MRI Characteristics of Autoimmune Encephalitis With Autoantibodies to GABAA Receptor
A Case Series

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Abstract

**Background and Objectives**
To characterize the clinical and neuroimaging phenotypes of patients with autoantibodies to γ-aminobutyric acid type A receptor (GABAA).R).

**Methods**
Ten patients with autoantibodies against GABAA.R from Huashan Hospital Autoimmune Encephalitis cohort were identified. We used MRI assessments and clinical examinations to summarize major clinical profile and visualize and quantify lesion distribution features. The relationship between clinical features, neuroimaging phenotypes, and topology of GABAA.R expression were further investigated.

**Results**
The median age at onset of 10 patients (8 male patients and 2 female patients) with anti-GABAA.R encephalitis was 41.5 years (range: 17–73 years). All patients had prominent seizures and multifocal spotted or confluent lesions involved in limbic, frontal, and temporal lobes on brain MRI. Bilateral but asymmetric lesions in cingulate gyri were observed in all patients. These involved lesions could change dynamically with immunotherapies and relapse. Distribution of patients’ brain MRI lesions was positively correlated with gene expression level of β3 subunit-containing GABAA.R (Spearman ρ = 0.864, p = 0.001), the main target of autoantibodies. According to topology of lesions, patients with anti-GABAA.R encephalitis could be classified into 2 clinical-radiological types: confluent type with bilateral confluent lesions involved in almost all limbic, frontal, and temporal lobes and spotted type with multiple scattered small-to-medium patchy lesions. Patients with confluent type exhibited worse clinical presentations and outcomes when compared with those with spotted type (maximum modified Rankin scale [mRS]: 5 [5–5] vs 3.5 [3–4], respectively, p = 0.008; follow-up mRS: 4 [2–6] vs 0.5 [0–1], respectively, p = 0.016).

**Discussion**
Anti-GABAA.R encephalitis has distinctive neuroimaging phenotype. Cingulate gyri were frequently involved in this disorder. The topology of lesions might be associated with the distribution of β3 subunit-containing GABAA.R and reflected patients’ disease severity and outcomes.

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Autoimmune encephalitis (AE) constitutes a large group of severe neurologic disorders associated with antibodies directed at neuronal synaptic receptors, ion channels, or cell surface proteins. Diagnosis of AE is important because an early diagnosis and prompt treatment with immunotherapy could dramatically improve patients’ outcomes. Detection of autoantibodies in serum and/or CSF could confirm the diagnosis. However, it usually takes several days or weeks before obtaining antibody testing results, which might delay the diagnosis. Moreover, many countries and regions have no access to antibody testing, especially for AE mediated by some rare antibodies. Therefore, to diagnose AE based on neuroimaging and clinical information is of great clinical significance.

Given the broadening spectrum of clinical presentations in AE, clinical phenotypic characterization focusing on clinical and neuroimaging features could help identifying precise subtyping of AE, thus achieving an early and accurate diagnosis. For example, central hypoventilation is one of the recognizable clinical phenotypes of anti-NMDAR encephalitis. More recently, comprehensive studies of leucine-rich glioma-inactivated (LGI)1 and contactin-associated protein-like (CASPR)2 antibodies–related diseases have revealed that the phenotype of faciobrachial dystonic seizures is highly specific for patients with LGI1 antibodies but not with CASPR2 antibodies. These studies have shed light on how a phenotypic study could facilitate the classification and diagnosis of different subtypes of AE.

AE associated with antibodies against γ-aminobutyric acid type A receptor (GABAAR) is 1 recently described disease entity. Of interest, some clinical and neuroimaging features of this disorder might make it different from other antibody-mediated encephalitis. However, anti-GABAAR encephalitis is an extremely rare disease. Since the first report in 2014, only approximately 50 cases were reported worldwide. A detailed clinical and neuroimaging description of this disorder was scarce, thus impeding an early diagnosis of this potentially treatable disorder based on disease phenotype.

