Thalamus	0 (0%)	0 (0%)	3 (10%)	31 (10%)
Extension to brainstem or cerebellum	1 (2%)	0 (0%)	2 (6%)	7 (2%)
Gliomatosis	5 (10%)	6 (12%)	20 (64%)	44 (15%)
Gyriform infiltration	0 (0%)	0 (0%)	16 (52%)	40 (14%)
Contrast enhancement	17 (34%)	22 (43%)	8 (26%)	284 (97%)
Multicentric locations	1 (2%)	1 (2%)	7 (23%)	66 (22%)
Edges				
Poorly defined	27 (54%)	22 (43%)	29 (94%)	239 (81%)
Well defined	23 (46%)	29 (57%)	2 (6%)	55 (19%)
WHO grade, n (%)				
Grade II	24 (48%)	24 (47%)	0 (0%)	0 (0%)
Grade III	21 (42%)	30 (59%)	0 (0%)	0 (0%)
Grade IV	5 (10%)	0 (0%)	31 (100%)	294 (100%)
Molecular characteristics, n (%)				
IDHmt	50 (100%)	51 (100%)	0 (0%)	0 (0%)
TERTp-mt	1 (2%)	50 (98%)	30 (97%)	274 (93%)
MGMT meth ^a	39 (78%)	49 (96%)	18 (58%)	151 (51%)
EGFR amp	0 (0%)	0 (0%)	16 (52%)	129 (44%) ^b
1p/19q codel	0 (0%)	51 (100%)	0 (0%)	0 (0%)

IDH isocitrate dehydrogenase, *IDHwt* IDH-wild-type, *TERT* telomerase reverse transcriptase, *TERTp-mt* TERT promoter mutation, *Astro* astrocytoma, *IDHmt* IDH mutant, *GBM* glioblastoma, *OD* oligodendroglioma, *yr* years, *WHO* world health organization, *1p/19q codel* 1p/19q codeletion, *MGMT meth MGMT* promoter methylation, *EGFR amp EGFR* amplification, *Histone mt* histone mutation, *KPS* karnofsky perfomance status, *IHS* intracranial hypertension symptoms

^a 5, 1, 4, and 19 missing values, respectively, for MGMT promoter methylation

^b One missing value

Table 2Summary and
comparisons of the
characteristics of gliomas
presenting or not an elective
gyriform infiltration on T2FLAIR MRI sequences

	Patients with gyriform infiltration	Patients without gyri- form infiltration	p-value
n	56	370	
Mean \pm SD age, yr	63.4 ± 10.2	59.6 ± 15.7	0.018
Sex, n (%)			0.086
Male	38 (68%)	206 (56%)	
Female	18 (32%)	164 (44%)	
KPS at diagnosis, n (%)			0.55
≥70%	44 (79%)	303 (82%)	
	12 (21%)	67 (18%)	
Clinical presentation, n (%)			0.28
Seizure	29 (52%)	142 (38%)	
Focal deficit	15 (27%)	101 (27%)	
Cognitive deficit	6 (11%)	60 (16%)	
IHS	4 (7%)	54 (15%)	
Casual	2 (4%)	13 (3%)	
Type of surgery, n (%)			< 0.001
Biopsy	50 (89%)	227 (61%)	
Partial or complete resection	6 (11%)	143 (39%)	
Radiological characteristics, n (%)			
Location			
Frontal	27 (48%)	199 (54%)	0.45
Parietal	19 (34%)	111 (30%)	0.55
Occipital	5 (9%)	28 (8%)	0.72
Temporal	28 (50%)	163 (44%)	0.40
Insula	23 (41%)	94 (25%)	0.014
Corpus callosum	16 (29%)	73 (20%)	0.13
Thalamus	13 (23%)	21 (6%)	< 0.001
Extension to brainstem or cerebellum	2 (4%)	8 (2%)	0.42
Gliomatosis	36 (64%)	39 (10%)	< 0.001
Contrast enhancement	43 (77%)	288 (78%)	0.97
Multicentric/multifocal locations	21 (38%)	54 (15%)	< 0.001
Edges	(*****)		0.002
Poorly defined	51 (91%)	266 (72%)	
Well defined	5 (9%)	104 (28%)	
WHO grade n (%)		101 (2070)	< 0.001
Grade II	0 (0%)	48 (13%)	(0.001
Grade III	0 (0%)	48 (13%)	
Grade IV	56 (100%)	274 (74%)	
Histomolecular subtypes, n (%)	50 (10070)	2/1 (/1/0)	
Oligodendroglioma	0 (0%)	51 (14%)	< 0.001
Astrocytoma IDHmt	0 (0%)	50 (13%)	< 0.001
Molecular GBM	16 (29%)	15 (4%)	< 0.001
Glioblastoma	40 (71%)	258 (69%)	0.71
Molecular characteristics n (%)	10 (11/0)	230 (05/10)	0.71
IDHmt	0 (0%)	101 (27%)	< 0.001
TERTn-mt	54 (96%)	301 (81%)	0.005
MGMT meth	28 (50%) ^d	230 (62%) ^e	0.005
EGER amp	33 (59%)	$112(30\%)^{f}$	< 0.001
1p/19a codel	0.00%	51 (14%)	0.003
	0(070)	51 (1470)	0.005

