

# Multi-Echo Gradient-Echo Spin-Echo with Echo-Planar Imaging for simultaneous quantification of $T_2^*$ and $T_2$

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## Abstract

**Motivation:**  $T_2^*$  and  $T_2$  contrast-based sequences provide complementary information for MR applications. The Gradient-Echo and Spin-Echo (GESE) sequence has been developed to facilitate the simultaneous acquisition of  $T_2^*$  and  $T_2$ -weighted images.

**Objective:** Corroborate the results reported in existing studies that tested the GESE sequence with EPI readout in the myocardium, WM and GM.

**Methods:** The GESE sequence was implemented through pulse programming by modifying the standard implementation of the ME-GE with EPI readout. Two variations tested - with simple and with composite refocusing pulses. Phantom scans and in-vivo scans of two healthy volunteers were conducted. Subsequently, a pixel-wise least squares fit was performed for computing quantitative  $T_2^*$  and  $T_2$  maps, from which ROIs were defined and mean values were extracted.

**Results:** Results obtained with simple and composite refocusing pulses, were respectively: for WM  $T_2^*$ : ( $55 \pm 1$  ;  $52 \pm 2$ ) ms ,  $T_2$ : ( $64 \pm 2$  ;  $64 \pm 4$ ) ms , for GM ( $T_2^*$ :  $57 \pm 2$  ;  $56 \pm 3$ ) ms ,  $T_2$ : ( $78 \pm 5$ ;  $80 \pm 8$ ) ms and for Myocardium  $T_2^*$ : ( $25 \pm 10$  ,  $27 \pm 12$ ) ms;  $T_2$ : ( $34 \pm 14$ ,  $46 \pm 32$ ) ms.

**Conclusions:** The values of  $T_2^*$  and  $T_2$  using the GESE sequence with composite pulse, for the myocardium and WM were similar to the ones reported in literature and generated relative errors between 1.6 and 6.1%. For GM, both values were more distant from to the ones reported in literature, with relative errors of 12.0% and 12.7%.

**Keywords:** Gradient-Echo Spin-Echo, EPI readout,  $T_2^*$  mapping,  $T_2$  mapping, Quantitative MRI;

## 1. Introduction

Gradient-Echo (GE)  $T_2^*$ -weighted sequences are widely used in several Magnetic Resonance (MR) applications such as Susceptibility-Weighted (SW) imaging, Perfusion Weighted Imaging (PWI) and Functional Magnetic Resonance Imaging (fMRI) [7]. There are two types of magnetic inhomogeneities that lead to faster  $T_2^*$  decay: macroscopic inhomogeneities and microscopic inhomogeneities. Macroscopic inhomogeneities are constant within voxels and can be caused by air-tissue boundaries, metallic implants, or deoxyhemoglobin in small veins. Microscopic inhomogeneities change within a voxel and can arise from blood products, iron deposits, or paramagnetic contrast agents.

Besides its wide range of clinical applications, GE sequences provide more advantages, such as short acquisition times through the use of short

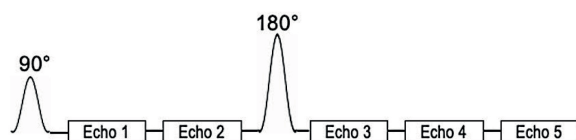
Echo Time (TE)s and valuable data for Quantitative Magnetic Resonance Imaging (qMRI) and more precise physiological and pathological evaluation of tissues and organs [15]. However, there are some limitations associated with GE sequences, such as low inherent Signal to Noise Ratio (SNR) due to the use of low Flip Angle (FA)s and higher sensitivity to  $B_0$  inhomogeneities.

$T_2$  contrast-based sequences can help assess the pathological status of the tissues, as it has been proven that the  $T_2$  relaxation times vary between healthy and pathological tissues [5]. An example of an MR application where  $T_2$ -based sequences are widely used is Cardiovascular Magnetic Resonance Imaging (CMR). Myocardial tissues with a high water content have higher  $T_2$  relaxation times (slower relaxation times) and therefore appear brighter in the images. As is known for myocardial tissues and other tissues, edema

is highly associated with acute injury.  $T_2$ -weighted sequences have also been useful for the pathological assessment of other anatomical regions, such as the brain, kidneys, and lungs.