In this study, we reported a case series of anti-GABAAR encephalitis in China. We investigated whether anti-GABAAR encephalitis exhibited recognizable clinical and neuroimaging phenotypes. We further examined the relationship between clinical features, neuroimaging phenotype, and gene expression pattern of GABAAR.

Methods

Patients
We enrolled 1,919 patients with suspected AE in Huashan Hospital, Fudan University, between January 2013 and February 2021. In 483 patients with neuronal cell surface antibodies, 10 were finally diagnosed with anti-GABAAR encephalitis. Patients’ clinical information was obtained from medical records and telephone interviews. All patients had at least 1 brain MRI scan, and their images were reviewed by a neurologist (X.J.C.) and neuroradiologist (S.G.C.). We used modified Rankin scale (mRS) to evaluate the severity of symptoms at acute stage. The outcome at the last follow-up was also assessed with mRS. An outcome was considered favorable with an mRS 0–2 according to the previous study. CT, B-mode ultrasound, or 18fluorodeoxyglucose PET was performed to detect underlying tumors.

Autoantibody Detection and Identification of Subunit-Binding Specificity With Cell-Based Assays
Previous studies demonstrated that α1, β3, and γ2 subunits of GABAAR were targets of autoantibodies in anti-GABAAR encephalitis. Therefore, we first used live HEK293T cells transfected with plasmids encoding human α1 (NM_000806.5), β3 (NM_000814.5), and γ2 (NM_198904.3) subunits of GABAAR (all were obtained from MiaoLing Plasmid Sharing Platform, Wuhan, China) in combination to screen whether patients’ serum and CSF samples had autoantibodies to GABAAR. Second, when a patient’s sample reacted with coexpression of α1β3γ2, we transfected α1, β3, or γ2 subunit individually to identify the subunit-binding specificity of autoantibodies from each patient. Because the γ2 subunit was not well expressed in the cell surface when transfected individually, autoantibodies to γ2 subunit was observed after fixation and permeabilization, as previously described.

HEK293T cells were transfected with an equivalent amount of indicated single α1, β3, or γ2 subunit of GABAAR or cotransfected with 3 subunits (ratio 2:2:1). Thirty-six hours after transfection, cells were used to detect autoantibodies. For a live cell-based assay (CBA), cells were washed 3 times with phosphate-buffered saline and then incubated with serum (1:10) or CSF (1:1) for 30 minutes at room temperature (RT). Cells were washed, fixed with 4% paraformaldehyde for 5 minutes, permeabilized with 0.2% Triton X-100 for 10

Glossary

AE = autoimmune encephalitis; AED = antiepileptic drug; CBA = cell-based assay; CBA = contactin-associated protein-like 2; FLAIR = fluid-attenuated inversion recovery; GABAAR = γ-aminobutyric acid type A receptor; GABAAR = γ-aminobutyric acid type B receptor; GAD = glutamic acid decarboxylase; HHV6 = human herpesvirus 6; ICU = intensive care unit; LGI1 = leucine-rich glioma-inactivated 1; MNI = Montreal Neurological Institute; MOGAD = myelin oligodendrocyte glycoprotein-antibody-associated disease; mRS = modified Rankin scale; RT = room temperature.
Detection of Coexisting Autoantibodies

Previous studies demonstrated that patients with anti-GABAergic encephalitis could have coexisting autoantibodies.7,8,12 Therefore, patients’ serum and/or CSF samples were tested for onconeuronal antibodies (Hu, Yo, Ri, amphiphysin, CRMP5, and Tr) and antibodies to NMDAR, LGI1, CASPR2, α-amino-3-hydroxy-5-methyl-4-isoxazolepropionic acid receptor, γ-aminobutyric acid type B receptor (GABAβR) with commercially available kits (Euroimmun, Lübeck, Germany). Serum glutamic acid decarboxylase (GAD) 65 antibodies were detected with ELISA according to the manufacturer’s instructions (RSR, Cardiff, United Kingdom). Only high serum level of GAD65 antibodies (>10,000 U/mL) was considered clinically relevant.