IDH isocitrate dehydrogenase, *IDHwt* IDH-wild-type, *IDHmt* IDH mutant, *TERT* telomerase reverse transcriptase, *TERTp-mt* TERT promoter mutation, *TERTp-wt* TERT promoter wild-type, *yr* years, *WHO* world health organization, *1p/19q codel* 1p/19q codeletion, *MGMT meth MGMT* promoter methylation, *EGFR*

Table 2 (continued)

amp EGFR amplification, KPS karnofsky perfomance status, IHS intracranial hypertension symptoms, SD standard deviation

- ^a Patients of the cohort presenting an elective gyriform infiltration on T2 FLAIR MRI sequences
- ^b Patients of the cohort without elective gyriform infiltration on T2 FLAIR MRI sequences
- ^c Comparison between the two groups
- ^d 6 missing values
- e 23 missing values
- f 1 missing value



Fig. 2 Representative axial FLAIR sequences of gyriform-infiltration-positive cases

case remained: the 31 discordant cases were finally considered as gyriform-infiltration-positive for 13 and gyriforminfiltration-negative for 18. In the latter cases, the reason for finally considering the gyriform as absent was related to the non-limitation of the infiltration to the cortex (it also involved the underlying white matter) or the limitation of the infiltration to the white matter (*see Supplementary Figure*).

Progression of gyriform infiltration

A total of 38 patients with gyriform infiltration had an MRI scan performed during follow-up for radiological

review to assess the progression of the sign, which was indeed observed in 28 cases: 20 patients developed contrast enhancement at the site of the gyriform infiltration within a median time of 2 months and 8 patients developed an infiltration of the underlying white matter. Only one patient displayed a partial regression of the gyriform infiltration 6 months after radiochemotherapy. The gyriform infiltration was stable for the remaining 9 patients.

Characteristics of gyriform-infiltration-positive patients

Among the patients with a gyriform infiltration, 54/56 (96%) patients had an IDH-wild-type TERTp-mutant glioma and 2 (4%) had an IDH-wild-type TERTp-wild-type glioma. A gyriform infiltration was observed in 16/31 (52%) patients with a molecular GBM, 40/294 (14%) patients with a GBM IDH-wild-type, but in none of the patients with an astrocytoma IDH-mutant or an oligodendroglioma IDH-mutant and 1p19q codeleted. The gyriform infiltration was significantly more frequent in the subgroup of molecular GBM than in other groups (p < 0.001).

Compared to gyriform-infiltration-negative patients, gyriform-infiltration-positive patients were older at diagnosis (mean age: 63.4 vs 59.6 years, p = 0.018), had tumors less accessible to surgical resection (11% vs 39%, p < 0.001), and had more frequently an *EGFR* amplified (59% vs 30%, p < 0.001) or TERTp-mutant glioma (96% vs 81%, p < 0.001). Regarding radiological characteristics, compared to gyriform-infiltration-negative patients, gyriform-infiltration-positive patients were more in proportion to display multicentric or multifocal tumors (38% vs 15%, p < 0.001), which were more frequently associated with gliomatosis (64% vs 10%, p < 0.001), more frequently located in the insula (41% vs 25%, p = 0.014) and the thalamus (23% vs 6%, p < 0.001), and displayed more frequently poorly delimited edges (91% vs 72%, p = 0.002; *Table 2*).

