

# Reproducibility of 2D GluCEST in healthy human volunteers at 7 T

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**Purpose:** To investigate the reproducibility of gray and white matter glutamate contrast of a brain slice among a small group of healthy volunteers by using the 2D single-slice glutamate CEST (GluCEST) imaging technique.

**Methods:** Six healthy volunteers were scanned multiple times for within-day and between-day reproducibility. One more volunteer was scanned for within-day reproducibility at 7T MRI. Glutamate CEST contrast measurements were calculated for within subjects and among the subjects and the coefficient of variations are reported.

**Results:** The GluCEST measurements were highly reproducible in the gray and white matter area of the brain slice, whether it was within-day or between-day with a coefficient of variation of less than 5%.

**Conclusion:** This preliminary study in a small group of healthy volunteers shows a high degree of reproducibility of GluCEST MRI in brain and holds promise for implementation in studying age-dependent changes in the brain.

## KEYWORDS

GluCEST, GluCEST reproducibility, glutamate

## 1 | INTRODUCTION

Glutamate is a major excitatory neurotransmitter and is involved in many aspects of brain function.<sup>1-3</sup> In addition, alterations of the glutamatergic system may contribute to the underlying pathophysiology of many neurological and neuropsychiatric disorders.<sup>4,5</sup> For example, elevated glutamate likely contributes to “excitotoxic” neuronal cell loss in Alzheimer’s disease through overstimulation of glutamatergic receptors.<sup>6-9</sup> Decreased glutamate is observed in animal model of Alzheimer’s disease and in patients with advanced

pathology,<sup>10-13</sup> particularly within the hippocampus, and is positively correlated with the progressive loss of glutamate transporter proteins in brain regions associated with memory and cognitive function.<sup>14,15</sup>

Although altered glutamate levels are implicated in the pathogenesis of neurological and neuropsychiatric diseases, glutamate has also been shown to decrease with age in the neurotypical brain.<sup>16-18</sup> To distinguish glutamate changes that may occur as a normal part of the aging from those associated with disease, it is vital to have a sensitive *in vivo* marker for brain glutamate. The ability to accurately monitor changes in

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glutamate will potentially provide better diagnostics and aid in the development of early therapeutic intervention.

Most published studies for measuring age-related changes in glutamate *in vivo* have used single-voxel proton MRS ( $^1\text{H}$  MRS) at field strengths of 4 T or less.<sup>16-19</sup> However, the utility of  $^1\text{H}$  MRS for accurately measuring glutamate is limited due to the severe overlap of glutamine resonances with glutamate along with contributions from macromolecules at field strengths of 4 T or less.<sup>19,20</sup> Recently there were 2 additional  $^1\text{H}$  MRS studies done at 7 T, in which the coefficient of variations for glutamate were reported to be 4 to 13%, whereas for glutamine it was between 10 and 33%.<sup>21,22</sup> With a recently developed imaging technique glutamate can be measured with higher sensitivity and at higher resolution by exploiting its CEST properties (glutamate-weighted CEST, or GluCEST) at 7 T.<sup>23-25</sup> Approximately 70 to 75% of GluCEST value has been shown based on theoretical as well as experimental studies is due to glutamate.<sup>23,26</sup> An advantage of this imaging method is that the glutamate measurements are devoid of any glutamine contamination. The GluCEST technique has been implemented in studies of mouse models of Alzheimer's disease, which demonstrated the sensitivity of the method to detect decreased concentrations of glutamate in diseased mice compared with healthy, age-matched controls.<sup>10,27,28</sup> In the mouse model of Parkinson's disease, increased GluCEST was measured from the striatum compared with the healthy controls.<sup>26,29</sup> This method was also used in studying the Huntington's disease mouse model in which the decreased GluCEST was measured in the striatal region of the brain.<sup>30</sup> This technique has also been demonstrated in a proof-of-principle study in human brain<sup>23</sup> and was recently used in small studies to show differences in brain glutamate levels between patients compared with healthy controls.<sup>25,31</sup>