MRI Data Acquisition

All included participants received T2-weighted fluid-attenuated inversion recovery (FLAIR) scans to identify their brain lesions. Of note, although the included participants were diagnosed in Huashan Hospital, we also retrieved these neuroimaging data from other hospitals where patients first visited at the onset of disease. Therefore, some of the participants had neuroimaging from different scanners and sequences. To visualize the lesion distribution and quantify regional volumes anatomically, we used the FLAIR scan with largest lesion volume for each case. FLAIR parameters with largest lesion volume for each case are listed in eTable 1 (links.lww.com/NXI/A705). All patients’ lesion volumes on FLAIR scan at each time point are listed in eTable 2.

Classifications of Lesions on FLAIR Imaging

According to the lesion patterns on FLAIR scans, we classified patients into 2 clinical-radiological types: confluent and spotted. Specifically, we semiquantitatively defined the confluent type when (1) at least half of the lesion areas grew together spanning 2 or more lobes and were not discrete with visual inspection in a consensus meeting and (2) the largest diameter of a lesion was ≥3 cm. By contrast, spotted type was primarily defined when at least half of the lesion areas were small spotted or irregular patchy lesions with scattered distribution pattern and the largest lesion diameter was <3 cm.

Besides, for patients who did not meet the criteria of confluent type, they were also defined as spotted type. Patient 2 had only head CT at acute stage, i.e., largest lesion. Because brain lesion volumes between CT and MRI scan are not easily comparable, this patient was not classified and therefore excluded when comparing clinical features, treatment, and outcome between patients in confluent type and spotted type group.

Quantification and Probability Map of Lesions on FLAIR Imaging

To increase the accuracy of subsequent lesion segmentation, FLAIR images were first skull-stripped using brain extraction tool to remove any nonbrain tissue component.13 Visible lesions on skull-stripped FLAIR scans were manually segmented by using the interactive software program ITK-SNAP14 blinded to the clinical data, followed by a consensus meeting when the disagreement occurred. A binary lesion mask for each participant was obtained.

To capture the spatial pattern of brain lesion, we generated the probabilistic lesion mapping, which is a widely used approach to visualize lesion distribution and frequency of lesion occurrence in a given brain location.15-17 Because high-resolution T1-weighted scans were not available for all participants, we used the FLAIR template provided online (brainder.org/download/flare/) as the reference image to register native space FLAIR images in Montreal Neurological Institute (MNI) standard space. This nonlinear spatial normalization was performed by the antsRegistrationSyN.sh tool,18 part of the Advanced Normalization Tools software package (snavagithub.io/ANTS/). First, we registered binarized lesion masks in MNI space for each patient. Probabilistic lesion mapping was generated by first merging and then averaging all the lesion masks onto MNI template, and it indicated the percentage of patients who had lesions localized in a certain brain region. The registration results were visually inspected, and manual adjustments were made in case of minor displacements.