Diagnostic value

The Sp, Se, PPV, and NPV of the presence of a gyriform infiltration for the diagnosis of molecular GBM were 90%, 52%, 29%, and 96%, respectively, in the whole series, and 97%, 48%, 85%, and 85%, respectively, among patients presenting non-enhancing tumors (n=95). In the whole series, the Sp, Se, PPV, and NPV of the presence of a gyriform infiltration for the diagnosis of an IDH-wild-type TERTp-mutant glioma were 97%, 15%, 96%, and 19%, respectively. In the subgroup of patients presenting non-enhancing tumors, all the tumors displaying the gyriform infiltration sign (n=13) were aggressive grade IV TERTp-mutant gliomas (11 molecular GBM and 2 IDH-wild-type GBM).

Impact of gyriform infiltration on outcome

At the time of analysis, 187 (44%) patients had died. The median follow-up duration was 14.9 months and the median overall survival (OS) was 23.2 months in the entire cohort. There was a significant difference in the median OS between patients with gyriform-infiltration-positive compared to gyriform-infiltration-negative diffuse gliomas (16.9 vs 25.6 months, p = 0.005). This difference was also

significant in the subgroup of patients with non-enhancing tumors (20.2 months vs not reached, p < 0.001; Fig. 3) but not maintained if only IDH-wild-type gliomas were considered. Also, the difference was not observed in the subgroup of patients with IDH-wild-type GBM and molecular GBM (Fig. 3c; 16.9 months vs 16.0 months, p = 0.07).

Discussion

Early identification of poor prognosis gliomas is of utmost importance to allow rapid diagnosis and treatment. Herein we showed that the presence of a gyriform infiltration is a specific imaging marker for molecular GBM and more generally for IDH-wild-type TERTp-mutant gliomas. To the best of our knowledge, our study is the first one to analyze the gyriform infiltration diagnostic value in gliomas.

Radiological presentation of molecular glioblastomas

Molecular GBM have been previously shown to have a radiological presentation different from that of IDH-mutant LGG and IDH-wild-type GBM (diagnosis based on the presence of microvascular proliferation or necrosis), characterized by frequent temporo-insular location, thalamic involvement, gliomatosis, and gyriform infiltration [4]. The analysis of the radiological presentation of molecular glioblastomas performed in the present study using an independent cohort validates these findings, which are also consistent with those of other studies showing a high frequency of temporoinsular location, thalamic involvement, and gliomatosis in molecular GBM [16–18]. However, except once [4], the presence of a gyriform infiltration has not been reported as a hallmark of molecular GBM elsewhere. Some authors have reported that nearly half of IDH-wild-type GBM displayed non-enhancing cortical signal abnormalities, defined as non-enhancing FLAIR hyperintensity in the cortex contiguous with the area of tumoral enhancement, and that these cortical abnormalities could be associated in 5-10% of cases with distant nonenhancing lesions [19, 20]. However, these abnormalities generally involved both the grey and white matter adjacent to the tumor, and are therefore different from the gyriform infiltration sign reported herein, which consists in an infiltration limited to the grey matter [21]. Gyriform infiltrations have been previously reported as present in a third of molecular GBM and 15% IDH-wild-type GBM, but virtually absent in IDH-mutant astrocytomas and IDH-mutant and 1p/19q codeleted oligodendrogliomas [4]. The present study validates these findings. Another characteristic of molecular GBM is that they frequently lack contrast enhancement, which can wrongly suggest a lower-grade glioma. In the present study, 74% of molecular GBM displayed no contrast

Fig. 3 Survival probability according to the presence of a gyriform infiltration. Kaplan-Meier overall survival curves for patients with (GI+) and without (GI-) gyriform infiltration in the entire cohort (a Log-rank test: 16.9 months vs 25.6 months, p = 0.005), in the subgroup of patients with non-enhancing lesions (b 20.2 months vs not reached, p < 0.001), and in the subgroup of patients with IDH-wild-type GBM and molecular GBM (c 16.9 months vs 16.0 months, p = 0.07)



enhancement, highlighting the need for imaging biomarkers to differentiate these tumors from actual low-grade gliomas.