It has been proven that  $T_2^*$  and  $T_2$  contrast-based sequences provide complementary information on MR applications such as Perfusion Susceptibility Imaging and CMR [12]. In Perfusion Susceptibility Imaging,  $T_2^*$ -based sequences provide relatively high Contrast to Noise Ratio (CNR) and sensitivity to macro vasculature diameters whereas  $T_2$ -based sequences provide lower sensitivity to contrast agent-induced signal changes but higher sensitivity to microvasculature changes. Therefore, the acquisition on both GE-PWI and Spin-Echo (SE)-PWI is often performed to improve diagnostic value [12]. Additionally, in CMR Blood Oxygenation Level-Dependent (BOLD) Magnetic Resonance Imaging (MRI) techniques can be used to assess myocardial oxygenation with no exogenous contrast agent. In this context, both  $T_2^*$  and  $T_2$  increase if there are higher concentrations of carbon dioxide in the blood that result in vasodilation. [6]. Hence, Gradient-Echo and Spin-Echo (GESE) sequence has been developed to facilitate the simultaneous acquisition of GE and SE [8, 10, 11, 13, 14, 16].

In summary, the GESE sequence, depicted in Figure 1, comprises the application of an excitation pulse, followed by two GE acquisitions and after the second acquisition a refocusing pulse is applied, followed by three mixed SE acquisitions. This sequence allows the simultaneous quantification of  $T_2^*$  and  $T_2$  through the data to the signal equation.



**Figure 1:** Sequence schematics of the GESE. The GESE acquires two GE echoes and three mixed SE echoes. Retrieved from [10].

To capture subtle Nuclear Magnetic Resonance (NMR) relaxation effects GESE has been implemented with Echo Planar Imaging (EPI) readout high temporal resolution. In order to reduce readout times and avoid geometric distortions due to  $T_2^*$  dephasing, Parallel Imaging (PI) techniques, such as SENSitivity Encoding (SENSE), can be combined with EPI [13].

### 1.1. Objectives

Given that very few reports have so far been published regarding the use of the GESE-EPI sequence for BOLD MRI techniques, this work has as

its principal objectives the expansion of the available body of literature and the corroboration of the results reported in existing studies that tested this sequence in the human brain and heart. In addition, this study aims to complement the previous development of GESE-EPI by its application in different MRI scanner.

Moreover, at the beginning of this study an additional objective was set, which consisted of implementing the GESE sequence with spiral readout. However, given the time restriction, this objective was dropped.

## 2. Methodology

Since the GESE sequence is not provided by the scanner's vendor (Philips), it was implemented through pulse programming by modifying the standard implementation of the multi-echo Fast-Field Echo (FFE) with EPI readout.

The FFE sequence consisted of a 90° excitation pulse with sinc profile, five-echo single-shot EPI readouts, and a Spectral Saturation with Inversion Recovery (SPIR) fat suppression pulse.