GABAergic Gene Expression Data and Brain Lesion Analysis

Previous studies demonstrated patients’ autoantibodies could bind to α1, β3, or γ2 subunit of GABAβR, but β3 subunit was the most frequent subunit that could be recognized by autoantibodies.8 We hypothesized that the distribution of patients’ MRI lesions could be related to the topology of GABAβR expression across the brain. Therefore, we first visualized the gene expression level of α1, β3, and γ2 subunits (GABRA1, GABRB3, and GABRG2), as previously described.19 In brief, complete microarray gene expression data sets were downloaded from the Allen Institute of Brain (human.brain-map.org/static/download).20 These data were originally obtained from 6 individuals. Gene expression was assayed by the Allen Institute with custom-designed Agilent arrays. Block-face images of tissue slabs were used to map samples of gene expression to anatomical locations. Before brain dissection, whole brain MRI images were obtained for each brain. The
<table>
<thead>
<tr>
<th>Patient no.</th>
<th>Age/sex</th>
<th>Prodromal symptoms</th>
<th>Onset symptom</th>
<th>Main clinical symptoms</th>
<th>Maximum mRS</th>
<th>ICU admission</th>
<th>Types of GABA&lt;sub&gt;R&lt;/sub&gt; subunit-binding specificity</th>
<th>CSF</th>
<th>EEG</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>17/M</td>
<td>Fever</td>
<td>Memory impairment</td>
<td>FS, GS, SE, cognitive deficits, irritability, psychosis, movement disorder, speech dysfunction, hearing loss, weakness and numbness in right limbs, and coma</td>
<td>5</td>
<td>Yes</td>
<td>Type 1 α1, β3, γ2</td>
<td>Elevated protein</td>
<td>Diffuse slowing, bilateral with R-dominant epileptiform activity</td>
</tr>
<tr>
<td>2</td>
<td>29/M</td>
<td>Excessive sweating, malaise</td>
<td>Seizure</td>
<td>FS, GS, SE, cognitive deficits, irritability, mood and behavioral change, speech dysfunction, autonomic dysfunction, muscle cramps, and coma</td>
<td>5</td>
<td>Yes</td>
<td>NA&lt;sup&gt;b&lt;/sup&gt; α1, β3</td>
<td>Normal</td>
<td>Bilateral with R-dominant slowing and epileptiform activity</td>
</tr>
<tr>
<td>3</td>
<td>68/M</td>
<td>None</td>
<td>Memory impairment</td>
<td>FS, GS, SE, cognitive deficits, psychosis, and coma</td>
<td>5</td>
<td>Yes</td>
<td>Type 1 α1, β3</td>
<td>Normal</td>
<td>Diffuse slowing, bilateral epileptiform activity</td>
</tr>
<tr>
<td>4</td>
<td>73/M</td>
<td>None</td>
<td>Seizure</td>
<td>FS, GS, SE, cognitive deficits, irritability, psychosis, movement disorder, and coma</td>
<td>5</td>
<td>Yes</td>
<td>Type 1 α1, β3</td>
<td>Elevated protein, OCBs</td>
<td>Bilateral with L-dominant frontal focal slowing and epileptiform activity</td>
</tr>
<tr>
<td>5</td>
<td>57/F</td>
<td>HHV6 encephalitis</td>
<td>Seizure</td>
<td>FS, GS, SE, cognitive deficits, hallucinations, and coma</td>
<td>5</td>
<td>Yes</td>
<td>Type 1 α1, β3</td>
<td>Normal</td>
<td>Diffuse slowing and bilateral PED</td>
</tr>
<tr>
<td>6</td>
<td>66/M</td>
<td>None</td>
<td>Confusion</td>
<td>FS, GS, SE, cognitive deficits, confusion, hallucinations, personality change, sleep disorder, speech dysfunction, and coma</td>
<td>5</td>
<td>Yes</td>
<td>Type 1 α1, β3</td>
<td>Normal</td>
<td>Diffuse slowing and bilateral epileptiform activity</td>
</tr>
<tr>
<td>7</td>
<td>45/M</td>
<td>None</td>
<td>Seizure</td>
<td>FS, GS, cognitive deficits, emotional instability, and behavioral change</td>
<td>3</td>
<td>No</td>
<td>Type 2 α1, β3</td>
<td>Elevated protein</td>
<td>R hemisphere slowing and epileptiform activity</td>
</tr>
<tr>
<td>8</td>
<td>29/F</td>
<td>Headache, fever, vomiting</td>
<td>Seizure</td>
<td>FS, GS, SE, cognitive deficits, apathy, behavioral change, speech dysfunction, right arm movement disorder, sleep disorder, and confusion</td>
<td>4</td>
<td>No</td>
<td>Type 2 α1, β3</td>
<td>Pleiocytosis</td>
<td>L PED and diffuse slowing</td>
</tr>
<tr>
<td>9</td>
<td>38/M</td>
<td>Headache</td>
<td>Memory impairment, and confusion</td>
<td>FS, GS, SE, cognitive deficits, confusion, irritability, bizarre behavior, movement disorder, and speech dysfunction</td>
<td>4</td>
<td>No</td>
<td>Type 2 α1, β3</td>
<td>Elevated protein</td>
<td>Bilateral slowing and epileptiform activity</td>
</tr>
<tr>
<td>10</td>
<td>24/M</td>
<td>None</td>
<td>Seizure</td>
<td>FS, GS, cognitive deficits, irritability, behavioral change, and dyseusia</td>
<td>3</td>
<td>No</td>
<td>Type 2 α1, β3, γ2</td>
<td>Normal</td>
<td>Diffuse slowing</td>
</tr>
</tbody>
</table>

