

Available online at www.sciencedirect.com



Journal of Magnetic Resonance 173 (2005) 169-174

JOURNAL OF Magnetic Resonance

www.elsevier.com/locate/jmr

# Increasing the speed of relaxometry-based compartmental analysis experiments in STEAM spectroscopy

Jack Knight-Scott\*, S. Andrea Dunham, Dattesh D. Shanbhag

Department of Biomedical Engineering, University of Virginia, Charlottesville, VA 22908, USA

Received 8 July 2004; revised 30 November 2004 Available online 20 January 2005

#### Abstract

In this work we present a method for improving the speed of spin-spin relaxation time  $(T_2)$  measurements for compartmental analysis in stimulated echo localized magnetic resonance spectroscopy without reducing the sampling density. The technique uses a progressive repetition time (TR) to compensate for echo time (TE) dependent variations in saturation effects that would otherwise modulate the received signal at short TRs. The method was validated in  $T_2$  studies on 10 young healthy subjects in spectroscopic voxels localized along either the right or left Sylvian fissure  $(2 \times 2 \times 1.5 \text{ cm}^3, 10 \text{ ms} \text{ mixing time (TM)}, 2048 \text{ data points, 819.2 ms}$ acquisition time). The TR was automatically adjusted so that TR-TM-TE/2 was kept constant as the TE was incremented. Compared to long TR  $T_2$  experiments, the progressive TR technique consistently replicated the  $T_2$  relaxation times and reference signals of the tissue water compartment while reducing the data acquisition time by more than 50%. The percent error was on average less than 2% for estimates of  $T_2$  and  $S_0$  for the tissue water, an indication that the progressive TR technique is a useful method for determining the tissue water signal for internal referencing.

© 2004 Elsevier Inc. All rights reserved.

Keywords: MRS; Transverse relaxation; T2; Spin-spin relaxation; Spin-lattice relaxation

### 1. Introduction

Voxel segmentation is an integral part of the quantitation process of single-voxel in vivo <sup>1</sup>H magnetic resonance spectroscopy (<sup>1</sup>H MRS) [1–3]. In <sup>1</sup>H MR brain spectroscopy, internal and external referencing schemes use spin–spin relaxometry to separate the volume contributions of tissue water and cerebral spinal fluid (CSF) in a localized volume-of-interest (VOI) to obtain a referencing signal for the calculation of the metabolite concentrations [1–7]. In internal schemes, the tissue water component serves as a reference, while in external schemes the CSF component serves as a measure of CSF partial volume. In the latter case, the CSF component is referenced to an external standard to calculate the tissue volume within the VOI. The presence of both CSF and tissue water within the VOI require a moderate-to-high sampling density to accurately separate the differential relaxation effects of the water compartments. In addition, the long spinlattice relaxation time  $(T_1)$  of CSF typically requires long repetition times to minimize the effects of magnetization saturation on the spin-spin relaxation time  $(T_2)$  measurements. These conditions necessitate sparse sampling of the  $T_2$  curve in clinical examinations to minimize patient scan times. Minimal sampling schemes have poor precision and accuracy [8]. In this work, we present a time efficient relaxation measurement technique that uses a variable repetition time (TR) and the acquisition of multiple relaxation curves to reduce the scan time for a stimulated echo (STEAM) spectroscopic  $T_2$  experiment while preserving a high sampling density.

<sup>\*</sup> Corresponding author. Fax: +1 434 982 3870.

E-mail address: jack.knight-scott@virginia.edu (J. Knight-Scott).

<sup>1090-7807/\$ -</sup> see front matter @ 2004 Elsevier Inc. All rights reserved. doi:10.1016/j.jmr.2004.12.001