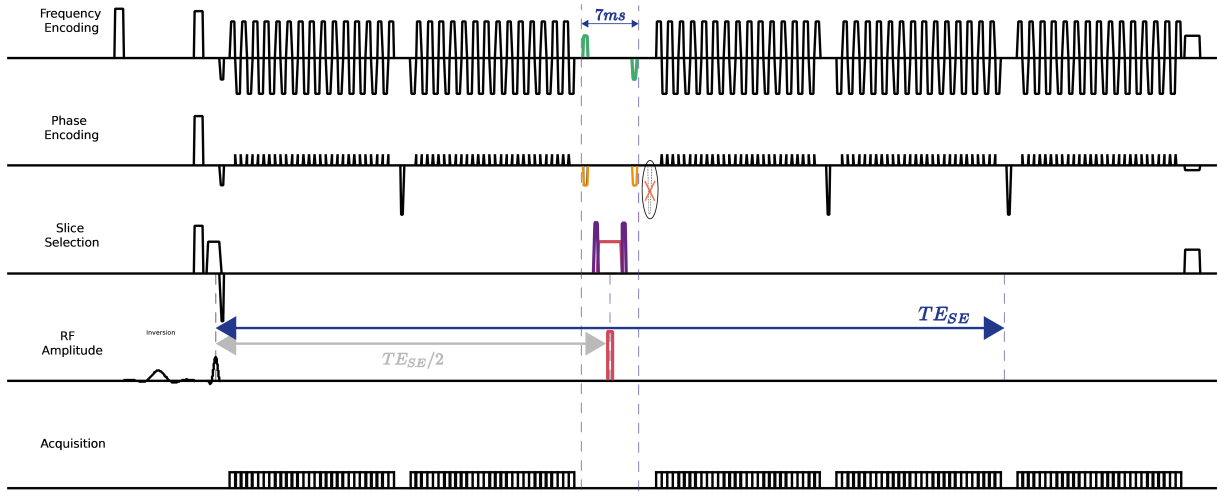
The modifications performed in the product implementation included the addition of a refocusing pulse between the second and third echoes, three slice selection gradients, two phase-encoding gradients, and two frequency encoding gradients, which will be further explained below. All these added elements increased the duration of the sequence by seven milliseconds. Additionally, one phase encoding gradient was removed. For debugging purposes and to test different combinations of sequence elements, the implementation allows the user to selectively choose which elements to include. The final implementation of the GESE sequence is shown in Figure 2, in which all modifications performed are colored.

### 2.1. Image Acquisition

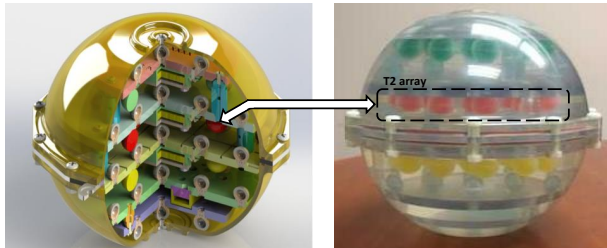
All scans acquired were performed on a 3.0T scanner (Ingenia, Philips, Best, The Netherlands)

#### 2.1.1 Phantom

Phantom imaging was performed using the International Society for Magnetic Resonance in Medicine (ISMRM)/National Institute of Standards and Technology (NIST) system phantom.



**Figure 2:** Sequence schematics of the five-echo GESE with EPI readout obtained through modifications performed to the standard implementation of the multi-echo FFE. The modifications include the addition of two frequency gradient blips (green), two phase gradient blips (yellow), two crusher gradients (purple), and a slice selection gradient and refocusing pulse (red). The yellow cross represents the removal of the first prephasing gradient along the phase encoding direction of the third echo acquisition. The added elements increased the sequence duration by 7 ms. The schematic was generated by the Philips' Graphical Viewer Environment (GVE).



**Figure 3:** ISMRM/NIST system phantom composed of three layers with sphere arrays filled with different concentrations of  $\text{NiCl}_2$ ,  $\text{MnCl}_2$  and  $\text{H}_2\text{O}$ , resulting in different values of  $T_1$ ,  $T_2$  and proton density. For this study only the  $T_2$  layer was used. Adapted from [2, 9]

**Table 1:** Imaging parameters used for the phantom experiments.

Imaging Parameters	GESE   FFE   SE	Ref. $T_2^*$	Ref. $T_2$
Technique	-	FFE	SE
N <sup>o</sup> of echoes	5	20	9
Fast Imaging mode	EPI	-	GRASE
Factor	EPI factor = 29	-	(Turbo Spin-Echo (TSE)+EPI)
Shot mode	single-shot	-	TSE factor = EPI factor = 9
SENSE factor	3	3	-
TE [ms]	10	1.19	7.9
$\Delta$ TE [ms]	19.4	1.5	-
FA	90°	50°	90°   180°
TR [ms]	10 000	600	1 000
Partial Fourier (Halfscan)	no	0.6	no

A 15 channel head coil was used for the phantom experiments. For all scans a single slice was selected with coronal orientation with the following imaging parameters: Field Of View (FOV) =  $270 \times 270 \text{ mm}^2$ , in-plane resolution =  $3 \times 3 \text{ mm}$ , and slice thickness =  $4 \text{ mm}$ . The TE for GESE, FFE, SE was set to the shortest value, of 10 ms, and Repetition Time (TR) was set to 10000 ms, to allow for recovery of the longitudinal magnetization. All sequences, with the exception of the  $T_2$  Reference sequence, were tested with SPIR fat suppression pulse. Additional imaging parameters are specified in Table 1. The proposed GESE sequence was tested with 2 different refocusing pulses: a 180y simple pulse or a 90y240x90y composite pulse.