Diagnostic and prognostic value of the gyriform infiltration

Herein we found that gyriform infiltration was a specific biomarker for molecular GBM that could be identified with a substantial inter-rater agreement, similar to that of the recently described T2-FLAIR mismatch sign in IDHmutant astrocytomas [9] or the Fluid attenuation sign in noncontrast-enhancing tumor, correlated with IDH glioblastoma [22]. We found that tumors displaying gyriform infiltration were more infiltrative and more frequently associated with gliomatosis likely explaining why patients with gyriform infiltration tumors more frequently underwent a biopsy than a surgical resection. However, differentiating gyriform infiltration from cortical and subcortical infiltration is not always easy on T2 FLAIR sequences, especially when the infiltration is located in the internal temporal lobe. In such cases, T2w coronal sequences may be more appropriate (see Supplementary Figure). A previous study showed that infiltration of grey matter (as patterns of non-contrast-enhancing tumor) in glioblastoma was correlated with the IDH1 mutation status [23]. In addition to its diagnostic value, we found that gyriform infiltration was associated with a poorer prognosis in the entire cohort and in the subgroup of patients presenting non-enhancing tumors. In this population, the gyriform infiltration sign could be a precious imaging marker for the early identification of non-enhancing gliomas with poor prognosis, not suitable to the wait-and-scan strategy that can be proposed for some LGG.

Consequences for radiotherapy planning

Currently, there is no recommendation regarding the inclusion of gyriform infiltration-and more generally remote unconnected and non-enhancing lesions-in the clinical target volume (CTV). One strategy could be to include these lesions and consider all FLAIR hypersignal in CTV, although it could lead to an important treatment volume. Another strategy could be to focus radiotherapy on the enhancing lesion which is the site of tumor recurrence in most cases [24, 25] and to consider that systemic treatment (temozolomide) will treat other distant non-enhancing lesions. However, in a recent study focusing on 12 patients with multicentric non-enhancing lesions in GBM, 12 of 16 identified non-enhancing lesions were included in the radiation field: during follow-up, 9 remained unchanged and 7 progressed (with the appearance of an enhancing component in 6 lesions)[20]. The 4 non-enhancing lesions not treated were all progressive within a short delay, appearing as aggressive enhancing lesions. These results, combined with ours showing a progression in 70% of cases of gyriform infiltration within a short delay, support the inclusion of gyriform infiltration in the treated volume. It could be an alternative option to surgery, which is very challenging in these cortical areas with a high risk of functional damages. When gyriform infiltration is associated with gliomatosis and multicentric lesions, a radiation protocol including the whole brain in association with temozolomide can be another strategy in selected patients [26, 27].

Limitations and perspectives

Our study is limited by its single-center design and by the fact that molecular GBM were only defined by the presence of a TERTp mutation and/or an EGFR amplification. There was no case only defined by the presence of a combined chromosome 7 gain and chromosome 10 loss. Although these cases are rare (about 10% of molecular GBM [3]), future studies are needed to determine whether these molecular GBM frequently present a gyriform infiltration. Additionally, some patients were not included in the analysis because of the lack of some MRI sequences (especially T2 FLAIR sequences). Confounding bias could be related to the prognosis value of gyriform infiltration, for example patients with tumors displaying gyriform infiltration were more in proportion to have had a biopsy rather than a complete surgery. Finally, we cannot exclude occurrence of a disease spectrum bias due to the retrospective design with missing data.

The strength of our study lies in the large sample size and in the blinded assessment of the gyriform infiltration sign. Nevertheless, future studies will be needed to validate our findings and to understand the pathophysiology of gyriform infiltration. Scherer has described invasion patterns in glioblastoma and defined "secondary structures" corresponding to mechanisms by which glioma cells spread from preexisting tissue elements [28]. We think that the gyriform infiltration could be explained by perineuronal satellitosis and surface (subpial) growth (2 of the 4 "secondary structures of Scherer"). There is probably microscopic communication between the principal tumor and this elective cortical infiltration that cannot be detected due to the imaging resolution. These hypotheses will need to be confirmed by a dedicated histopathological study.

Conclusion

An elective gyriform infiltration on T2 FLAIR MRI sequences constitutes a highly specific imaging marker of IDH-wild-type TERTp-mutant gliomas especially in the molecular GBM subgroup, with a good inter-rater agreement. This sign is associated with a poor prognosis and

could be helpful to guide the clinical decision preoperatively for patients with non-enhancing tumors.

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Data availability The datasets generated during and/or analysed during the current study are available from the corresponding author on reasonable request.

Declarations

Conflict of interest The authors report no conflict of interest.

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