The GluCEST tool is promising for assessing brain glutamate. However, before GluCEST can be implemented in larger clinical studies, the reproducibility of the measurement in healthy subjects must be established. This is a particularly important consideration, as neurotypical brains may also experience age-dependent changes in glutamate, which vary by sex and reproductive status of the individual.<sup>16-18</sup> The goal of the current work is to demonstrate reproducibility of GluCEST measurements in the brains of healthy volunteers. Here we use single-slice 2D GluCEST imaging to measure the reproducibility of glutamate in healthy human subjects at 7 T. For this initial reproducibility study of GluCEST in human brain, we performed multiple scans on 6 healthy volunteers over a maximum period of 8 months. We show that GluCEST MRI provides highly reproducible measurements in the human brain, both within individual subjects and across a small group of healthy adults.

Future work will focus on showing age-dependent changes and possible effects of sex and reproductive status

on GluCEST in healthy subjects. Ultimately, this work will provide a foundation for future clinical studies seeking to show glutamate alterations in patient populations.

## 2 | METHODS

All human studies were conducted under an approved University of Pennsylvania Institutional Review Board protocol. Written, informed consent was obtained from each volunteer. Seven healthy volunteers (6 males, 1 female) aged 28 to 66 years old ( $45 \pm 14.54$  years) participated in the study. Imaging was performed on a 7T Siemens scanner (Erlangen, Germany) with a Siemens volume coil transmit/32-channel receive proton head phased-array coil. Four scans were acquired on each subject ( $n = 6$ ). Two of these scans were acquired on the same day for assessment of within-day repeatability. The time period to complete all scans ranged from 3.5 months to 8 months (Table 1). For 3 of the subjects, 1 additional scan was acquired. A seventh subject was also included and had 2 scans acquired on the same day. This subject is included only in the within-day results.

To accurately obtain consistent slice locations between imaging sessions, we used a coregistration program, ImScribe (accessible at <https://www.med.upenn.edu/cmroi/imscribe.html>). It uses high-resolution  $T_1$ -weighted images and the slice information from a subject's initial scan as a target template for subsequent scans and performs affine coregistration that provides information for slice placement as illustrated in Supporting Information Figure S1.

For each subject, an axial slice located about 3 to 4 mm above the corpus callosum for adequate gray/white matter separation was selected. For every subsequent scan, ImScribe was used to identify slice coordinates to maintain consistent slice positioning. The 2D GluCEST imaging parameters were slice thickness = 5 mm, in-plane resolution =  $1 \times 1 \text{ mm}^2$ , matrix size =  $256 \times 256$ , gradient-echo readout TR = 7.4 ms, TE = 3.5 ms, read-out flip angle =  $10^\circ$ , averages = 2, SHOT TR = 8000 ms, shots per slice = 1, and a saturation pulse of  $B_{1\text{rms}} = 3.06 \mu\text{T}$  with 800-ms-long saturation pulse train consisting of a series of 99.8-ms Hanning-windowed saturation pulses with a 0.2-ms interpulse delay (100-ms pulse train). Saturation flip angle was  $3772^\circ$ . The CEST images were acquired at varying saturation offset frequencies from  $\pm 1.8$  to  $\pm 4.2$  ppm (relative to the water resonance set to 0 ppm) with a step size of 0.2 ppm. For gray and white matter segmentation, we have also collected the data with these same parameters at offset frequencies of  $\pm 20$  ppm and  $\pm 100$  ppm to generate magnetization transfer ratio map. To compute  $B_0$  maps for correction of  $B_0$  field inhomogeneity, water saturation shift referencing images<sup>32</sup> were collected from  $\pm 0$  to  $\pm 1$  ppm (step size = 0.1 ppm) with a saturation pulse of  $B_{1\text{rms}} = 0.29 \mu\text{T}$  with 200-ms duration and imaging parameters identical to those used for CEST as described

**TABLE 1** Individual GM and WM GluCEST mean and SD for each scan along with the coefficient of variation from the mean and SD of all scans for each volunteer