### 2.1.2 Brain

The 15 channel head coil used in the phantom scans was also used for the brain scans. The sequences tested on the human brain were the proposed 2D five-echo GESE, FFE and SE. Due

to time constraints, no Reference  $T_2^*$  or  $T_2$  mapping sequences were acquired. A single-slice in transverse orientation was selected with: FOV =  $230 \times 230 \text{ mm}^2$ , in-plane resolution of  $2 \times 2 \text{ mm}$ , and slice thickness =  $3 \text{ mm}$ , EPI factor = 39, SENSE factor = 3 and with a SPIR fat suppression pulse. Additional acquisition parameters are shown in Table 2.

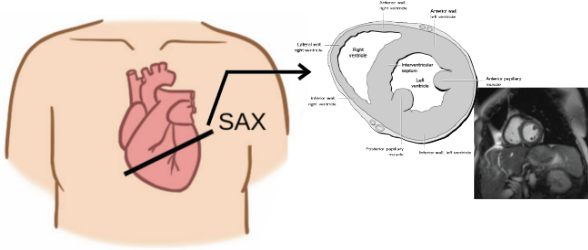
**Table 2:** Imaging parameters used for the brain experiments.

Imaging Parameters	GESE   FFE   SE
Fast Imaging mode	EPI
Shot mode	single-shot
N <sup>o</sup> of echoes	5
TE [ms]	14 (GESE and FFE); 31 (SE)
$\Delta$ TE [ms]	27.4
FA	90°
TR [ms]	3 000
Halfscan	no

### 2.1.3 Heart

Cardiac scans were performed using a 16-channel anterior coil and a 12-channel posterior coil. For all scans, a single slice was selected in the short axis plane (see Figure 4).

The five-echo GESE, FFE, SE, as well as, the Reference  $T_2^*$  sequence and  $T_2$  mapping sequences were acquired with the acquisition parameters shown in Table 3. For GESE, FFE and SE, a Spectrally Selective Attenuated Inversion Recovery (SPAIR) fat suppression pulse was selected. Additionally, for the Reference  $T_2^*$  and  $T_2$  mapping sequence, a black-blood prepulse was selected. All scans were set with a trigger delay of end diastole (903 ms) to minimize the cardiac motion.



**Figure 4:** Short axis view of the heart with a clear depiction of the left and right ventricles. Adapted from [1, 3, 4]

### 2.2. Image Post Processing

All post-processing steps were carried out in Matlab R2021a using original scripts unless mentioned otherwise.

The reference  $T_2^*$  and  $T_2$  maps were generated inline by the scanner and the GESE  $T_2^*$  and  $T_2$  maps were computed offline using a 4-parameter [6] pixel-by-pixel least squares fit, as follows:

$$S(\tau) = \begin{cases} S_0^I \cdot e^{-\tau \cdot R_2^*} & , 0 < \tau < TE_{SE}/2 \\ S_0^{II} \cdot e^{TE_{SE} \cdot (R_2^* - R_2)} \cdot e^{-\tau \cdot (2 \cdot R_2 - R_2^*)} & , TE_{SE}/2 < \tau < TE_{SE} \end{cases} \quad (1)$$

where  $S_0^I$  and  $S_0^{II}$  are the equilibrium signals before and after the refocusing pulse, respectively,  $\tau$  is the echo time and  $TE_{SE}$  is the time at which the spin-echo occurs.

For the phantom experiments, circular masks were created and mean  $T_2$  and  $T_2^*$  values for each vial were computed. Correlation and Bland-Altman plots were performed using the values obtained with the proposed five-echo GESE and the proposed Reference sequence. For the in-vivo scans, individual masks were computed for each sequence presented, to avoid incorrect Region Of Interest (ROI) positioning due to volunteer motion. The left-ventricular (heart) and white- and gray-matter (brain) masks were manually drawn using an in-house developed MATLAB script.