Abbreviations: FS = focal seizure; GABA<sub>R</sub> = γ-aminobutyric acid type A receptor; GS = generalized seizure; HHV6 = human herpesvirus 6; ICU = intensive care unit; mRS = modified Rankin scale; NA = not available; OCB = oligoclonal band; PED = periodic epileptiform discharges; SE = status epilepticus.

<sup>a</sup> According to the extent of lesions, brain MRI of patients were classified into 2 types: confluent type (type 1) and spotted type (type 2).

<sup>b</sup> Patient 2 was not classified because of lack of head MRI at acute stage, i.e., largest lesion.
MRI images were registered to the MNI152 space using FreeSurfer (help.brain-map.org/display/humanbrain/Documentation).20 The downloaded raw data (expression values and the MNI152 X, Y, and Z coordinates for each brain location) were processed with data reduction steps and further mapped into FreeSurfer space from MNI152 space. Last, values of gene expression were averaged across all voxels mapped into the Desikan-Killiany cortical atlas built into the FreeSurfer software for automatic labeling of regions of interest. These steps have been constructed into a fully established pipeline by French et al.19 They offered an R script to convert the gene expression data into 3D images. This script produces a gene-specific color lookup table for the FreeSurfer annotation files.

Moreover, the correlation between patients’ distribution of lesions and brain regional gene expression level was investigated, as previously described.21 In brief, gene expression levels of α1, β3, and γ2 subunits, which were expressed as z scores, were downloaded from the Allen Institute of Brain.20 The mean expression levels of 3 subunits in the limbic lobe, frontal lobe, temporal lobe, insular lobe, parietal lobe, occipital lobe, basal ganglia, thalamus, cerebellum, and brainstem were calculated. The proportions of patients who had lesions fell into these anatomical regions were also calculated. Finally, the relationship between gene expression level and proportion of patients who had MRI lesions in these brain regions was analyzed by the Spearman correlation analysis.

**Statistical Analysis**
Statistical analysis was performed with SPSS software, version 20 (IBM, Chicago, IL) and GraphPad Prism, version 6 (GraphPad Software, San Diego, CA). Data were presented as mean ± SD for normal distribution data and median (range) for non-normal distribution data. Normal distribution data were compared by the Student t test, and nonnormal distribution data were compared by Mann-Whitney U test. We used the Spearman correlation to examine the correlation
between gene expression level and distribution of MRI lesions. A 2-tailed \( p < 0.05 \) was considered as statistical significance.

**Standard Protocol Approvals, Registrations, and Patient Consents**

This study was approved by the institutional review board of Huashan Hospital, Fudan University. Written informed consent was obtained from each participant.

**Data Availability**

Anonymized data are available on reasonable request from any qualified investigator.

**Results**

**Clinical Characteristics**

We identified 10 patients (age 17–73 years, median 41.5 years; 8 male patients and 2 female patients) with anti-GABA\(_A\)R encephalitis from Huashan Hospital Autoimmune Encephalitis cohort (Table 1). GABA\(_A\)R autoantibodies were detected in both serum and CSF samples from all patients. Representative live CBAs of patients’ samples are shown in eFigure 1 (links.lww.com/NXI/A705). The flow diagram of diagnostic process is shown in eFigure 2.

Seizures were the dominant neurologic symptoms at the onset, which occurred in 6 patients. All patients experienced both focal and generalized convulsive seizures. Eight patients experienced status epilepticus. Six patients were admitted to intensive care unit (ICU) because of uncontrolled seizures. All patients also had cognitive deficits and psychiatric symptoms during the clinical course. Other clinical manifestations included decreased level of consciousness (8/10), speech dysfunction (5/10), movement disorder (4/10), sleep disorder (2/10), autonomic dysfunction (1/10), and peripheral nerve hyperexcitability (1/10). Demographics and clinical features of patients are summarized in Table 1.