Subject	Age (years), gender	GM		GM		GM		WM		WM		
		GluCEST (% asymmetry) <sup>a</sup>	mean	GluCEST SD	GluCEST mean	GluCEST SD	GluCEST COV (%)	GluCEST (% asymmetry) <sup>a</sup>	GluCEST mean	GluCEST SD	GluCEST COV (%)	
1	26, M	Day 1	7.87 ± 1.36	7.92	0.26	0.26	3.27	Day 1	4.83 ± 1.91	4.59	0.17	3.74
		Week 3	8.3 ± 1.17					Week 3	4.55 ± 1.75			
		Month 6 <sup>b</sup>	7.72 ± 1.34					Month 6	4.42 ± 1.76			
		Month 6 <sup>b</sup>	7.8 ± 1.36					Month 6	4.57 ± 1.89			
		Day 1	7.7 ± 1.26	7.73	0.24	0.24	3.13	Day 1	4.41 ± 1.76	4.39	0.11	2.6
		Week 2	8.04 ± 1.32					Week 2	4.54 ± 1.82			
2	32, M	Month 3.5 <sup>b</sup>	7.45 ± 1.32					Month 3.5	4.3 ± 1.65			
		Month 3.5 <sup>b</sup>	7.72 ± 1.42					Month 3.5	4.3 ± 1.79			
		Day 1	7.54 ± 1.68	7.62	0.20	0.20	2.65	Day 1	4.46 ± 1.8	4.48	0.14	3.14
		Month 1.5	7.97 ± 2.16					Month 1.5	4.67 ± 2.07			
		Month 4 <sup>b</sup>	7.46 ± 2					Month 4	4.45 ± 2.08			
		Month 4 <sup>b</sup>	7.53 ± 1.86					Month 4	4.28 ± 1.89			
3	37, M	Month 8	7.59 ± 1.8					Month 8	4.52 ± 1.94			
		Day 1 <sup>b</sup>	7.49 ± 1.1	7.56	0.07	0.07	0.91	Day 1	4.37 ± 1.69	4.46	0.16	3.68
		Day 1 <sup>b</sup>	7.64 ± 1.4					Day 1	4.26 ± 1.68			
		Month 2	7.48 ± 1.24					Month 2	4.5 ± 1.68			
		Month 2	7.58 ± 1.19					Month 2	4.48 ± 1.69			
		Month 6	7.59 ± 1.25					Month 6	4.7 ± 1.78			
4	52, F	Day 1 <sup>b</sup>	7.81 ± 1.69	7.82	0.16	0.16	2.06	Day 1	4.84 ± 1.6	4.88	0.12	2.44
		Day 1 <sup>b</sup>	7.65 ± 1.2					Day 1	4.87 ± 1.55			
		Month 4.5	7.79 ± 1.48					Month 4.5	4.77 ± 1.54			
		Month 5	8.04 ± 1.38					Month 5	5.05 ± 1.62			
		Month 5	8.04 ± 1.38					Month 5	5.05 ± 1.62			
		Month 5	8.04 ± 1.38					Month 5	5.05 ± 1.62			

(Continues)

TABLE 1 (Continued)

Subject	Age (years), gender	GM			WM						
		GluCEST mean	GluCEST SD	GluCEST COV (%)	GluCEST mean	GluCEST SD	GluCEST COV (%)				
6	66, M	GM GluCEST ± SD (% asymmetry) <sup>a</sup>	7.46 ± 1.29	7.49	0.31	4.09	WM GluCEST (% asymmetry) <sup>a</sup>	5.05 ± 1.57	5.16	0.14	2.68
		Day 1					Day 1				
		Month 1	7.16 ± 1.15				Month 1	4.99 ± 1.41			
		Month 4	7.96 ± 1.26				Month 4	5.33 ± 1.66			
		Month 6 <sup>b</sup>	7.58 ± 1.03				Month 6	5.24 ± 1.53			
		Month 6 <sup>b</sup>	7.3 ± 1.87				Month 6	5.18 ± 1.75			
7	37, M	Day 1 <sup>b</sup>	7.64 ± 1.07	7.68	0.06	0.74	Day 1	4.55 ± 1.71	4.60	0.06	1.38
		Day 1 <sup>b</sup>	7.72 ± 1.45				Day 1	4.64 ± 1.81			

<sup>a</sup>The mean and SD values reported here were without applying any smoothing filter.