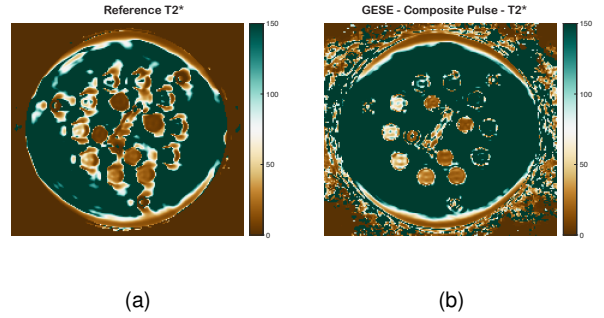
## 3. Results

### 3.1. Phantom

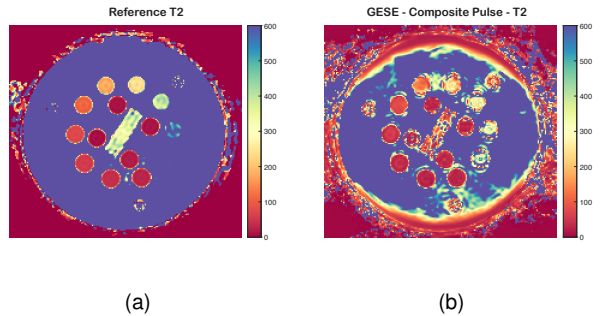
Figure 5(a) and 5(b) contain the  $T_2^*$  maps obtained with the reference and with the proposed GESE sequence, respectively. Figure 6(a) and 6(b) shows the  $T_2$  maps computed using the reference and the proposed GESE sequence, respectively.

To assess the accuracy of  $T_2^*$  and  $T_2$  parameter estimation using the GESE sequence with composite pulse,  $T_2^*$  (Figure 5(b)) and  $T_2$  (Figure 6(b)) maps were computed and compared with scanner generated  $T_2^*$  and  $T_2$  maps acquired with standard vendor sequences used as reference, shown in Figures 5(a) and 6(a).

From the maps of the GESE, Reference  $T_2^*$  and Reference  $T_2$  sequences, mean values for  $T_2^*$  and  $T_2$  inside each of the fourteen ROIs were computed. The values obtained are represented in the format of correlation plots, in Figures 7(a) and 8(a), for ease of comparison. Some outlier points, that correspond to phantom spheres with higher values of  $T_2^*$  and  $T_2$ , were excluded from this analysis, as they are not expected to be seen in-vivo. For outlier exclusion, a threshold of  $T_2^* > 100$  ms and  $T_2 > 200$  ms was established. The variable "n" displayed in the correlation plots represents the number of ROIs taken into account.



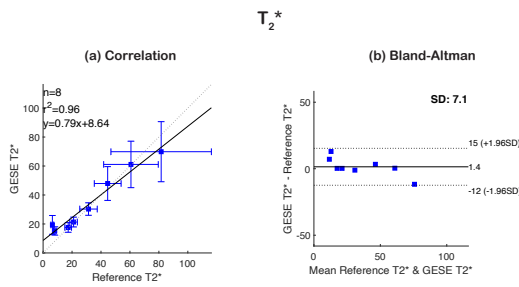
**Figure 5:**  $T_2^*$  maps computed from the (a) Reference  $T_2^*$  sequence and (b) GESE with composite pulse sequence for the phantom.



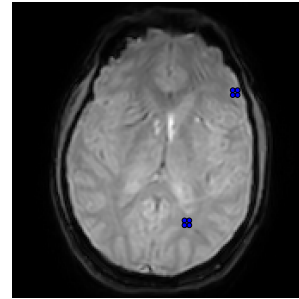
**Figure 6:**  $T_2$  maps computed from the (a) Reference  $T_2$  sequence and (b) GESE with composite pulse sequence for the phantom.

