

# Measuring $T_2$ and $T_1$ simultaneously in the abdomen using $T_2$ -prepared bTFE at 3T

E. F. Cox<sup>1</sup>, C. L. Hoad<sup>1</sup>, and P. A. Gowland<sup>1</sup>

<sup>1</sup>Sir Peter Mansfield Magnetic Resonance Centre, School of Physics & Astronomy, University of Nottingham, Nottingham, Nottinghamshire, United Kingdom

## Introduction

A technique that could measure  $T_2$  and  $T_1$  simultaneously in the abdomen in an acceptable time would have significant benefits for investigating the gastrointestinal (GI) tract, e.g. to provide a quantitative measure of changes in tissue either due to normal function or to pathology. Changes in GI contents, such as dilution and viscosity changes, can also be monitored using  $T_1$  and  $T_2$  respectively<sup>1,2</sup>. However, it is often difficult to measure  $T_1$  and  $T_2$  accurately using the same sequence and in a reasonable time because of sensitivity to flip angle errors<sup>3,4</sup>. Images of high SNR with minimal distortions particularly in the abdomen can be produced using the balanced turbo field echo sequence (bTFE or TrueFISP). bTFE is particularly useful in the abdomen at 3T where EPI techniques are affected by susceptibility artifacts and HASTE or RARE techniques have a very high SAR. We have previously shown that it is possible to measure  $T_2$  using bTFE with a suitable preparation phase ( $T_2$ -prep bTFE)<sup>5</sup>. This work aims to show that  $T_1$  and  $T_2$  can be measured simultaneously using  $T_2$ -prepared bTFE by adjusting the sequence timing and bTFE flip angles, and hence to measure the transverse and longitudinal relaxation times of abdominal tissues at 3T in normal subjects.

## The $T_2$ -prepared bTFE sequence

The sequence consisted of a  $\{90^\circ\text{-TE}/2\text{-}180^\circ\text{-TE}/2\text{-}90^\circ\}$  pulse  $T_2$ -preparation phase followed by a half Fourier bTFE acquisition block consisting of a  $\alpha/2$  preparation pulse and successive alternating  $\pm \alpha^\circ$  pulses with a TR of 3ms. There was a time delay of 15 s at the end of the bTFE acquisition to allow the system to fully relax to maximize SNR and avoid having to take account of longitudinal suppression between shots in the fit. The signal intensity at the centre of k-space was measured for several TEs at 3 different bTFE flip angles. The fitting program modelled the signals by repeated application of rotation (1) and relaxation (2) functions:

$$M_x(t) = M_x(0) \quad M_y(t) = M_y(0)\cos(\alpha) + M_z(0)\sin(\alpha) \quad M_z(t) = -M_y(0)\sin(\alpha) + M_z(0)\cos(\alpha) \quad (1)$$

$$M_x(t) = M_x(0)\exp(-t/T_2) \quad M_y(t) = M_y(0)\exp(-t/T_2) \quad M_z(t) = M_z(0)\exp(-t/T_1) + M_0(1 - \exp(-t/T_1)) \quad (2)$$

(Eq. 2 does not include additional rotation due to off resonance which was not taken into account). The Powell algorithm<sup>6</sup> was used to minimise the difference between the measured and modelled data in order to fit  $T_2$ ,  $T_1$ ,  $M_0$  and  $\beta$ . The  $T_2$ -preparation pulses are non-ideal and so a  $\beta$  term was also included in the fit for experimental data as  $\{\beta \cdot 90^\circ - \text{TE}/2 - \beta \cdot 180^\circ - \text{TE}/2 - \beta \cdot 90^\circ\}$  for the  $T_2$ -preparation phase.

## Imaging Methods

Imaging was carried out on a 3T Philips Achieva whole body MRI scanner using the SENSE Torso coil. The  $T_2$ -prep bTFE sequence was implemented with half Fourier acquisitions to increase sensitivity to the preparation phase for both phantom and *in vivo* experiments. IR-EPI and SE-EPI were used as the gold standard techniques, since if they are run with a long TR they are insensitive to RF pulse errors for both  $T_1$  and  $T_2$  measurements.