**Neuroimaging Findings**

All patients presented with remarkable multiple or confluence cortical and subcortical lesions on T2-FLAIR imaging without evident mass effect, gadolinium enhancement, and restricted diffusion (Figure 1). No evidence of lactate accumulation or increased ratios of N-acetyl aspartate to choline is shown on proton magnetic resonance spectroscopy (data not shown).

According to the topology of lesions on FLAIR scans, we classified patients into 2 clinical-radiological types: confluent and spotted types. The confluent type was characterized by bilateral frontal, temporal, and limbic confluent lesion distribution and identified in 5 patients; whereas spotted type (in 4 patients) comprised multiple scattered small spotted or irregular patchy lesions. We quantified largest lesion volumes for each patient (except for patient 2) and found the mean lesion volume in confluent type group was much larger compared with that in the spotted lesion group (313 ± 114 cm\(^3\) vs 80 ± 19 cm\(^3\), \( p = 0.005 \)). Representative brain MRI scans of 2 types are shown in Figure 2. Compared with spotted type, patients with confluent type had more severe disease, reflected by maximum mRS (5 [5–5] vs 3.5 [3–4], \( p = 0.008 \)) and a higher rate of ICU admission (100% vs 0).

In our series, patients’ brain lesions frequently appeared in the limbic lobe (10/10), frontal lobe (10/10), and temporal lobe (10/10), but no patient had lesion in the cerebellum and brainstem (0/10) (eFigure 3, links.lww.com/NXI/A705). Lesion probability map also revealed that all lesions were distributed in supratentorial region but not likely in...
infratentorial regions (Figure 3). Of interest, all patients were featured by bilateral but asymmetric cingulate gyrus involvement (Figures 2 and 3).

Autoantibodies from all 10 patients in our cohort reacted to α1 and β3 subunits but only 2 reacted to γ2 subunit (Table 1 and eFigure 4, links.lww.com/NXI/A705). To explore whether the distinctive distribution of MRI lesions in anti-GABA_A encephalitis is related to the expression level of GABA_A (especially β3 subunit), we first visualized gene expressions of α1, β3, and γ2 subunits projected onto the FreeSurfer cortical regions. These results demonstrated that β3 subunit, but not α1 or γ2 subunit, was highly expressed in limbic, frontal, and temporal lobes (Figure 4). We also found z scored mean expression of β3 subunit gene (GABRB3) in supratentorial region (0.81 ± 0.45) was higher than that in infratentorial region (−0.74 ± 0.57) \( (p < 0.001) \). Meanwhile, a positive correlation was demonstrated between the expression level of GABRB3 and distribution of MRI lesions of 10 patients in each brain region (Spearman \( \rho = 0.864, p = 0.001 \)). However, neither α1 nor γ2 subunit expression level was correlated with the topology of MRI lesions (α1: Spearman \( \rho = 0.342, p = 0.334 \); γ2: Spearman \( \rho = −0.565, p = 0.089 \)).

Of interest, follow-up FLAIR scans demonstrated that the brain lesions of patient dynamically changed during the

**Figure 3** Lesion Probability Map of Anti-GABA_A Encephalitis

Note that limbic (especially bilateral anterior cingulate gyri, white arrow), frontal, and temporal lobes were dominantly involved. GABA_A = γ-aminobutyric acid type A receptor.

**Figure 4** Lateral and Medial Views of GABRA1, GABRB3, and GABRR2 Gene Expression Data Projected Onto the FreeSurfer Cortical Regions

Expression level was log2 transformed. Note the high expression of GABAB3 in limbic, frontal, and temporal lobes but a relative low expression in parietal and occipital lobes.
clinical course. In short-term follow-up (e.g., usually within 6 months after onset), some lesions diminished or disappeared, but some lesions could enlarge or new lesions could appear at the same time, although patients’ clinical status was stable or improved after treatment. In long-term follow-up (e.g., over 1 year after onset), brain lesions usually disappeared with clinical improvement. In 3 patients with relapses, new lesions always appeared in different brain regions. Long-term follow-up FLAIR scans (in 2 patients) at 3 and 4 years showed cortical atrophy and enlarged lateral ventricle. Representative follow-up scans are shown in Figure 1.