**Table 3:** Imaging parameters used for the heart experiments.

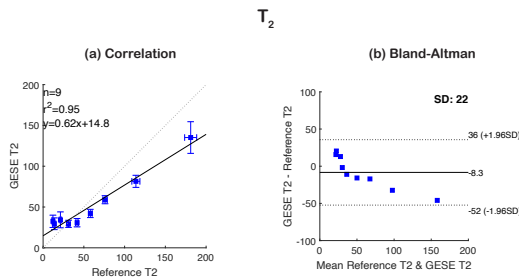
Imaging Parameters	GESE   FFE   SE	$T_2^*$ ref	$T_2$ ref
Technique	-	FFE	SE
FOV [ $mm^2$ ]	240 × 279	300 × 300	300 × 300
In-plane resolution [ $mm$ ]	3.1 × 3.1	3.6 × 3.6	3.1 × 3.1
Slice thickness [ $mm$ ]	8	10	8
Fast Imaging mode	EPI	Turbo Field Echo (TFE)	GRASE (TSE + EPI)
Factor	EPI factor = 31	8	TSE factor = 9   EPI factor = 8
Shot mode	single-shot	multi-shot	multi-shot
N° of echoes	5	15	9
TE [ms]	10	0.97	7.4
$\Delta TE$ [ms]	19.1	0.9	-
FA	90°	25°	90°   180°
TR [ms]	1 200 (1 beat)	15	1 200 (1 beat)
SENSE factor	3	2	2



**Figure 7:** (a) Correlation and (b) Bland-Altman plots comparing the mean  $T_2^*$  values obtained with the Reference  $T_2^*$  and GESE with composite pulse sequences. Only nine out of the fourteen ROIs were considered.



**Figure 9:** Representation of the two ROIs selected for analysis from the brain scans. One ROI corresponds to a grey matter region and the other corresponds to a white matter region.

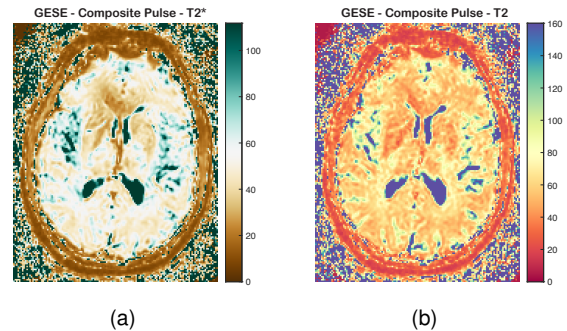


**Figure 8:** (a) Correlation and (b) Bland-Altman plots comparing the mean  $T_2$  values obtained with the Reference  $T_2$  and GESE with composite pulse sequences. Only thirteen out of the fourteen ROIs were considered.

### 3.2. Brain

For the analysis of the brain scans, two ROIs with circular shape were selected, white matter (ROI 1) and grey matter (ROI 2) regions. The ROIs are represented in Figure 9.

$T_2^*$  and  $T_2$  maps were computed from the scans obtained with the GESE sequence with both simple and composite pulses (for simplification, only the maps acquired with the composite pulse are shown, in Figures 10(a) and 10(b)), however, due to time restrictions, no reference  $T_2$  and  $T_2^*$  mapping scans were acquired. Instead, the reported values will be compared against available literature. The mean values for  $T_2^*$  and  $T_2$  inside both ROIs are shown on Table 4, as well as the values extracted from the literature.



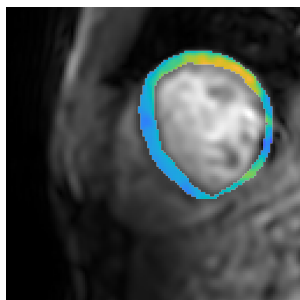
**Figure 10:** In vivo brain (a)  $T_2^*$  and (b)  $T_2$  maps computed from the GESE with composite pulse sequence.

**Table 4:** Mean  $T_2^*$  and  $T_2$  values measured for the two brain ROIs using the GESE with either simple or composite pulse.