## 2. Materials and methods

#### 2.1. A progressive repetition time

A common method for measuring  $T_2$  in in vivo <sup>1</sup>H MRS is the repetitive single-echo technique, where the TR is sufficiently long to allow the longitudinal magnetization to fully recovery between repetitions [9]. We will henceforth refer to this method as the fully relaxed  $T_2$  technique. The technique yields a signal  $S_{XY}$  from the transverse magnetization that varies as a function of the echo time (TE) as:

$$S_{XY}(\mathrm{TE}_n) = \sum_{m=1}^{M} S_{0\_m} \exp\left(-\frac{\mathrm{TE}_n}{T_{2\_m}}\right),\tag{1}$$

where the summation allows for M different relaxation compartments with various signal contributions  $S_{0_m}$ . In the spectroscopic brain model, M is generally restricted to two, CSF and tissue water [1–7]. If the TR is less than five times the  $T_1$  of CSF, then Eq. (1) must be modified to include  $T_1$  saturation effects. For the STEAM sequence, this inclusion yields the familiar equation (Fig. 1) [10–13]:



Fig. 1. Schematic of the stimulated echo localization sequence. The mixing time (TM) and delay time (TD) are kept constant throughout each progressive TR experiment. The time interval between the sets of concatenated gradients (positioned before and after the TM period) is kept constant as the TE increases to minimize TM-dependent diffusion-weighting. The concatenation of the refocusing gradient of the first slice-select gradient with the TE spoiler gradient is removed after the dead time during the first TE/2 interval becomes equal to or longer than the length of the RF pulse. This procedure minimizes the diffusion effects from the first slice-select gradient, especially at long echo times, since the rephasing gradient can be applied immediately following slice selection at long echo times.

$$S_{XY}(\mathrm{TE}_n) = \sum_{m=1}^{M} \frac{S_{0\_m}}{2} \left[ 1 - \exp\left(-\frac{\mathrm{TD}_{n-1}}{T_{1\_m}}\right) \right] \\ \times \exp\left(-\frac{\mathrm{TE}_n}{T_{2\_m}} - \frac{\mathrm{TM}}{T_{1\_m}}\right), \tag{2}$$

where  $TD_n = TR_n - TM - TE_n/2$ , and the intervals for TR, TM, and TE/2 are defined in Fig. 1. If Eq. (2) is to be employed for  $T_2$  measurements in a similar fashion as Eq. (1), then the following three conditions must be satisfied: (i) TM must be kept constant, (ii) diffusion-related signal attenuation must also be constant or negligible [14], and (iii) the focus of this work, TD must be kept constant by progressively changing the TR with the TE so that saturation effects are also kept constant. Thus representative  $T_2$  curves of Eqs. (1) and (2) will differ in intensity but not in decay rate.

Compared to the fully relaxed  $T_2$  technique, a relaxometry experiment with a short and progressive TR will yield estimates of the  $T_2$  and  $T_1$  weighted  $S_{0_m}$  values. To obtain the true reference  $S_{0_m}$  values needed in metabolite concentration calculations, two progressive TR relaxometry experiments with different TD values must be performed so that Eq. (2) can be employed. Although this requires two sets of data, because the minimum TR for a fully relaxed  $T_2$  experiment in the brain is 9 s or greater,  $T_2$  experiments with a progressive TR can be performed considerably faster.

#### 2.2. Relaxometry experiments

To validate the progressive TR technique, spectroscopic relaxometry data were collected from 10 healthy voung volunteers using STEAM (4 men/6 women, mean age:  $23.5 \pm 3.9$  years, range: 18–33 years). All experiments were performed on a 1.5 T Magnetom Sonata whole-body MRI system (Siemens Medical Systems, Iselin, NJ) using a standard Siemens circularly polarized proton head coil. Human studies were performed under protocols approved by the institutional review board and with the signed and informed consent of each participant. Each participant underwent a fully relaxed  $T_2$ experiment with TM/TR = 10 ms/11 s, a homospoil saturation recovery (HSR)  $T_1$  experiment [15], two progressive TR  $T_2$  experiments (Table 1), and then a repeat of the fully relaxed  $T_2$  experiment, in that order. The second fully relaxed  $T_2$  data were collected to determine whether a participant had significantly moved during the scan.

The HSR sequence was created by inserting a variable delay time between a three-pulse frequency-selective water suppression preparation sequence and the first slice-selective RF pulse in the STEAM sequence. The progressive TR  $T_2$  sequence was created by inserting variable delay times in the TE/2 periods and at the end of the sequence. The variable delay Download English Version:

# https://daneshyari.com/en/article/9587590

Download Persian Version:

https://daneshyari.com/article/9587590

Daneshyari.com