**Treatment and Outcome**

All patients received immunotherapies, of whom 7 had at least 2 kinds of first-line immunotherapeutic agents, whereas 2 received second-line immunotherapeutic agents. All patients received prescription with multiple antiepileptic drugs (AEDs). Notably, when immunotherapies were given to patients, seizures became controllable with reduced dosage of AEDs. The median follow-up time was 17.5 months (range 2–99 months). Eight patients became seizure free with oral AEDs at the last follow-up. Among 6 patients achieving favorable outcome, 4 had spotted lesion type. Two patients with confluent lesion type deceased, 1 died of severe pulmonary infection and the other died of complications of disseminated intravascular coagulation. Age, interval from disease onset to immunotherapy, proportion of underlying tumor, and follow-up time were not significantly different between patients in confluent type and spotted type group (all p > 0.05). However, patients with confluent type had poorer outcome than patients with spotted type (mRS: 4 [2–6] vs 0.5 [0–1], p = 0.016). Detailed treatment and outcome data are summarized in Table 2.

**Discussion**

AE associated with antibodies against GABA_{A}R was first described in 2014. In this study, we identified 10 patients with anti-GABA_{A}R encephalitis with a detailed description of clinical and neuroimaging phenotypes, thus providing a better understanding of this disorder: (1) Lesions in patients with anti-GABA_{A}R encephalitis had the dominant predilection site in supratentorial regions (i.e., limbic, frontal, and temporal), and lesion involvement in bilateral cingulate gyri was observed in all patients. This distinctive lesion distribution pattern could be positively associated with the expression level of β3 subunit–containing GABA_{A}R. (2) Anti-GABA_{A}R encephalitis could be classified into 2 topological types, namely confluent type and spotted type. Patients with confluent type had worse clinical course and poorer outcomes. (3) On temporal evolution, brain lesions of anti-GABA_{A}R encephalitis were dynamic during the clinical course, with diminishment after immunotherapies and new lesions during relapse.

In clinical practice, multifocal brain lesions in MRI could be caused by a variety of underlying etiologies, such as tumors,
denervating diseases, infections, and mitochondrial encephalomyopathy, lactic acidosis, and stroke-like episodes. As a consequence, correct differential diagnosis is challenging. Our study validated that multifocal cortical-subcortical MRI abnormalities were the core imaging features of anti-GABA\(_A\)R encephalitis.\(^7,8,12\) In fact, all patients in our study and most cases in previous reports presented with multifocal or confluent lesions on brain MRI.\(^{10}\) Therefore, anti-GABA\(_A\)R encephalitis should be considered as a possible etiology for this condition. By extensively reviewing cases from previous studies\(^7,9,22,23\) and quantitative analysis of lesions in our series, 3 distinctive imaging features of anti-GABA\(_A\)R encephalitis could be summarized. First, lesions of anti-GABA\(_A\)R encephalitis presented without mass effect, enhancement, or restricted diffusion. Second, these lesions were mainly distributed in supratentorial regions, and the predilection regions were limbic, frontal, and temporal lobes. Third, lesion in bilateral cingulate gyri could be a recognizable feature of anti-GABA\(_A\)R encephalitis because this sign was presented in all patients of this study and cases from previous reports.\(^7,9,22,23\) To the best of our knowledge, only 3 reported cases of myelin oligodendrocyte glycoprotein antibody–associated disease (MOGAD) showed similar lesions in cingulate gyri.\(^{24,25}\) However, lesions in anti-GABA\(_A\)R encephalitis had asymmetric distribution, more frequently involved in anterior cingulate gyrus, still making it different from MOGAD. In summary, the above-mentioned neuroimaging phenotype was a strong indicator of anti-GABA\(_A\)R encephalitis, which could be useful to distinguish this disorder from other mimics.