**Calibration:** 2 spherical phantoms, divided into quadrants containing NaCl and 8 different concentrations of agar gel and Gadolinium ions were scanned using the  $T_2$ -prep bTFE sequence and both SE and IR single shot EPI sequences.  $T_2$ -prep bTFE decay curves were generated at  $\alpha=30^\circ$  using 19 different TEs (0.02-10s) and also at  $\alpha = 15^\circ$  &  $45^\circ$  at TE = 2s and 8s ( $T_2$ -prep bTFE imaging time = 6mins). The data was fitted using the minimization technique described above. A non-linear fit was used to measure  $T_1$  from the IR-EPI data and a linear log fit was used to measure  $T_2$  from the SE-EPI data, correcting for background noise in the lowest signal to noise images.

***In vivo:*** 5 healthy volunteers (24-44 years) were scanned on a single occasion and measurements of  $T_2$  and  $T_1$  in several abdominal tissues were acquired from 2 different slices (1 transverse, 1 coronal) through the abdomen.  $T_2$ -prep bTFE decay curves were generated at  $\alpha=30^\circ$  using 19 different TEs (0.02-10s) and also at  $\alpha = 10^\circ$  &  $20^\circ$  at TE = 2s and 8s ( $T_2$ -prep bTFE imaging time = 6mins). The same minimization technique was used to obtain  $T_2$  and  $T_1$  from the  $T_2$ -prep bTFE data. Regions were drawn in both kidneys, and in the liver and spleen in both the transverse and coronal plane where possible. It was not possible to distinguish between the cortex and the medulla in the kidney due to the slice chosen.

## Results

**Calibration:** Fig. 1 plots the measured  $T_2$  and  $T_1$  values from each quadrant of the phantoms from  $T_2$ -prep bTFE against EPI (SE and IR respectively).

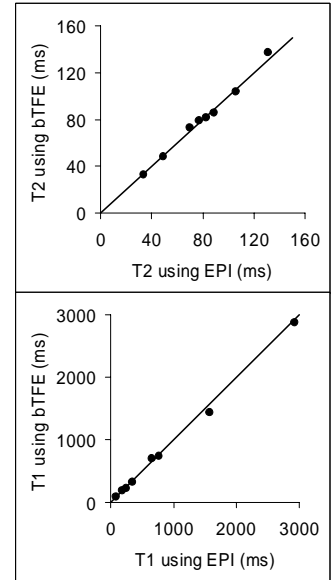
***In vivo:*** Fig. 2 shows a typical transverse image obtained for the shortest TE used (TE=20ms). Table 1 shows the mean  $T_2$  and  $T_1$  values measured in the 5 volunteers.

## Conclusions

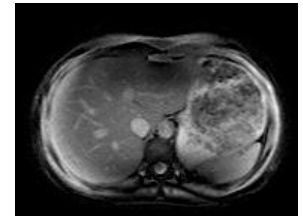
There was good agreement in the measured relaxation times between EPI and  $T_2$ -prep bTFE in the phantom and the *in vivo* data compares well with the literature<sup>7</sup>. By having a single imaging sequence to measure  $T_2$  and  $T_1$ , the analysis process is simplified and is more time efficient than using separate imaging sequences for the measurements. Further work will optimise the SNR per unit time in terms of both the number and values of TEs used to minimize the total imaging time. This sequence will be incorporated into GI studies where it is often important to be able to measure the dilution of contrast agents or food stuffs in the GI tract.

**References** 1. L Marciani *et al.* J. Nutr. **130**, 122-7 (2000), 2. CH Hoad *et al.* J. Nutr. **134**, 2293-300 (2004), 3. CA McKenzie *et al.* MRM. **41**, 208-12 (1999), 4. CS Poon *et al.* JMRI. **2**, 541-53 (1992), 5. CL Hoad. *et al.* Proc. ISMRM P2506 (2006), 6. WH Press *et al.* Num. Rec. in C (2nd Ed.), 7. CMJ de Bazelaire *et al.* Radiol. **230**, 652-9 (2004).

**Acknowledgements** This work was funded by the University of Nottingham and a grant from the MRC in partnership with EPSRC.



**Figure 1:** Phantom calibration for  $T_2$ -prep bTFE against single shot EPI (identity line shown)



**Figure 2:** Transverse slice using  $T_2$ -prep bTFE at TE=20ms

	$T_2$ (ms)	$T_1$ (ms)
Kidney	77 ± 7	1714 ± 257
Liver	32 ± 6	985 ± 143
Spleen	58 ± 12	1519 ± 160

**Table 1:**  $T_2$  and  $T_1$  (in ms) values for N=5 (mean ± std. dev)