Sequence	Tissue	$T_2^*$ (ms)	$T_2$ (ms)
		mean $\pm$ SD	mean $\pm$ SD
GESE - Simple Pulse	White matter (ROI 1)	55 $\pm$ 1	64 $\pm$ 2
	Gray matter (ROI 2)	57 $\pm$ 2	78 $\pm$ 5
GESE - Composite Pulse	White matter (ROI 1)	52 $\pm$ 2	64 $\pm$ 4
	Gray matter (ROI 2)	56 $\pm$ 3	80 $\pm$ 8

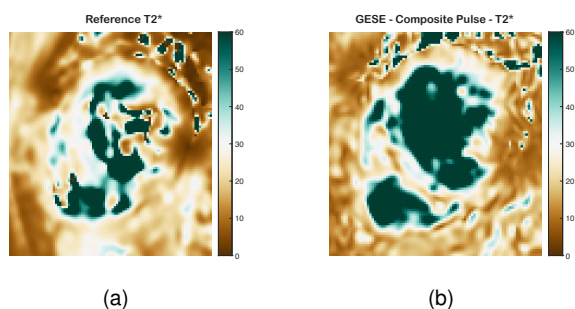
### 3.3. Heart

For the analysis of the cardiac scans, a single ROI was chosen within the myocardium, represented in Figure 11.

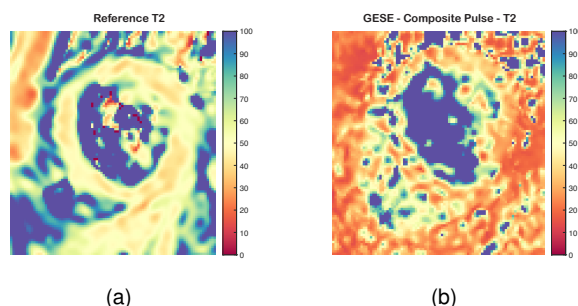


**Figure 11:** Representation of the ROI selected for analysis from the heart scans. The ROI corresponds to a region of the myocardium.

The  $T_2^*$  and  $T_2$  maps retrieved from the GESE with composite pulse and the reference sequences are shown in Figures 12 and 13 and the extracted mean ROI values are shown in Table 5.



**Figure 12:**  $T_2^*$  maps computed from the (a) Reference  $T_2^*$  sequence and (b) GESE with composite pulse sequence for the heart.



**Figure 13:**  $T_2$  maps computed from the (a) Reference  $T_2$  sequence and (b) GESE with composite pulse sequence for the heart.

**Table 5:** Mean  $T_2^*$  and  $T_2$  values measured for the heart ROI using the GESE with either simple or composite pulse.

Sequence	$T_2^*$ (ms)	$T_2$ (ms)
	mean $\pm$ SD	mean $\pm$ SD
Reference	24 $\pm$ 9	47 $\pm$ 7
GESE - Simple Pulse	25 $\pm$ 10	34 $\pm$ 14
GESE - Composite Pulse	27 $\pm$ 12	46 $\pm$ 32

## 4. Discussion

### 4.1. Phantom

The  $T_2^*$  Correlation (Figure 7(a)) plot shows a correlation coefficient of 0.96, indicating a strong linear association between the  $T_2^*$  obtained with GESE sequence and the  $T_2^*$  obtained with Reference  $T_2^*$  sequence. Five intermediate values, comprised roughly between a Reference  $T_2^*$  of 18 and 62 ms, are very close to the  $y=x$  equation (dashed line on the plot), which indicates strong proximity of values between both sequences. The Standard Deviation (SD) is visibly higher for the phantom spheres with higher values of  $T_2^*$  which indicates that the measurements are less reliable for high  $T_2^*$  values. Regarding the  $T_2^*$  Bland-Altman (Figure 7(b)) plot, all values are inside the limits of agreement.

For  $T_2$  even though there is a strong linear association between the  $T_2$  values obtained with the GESE sequence and those obtained with the Reference sequence, as supported by the  $r^2$  value of 0.95, this does not translate into the desired trend of having identical  $T_2$  values for both sequences. The obtained linear equation's slope of 0.62 and offset of 14.8 shows that it differs significantly from the ideal linear equation, whose slope would be 1 and offset would be 0.

The less satisfactory results obtained for  $T_2$  might be related to some limitations inherent to the GESE sequence. Since only 3 echoes with short echo spacing are used to estimate  $T_2$ , GESE sequence has limited capacity to estimate higher values of  $T_2$ .