In view of our findings, it would be intuitive to raise the question about why anti-GABA\(_A\)R encephalitis presented a distinctive lesion distribution pattern. Therefore, we explore whether this phenomenon is related to the topology of GABA\(_A\)R expression. Indeed, our study found that in supratentorial regions where there were multiple brain MRI lesions, the mean \(z\) score of GABRB3 gene expression level was much higher compared with that in the infratentorial region (no MRI lesion). Moreover, the expression level of \(\beta_3\) subunit, but not \(\alpha_1\) or \(\gamma_2\) subunit, was positively correlated with the distribution of brain MRI lesions of patients. Subunit-binding specificity studies of previous reported cases\(^7,8,12\) and our cases demonstrated that 92.1% (35/38), 81.6% (31/38), and 28.9% (11/38) of the patients with anti-GABA\(_A\)R encephalitis had autoantibodies reacted to \(\beta_3\), \(\alpha_1\), and \(\gamma_2\) subunit, respectively (eFigure 5, links.lww.com/NXI/A705), indicating \(\beta_3\) subunit was the main subunit that could be recognized by autoantibodies of patients. This result further corroborated the close relationship between \(\beta_3\) subunit expression, autoantibody-binding preference, and MRI lesion distribution. Of interest, a trend of negative correlation between expression level of \(\gamma_2\) subunit and patients’ distribution of lesions was found (Spearman \(\rho = -0.565, p = 0.089\)), suggesting that lower \(\gamma_2\) subunit expression might correlate with lesion burden. However, this finding from our study was limited by small sample size. Future larger studies are warranted to confirm this observation. Taken together, our study demonstrated that cerebral lesion distribution pattern in anti-GABA\(_A\)R encephalitis could reflect the topology of \(\beta_3\) subunit–containing GABA\(_A\)R, thus providing a new insight into the distinctive lesion features of this disorder.

Five patients in our series showed extensive confluent lesions spanning 2 or more lobes. These patients had prominent seizures and decreased consciousness and needed an ICU admission. We classified these patients into confluent type group. However, the other 4 patients with multiple scattered spotted or patchy lesions had less severe symptoms and did not need an ICU admission. These patients were classified into spotted type group. In the long-term follow-up, patients with confluent type had poorer outcomes as well. Therefore, the extent of brain MRI lesions could be a potential useful marker of disease severity and prognosis. In light of the relationship between MRI lesion distribution, the topology of \(\beta_3\) subunit–containing GABA\(_A\)R, vanishing lesions after immunotherapies, and emergence of new lesions at relapse, we considered \(\beta_3\) subunit–containing GABA\(_A\)R might be dominantly affected in this disorder. Previous studies have found that autoantibodies could reduce the number of synaptic \(\beta_3\) subunit–containing GABA\(_A\)R clusters and decrease the current mediated by GABA\(_A\)R in cultured hippocampal neurons.\(^7,12\) However, further studies are warranted to clarify the relationship between the pathogenic effects of autoantibodies and imaging phenotype of multiple brain lesions and whether the extent of brain lesions could reflect the intensity of antibody-mediated pathogenic reaction.

Our study had several limitations. First, this was a retrospective, single-center study with relatively small sample size. Multivariable analysis or stratified analysis to identify possible confounders is confined by sample size. Further studies, preferably with larger sample size, are needed to validate the relationship between 2 types of MRI lesions and clinical outcomes. Second, we did not include children and could not compare the phenotypic difference between children and adults. Third, we did not quantify brain atrophy due to the lack of high-resolution T1 scans. High-resolution MRI data and advanced imaging techniques are needed to further elucidate the specific abnormal imaging phenotype and its association with clinical findings.

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**Disclosure**

The authors report no disclosures relevant to the manuscript. Go to Neurology.org/NN for full disclosures.

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## References

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In the Research Article “MRI Characteristics of Autoimmune Encephalitis With Autoantibodies to GABAA Receptor: A Case Series” by Deng et al.1, it should be noted that Shuguang Chu and Xiangjun Chen are co–senior authors. The authors regret the omission.

REFERENCE