<sup>b</sup>Within-day scans.

COV, coefficient of variation

previously. A relative  $B_1$  map was generated from 2 images obtained using square preparation pulses with flip angles of  $30^\circ$  and  $60^\circ$ . The total acquisition time including the anatomical image, CEST images, and  $B_0$  and  $B_1$  field maps was approximately 12 minutes for each imaging session. The CEST contrast maps for the imaging slice were generated using in-house MATLAB (MathWorks, Natick, MA) routines as described by Cai et al.<sup>23</sup> Glutamate-weighted CEST image contrast values are reported in percent asymmetry. The  $B_0$ -corrected and  $B_1$ -corrected GluCEST contrast maps were averaged for each subject's entire gray matter (GM) and white matter (WM) to compute the coefficient of variation (COV) for the reproducibility studies. Gray and white matter segmentation was done on the same slice with the magnetization transfer ratio map, using a K-means cluster algorithm<sup>33</sup> with the number of segments set to 3 (gray, white, and CSF).

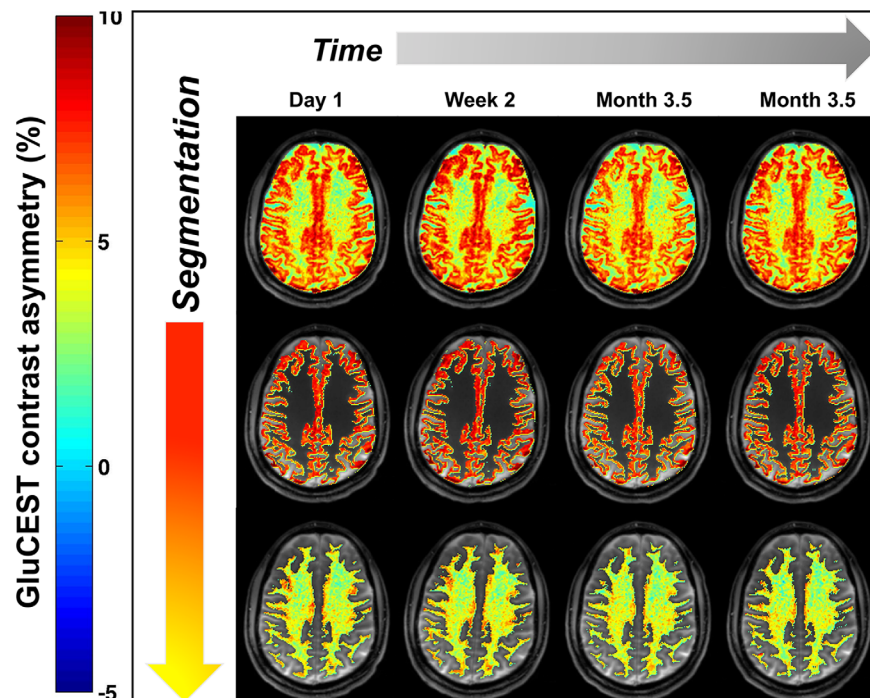
Gray matter and WM measures were also assessed between days and within days using linear mixed models to account for the dependency of the repeated measures. For the between-day models, GM and WM were modeled using time as a grouped discrete covariate in which days less than or equal to 50 were the first group, days 51 to less than or equal to 150 comprised the second, and the third group consisted of days greater than 150. This was done to help balance the sample as well as to prevent assuming a linear trend in the response over time, while conserving degrees of freedom. Repeated measures were adjusted for using subject-level random intercepts. For days in which there were 2 scans, only the first was included. The subject who only had 1 day with 2 scans was not included in the between-day analysis. For the within-day models, the session number (first or second) was included as a categorical covariate and repeated measures were also adjusted for using subject-level random intercepts. F-statistics were calculated using Satterthwaite approximations of degrees of freedom. All of the statistical analyses were performed on R v3.4.3 using the lme4 and lmerTest packages.<sup>34-36</sup>

### 3 | RESULTS

The  $B_0$ -corrected and  $B_1$ -corrected CEST maps for 1 of the subjects are shown in Figure 1 and for all subjects/time points are shown in Supporting Information Figure S2; the results for each subject and each scan are listed in Table 1.