### 4.2. Brain

After analysis of the results obtained for White matter (WM), it is possible to conclude that the GESE with composite pulse provided values of  $T_2^*$  and  $T_2$  were very similar to the values retrieved from literature using the same sequence, with relative errors of 6.1% and 1.6%, respectively. The values obtained deviated more from the literature values using other standard quantification sequences.

The values obtained for Gray matter (GM) generated higher relative errors when compared to literature using the same sequence. A possible cause might be related to inaccurate ROI position-

ing. However, the relative errors were reduced when compared to the values from literature using standard sequences.

#### 4.3. Heart

After a more detailed inspection of the results obtained for the myocardium, it is possible to conclude that the GESE sequence with composite pulse provided values of  $T_2^*$  and  $T_2$  very similar to the values retrieved from literature using the same sequence, with relative errors of 1.8% and 6%, respectively.

#### 5. Conclusions

Several limiting factors affected the results obtained, such as the reduced number of volunteers tested, which limits the comparison of the  $T_2^*$  and  $T_2$  values obtained with the GESE sequence versus the values retrieved from the literature. In addition, the limited number of echoes acquired limits the accuracy of the least-squares fit performed to obtain the quantitative maps. However, acquiring a higher number of echoes would compromise temporal resolution which could be a downside for applications where a temporal resolution is desired, such as CMR. Furthermore, the definition of ROIs was performed manually; therefore, a more robust technique should be used, such as automatic tissue segmentation, for a more precise selection of ROIs. In addition, because the  $TE_{SE}$  is not centered with any acquisition window, none of the echo images acquired is only  $T_2$ -weighted.

Further sequence optimization steps should be taken to improve the accuracy of  $T_2^*$  and  $T_2$  quantification using the developed sequence. First,  $TE_{SE}$  should be centered on the fifth acquisition window. This can be done by increasing the time interval between the second echo acquisition and the refocusing pulse. Second, different values of echo spacings ( $\Delta TE$ s) should be tested for a wider range of tissue quantification. Lastly, the design of the crusher gradient pair and the slice selection gradient should be improved and more designs of composite pulses should be tested to fully mitigate the problems arising from the mismatch of the slice profile.

Further research steps include testing the sequence on the brain and heart in an increased number of volunteers for a more sustained comparison of the results obtained with the values retrieved from literature. Additionally, breath hold experiments should be conducted to further validate the use of the GESE sequence for BOLD CMR techniques.

#### Acronyms

<b>NMR</b>	Nuclear Magnetic Resonance
<b>MRI</b>	Magnetic Resonance Imaging
<b>MR</b>	Magnetic Resonance
<b>FA</b>	Flip Angle
<b>BOLD</b>	Blood Oxygenation Level-Dependent
<b>SNR</b>	Signal to Noise Ratio
<b>CNR</b>	Contrast to Noise Ratio
<b>EPI</b>	Echo Planar Imaging
<b>GE</b>	Gradient-Echo
<b>FFE</b>	Fast-Field Echo
<b>TE</b>	Echo Time
<b>SE</b>	Spin-Echo
<b>GESE</b>	Gradient-Echo and Spin-Echo
<b>qMRI</b>	Quantitative Magnetic Resonance Imaging
<b>ISMRM</b>	International Society for Magnetic Resonance in Medicine
<b>NIST</b>	National Institute of Standards and Technology
<b>FOV</b>	Field Of View
<b>TR</b>	Repetition Time
<b>SENSE</b>	SENSitivity Encoding
<b>GVE</b>	Graphical Viewer Environment
<b>ROI</b>	Region Of Interest
<b>SD</b>	Standard Deviation
<b>SPIR</b>	Spectral Presaturation with Inversion Recovery
<b>SPAIR</b>	Spectrally Selective Attenuated Inversion Recovery
<b>TSE</b>	Turbo Spin-Echo
<b>TFE</b>	Turbo Field Echo
<b>SW</b>	Susceptibility-Weighted
<b>fMRI</b>	Functional Magnetic Resonance Imaging
<b>CMR</b>	Cardiovascular Magnetic Resonance Imaging
<b>PI</b>	Parallel Imaging
<b>PWI</b>	Perfusion Weighted Imaging
<b>WM</b>	White matter
<b>GM</b>	Gray matter

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