#### 3.1 | Within-day results

The results for same-day scans are shown in Figure 2 ( $n = 7$ ). Subject 6 is included despite the presence of motion in their second scan of within-day, visible in the CEST maps (Supporting Information Figure S2). The GluCEST contrast



**FIGURE 1** Glutamate CEST (GluCEST) map of the entire slice (top) of a young healthy volunteer (32 years old) and segmentation of GluCEST map based on gray matter (center) and white matter (bottom) for within-day and between-day scans

was calculated for each slice acquired on the same day for each subject, and the GM and WM were segmented. For GM, the COV for within-day scans from the subjects was about 2% (with an individual subject range of 0.66% to 2.66%) and similar for WM (COV about 2% [0% to 2.75%]). The linear mixed models showed that there were no differences between the 2 within-day scans for both GM ( $F = 0.179$ ,  $P = .687$ ) and WM ( $F = 0.056$ ,  $P = .821$ ).

### 3.2 | Between-day results

For the between-day results, the first scan of the within-day and all of the other single scans from different days were used for calculating the COV. For GM, the COV for between-day scans from the subjects was about 4% (with an individual subject range of 0.81% to 4.39%, and similar for WM (COV of about 4% [2.24%–4.55%]). The linear mixed models showed that there were no differences between scans for both GM ( $F = 0.202$ ,  $P = .820$ ) and WM ( $F = 0.417$ ,  $P = .668$ ).

### 3.3 | Intrasubject scan results

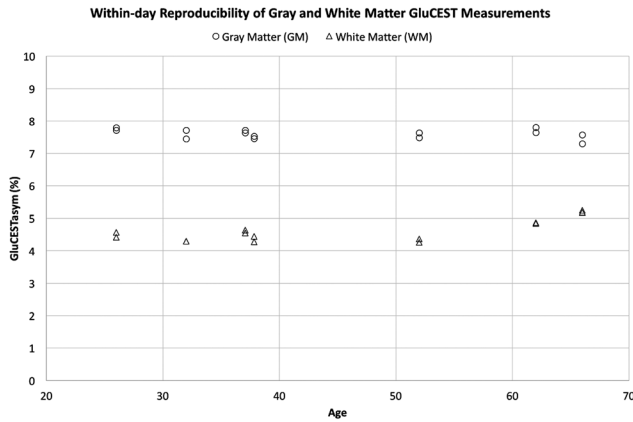
The GluCEST contrast values for each subject ( $n = 6$ ) were calculated for every scan (Figure 3) (either 4 or 5 scans, see Table 1), and the values for GM and WM were averaged to calculate each individual subject's COV. For GM, the individual COV ranged from 0.91% to 4.09%. For WM, the individual COV ranged from 2.44% to 3.74%.

### 3.4 | Intersubject scan results

The GluCEST group averages ( $n = 6$ ) were calculated for both GM ( $7.69 \pm 0.16$ ) and WM ( $4.66 \pm 0.3$ ). The intersubject COV was determined from these averages: GM COV = 2.15%, WM COV = 6.44%. Individual GluCEST values from each subject are given in Table 1. Modeling all GM measures for these 6 subjects by both within-day scan session as well as between-day time groups, the  $P$  values for session and time are .431 ( $F = 0.649$ ) and .852 ( $F = 0.161$ ). For WM, the  $P$  values for the effects of session and time are .142 ( $F = 2.355$ ) and .382 ( $F = 1.014$ ).

## 4 | DISCUSSION AND CONCLUSIONS

Glutamate CEST imaging has emerged as an ideal tool for use in neurological studies. This technique provides greatly enhanced spatial resolution compared with traditional spectroscopic methods, allowing for a whole-slice assessment of glutamate levels across varying regions of anatomy. To the best of our knowledge, this is the first human GluCEST study at 7 T that determines the reproducibility of the measurements. This study shows that GluCEST MRI provides highly reproducible measurements both within individual subjects and across a small group of individuals. Specifically, the COV was less than 5% in GM and WM regions of the brain for all subjects, both within and across days. Based on

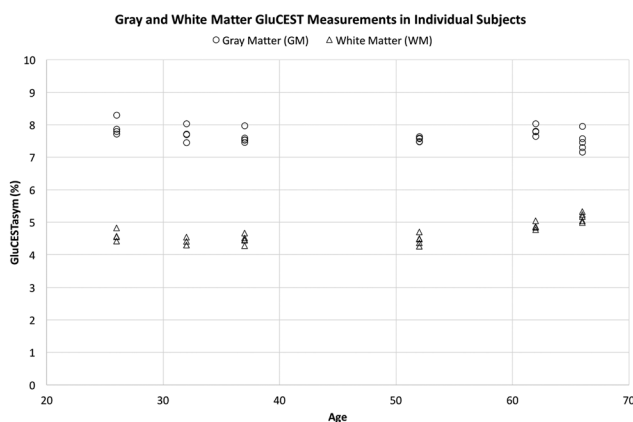


**FIGURE 2** Within-day GluCEST measurements of gray matter (GM, circles) and white matter (WM, triangles) for each individual subject

the small-reproducibility COV, this method is sensitive to detect any physiological changes above 5%.

The GluCEST technique has been used in studying the seizure foci in temporal lobe epileptic patients, and showed increased GluCEST contrast on the ipsilateral hippocampus compared with the contralateral hippocampus.<sup>25</sup> In addition, the GluCEST was shown to be a potential biomarker for patients on the psychosis spectrum.<sup>31</sup> In these studies, the GluCEST saturation pulse parameters used were slightly different than those used in the current study. Furthermore, the regions of the brain that were investigated in the prior studies were also different from those investigated in the current study. Accounting for these differences, the GluCEST results from this study are consistent with those reported in previous studies.

As previous work<sup>37</sup> has noted, there are likely several sources that contribute to variation in our measurements. These include motion artifact and  $B_1$  correction based on segmentation. We attempted to reduce motion artifact by using an approach that comfortably immobilized the head while the individuals were being scanned. However, we cannot rule out possible effects of motion. In the future, the



**FIGURE 3** GluCEST measurement of GM (circles) and WM (triangles) from all scans of each individual subject

inclusion of advanced, fast navigator sequences (e.g., EPI) collected throughout GluCEST acquisition will enable estimation of subject motion and correction during each scan. As our  $B_1$  correction used in this study depends on the accuracy of the GM and WM segmentation, we are working on a voxel-based  $B_1$  correction strategy. We acknowledge several other limitations of the current manuscript, including a limited sample to assess age or gender effects; however, we anticipate that future studies will systematically investigate these important questions.

Here, our initial attempt at establishing reproducibility demonstrates that GluCEST MRI provides highly reproducible measurements in human brain, both within individual subjects and across a small group of healthy adults. Additional work needs to establish the reproducibility of GluCEST within and across specific regions of the brain. Nonetheless, we anticipate that this assessment of reproducibility lays the groundwork for further age-dependent and sex-differences studies of glutamate, and for detecting subtle changes in glutamate in psychiatric and neurological disorders.

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## SUPPORTING INFORMATION

Additional Supporting Information may be found in the supporting information tab for this article.

**FIGURE S1** Snap shot of the Imscribe software showing the coregistration process of the template and target T<sub>1</sub>-weighted images along with the slice-of-interest information in the target space.

**FIGURE S2** The GluCEST maps of all scans (within-day and between-day) for each individual for the entire slice (top) and segmentation of GluCEST map based on gray (center) and white matter (bottom).

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