

# Robust B<sub>1</sub>-Insensitive Whole-Brain T<sub>1</sub> Mapping with 3-TI MP-RAGE: Validation and Acquisition Strategy

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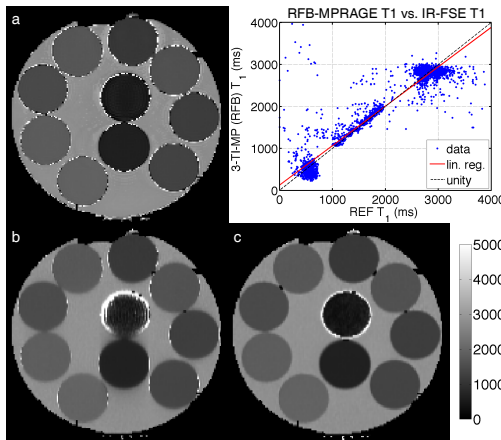
**Target audience:** Researchers, clinicians, and neuroscientists interested in robust volumetric high-resolution T<sub>1</sub> mapping.

**Purpose:** Fast T<sub>1</sub> mapping is potentially useful for segmentation of brain structures<sup>1,2</sup> and for myelin imaging<sup>3</sup>. Accurate, whole-brain, high-resolution T<sub>1</sub> maps have been obtained in monkeys at 7 T in a clinically relevant time, from 3 MPRAGE images with carefully selected inversion times (TI)<sup>4</sup>. This approach, which we will refer to as 3-TI-MP T<sub>1</sub> mapping, is free from B<sub>1</sub> heterogeneity effects, a particularly attractive feature for high field (≥ 3 T) applications. We implemented 3-TI-MP for human imaging at 7 T based on a MPRAGE sequence with 1D-centric (k<sub>z</sub>) ordering. We also implemented a 2D-centric (k<sub>y</sub>-k<sub>z</sub>) phase encode ordering scheme (radial fanbeam, or 2D-RFB<sup>5</sup>) to improve scan efficiency. In this work, we validated the method, and compared the accuracy and blur of the 3-TI-MP method for different k-space ordering and parallel imaging factors.

**Methods:** 3-TI-MP data were acquired using 3 serial MPRAGE scans with optimally selected TIs (= 150, 1280, 4000 ms). One k-space segment was acquired after each inversion pulse (inversion pulse spacing TS = TI + N\*TR + TD), using N=180-240 readouts, each at small flip angle (α = 5°) and short TR (= 7.7 ms), with other parameters as in Table 1. TS was held constant for the different TIs by altering the final delay TD; this removes dependence on M<sub>0</sub>, T<sub>2</sub><sup>\*</sup>, and B<sub>1</sub> and allows rapid T<sub>1</sub> estimation based on a simple lookup table<sup>4</sup>. All data were collected using a GE Discovery MR950 7 T scanner (GE Healthcare, Waukesha WI USA) with a 32-channel head coil (Nova Medical, Wilmington, MA USA). Data were collected in a multi-compartment phantom constructed using a range concentrations of MnCl<sub>2</sub> in 0.9 % saline solution, to provide T<sub>1</sub> values expected at 7 T in human brain (roughly 1000-5000 ms), and one peanut oil compartment. Three male volunteers (A, B, and C; ages 35, 35, and 31) were scanned each with a different k-space-ordering variant of the protocol, after providing informed consent. For all experiments, a reference T<sub>1</sub> map was acquired with a single-slice IR-FSE sequence, using four TIs (= 200, 600, 1500, 4000 ms), and freely available T<sub>1</sub> estimation software<sup>6</sup> that takes into account RF pulse imperfections and finite TR (= 5000 ms). 3-TI-MP T<sub>1</sub> lookups were performed offline without additional B<sub>1</sub> correction using either 1) scanner-reconstructed magnitude data with polarity restoration<sup>6</sup> or 2) coil-wise complex raw data as recommended by Liu *et al.*<sup>4</sup>

**Results:** T<sub>1</sub> estimation based on magnitude-data with polarity restoration was simple to compute and generally correct in white matter, but led to large errors in long-T<sub>1</sub> regions such as cortical grey matter near cerebrospinal fluid (data not shown), and for this reason was subsequently abandoned. The reference and coil-wise-complex 3-TI-MP T<sub>1</sub> maps agreed well in the phantom (*linear regression*: T<sub>1,MPRAGE</sub> = 0.97\*T<sub>1,REF</sub> + 66 ms, r = +0.98), and 2D-RFB ordering produced better quality T<sub>1</sub> maps than 1D-centric, as seen in Fig. 1. In volunteers, the correspondence of reference and 3-TI-MP T<sub>1</sub> maps was very good (r ≥ +0.78), and 2D-RFB resulted in maps with lower spatial variability (COV range=4-9% for 2D-RFB vs. 6-9% for 1D-centric) in a shorter scan time (Fig. 2).

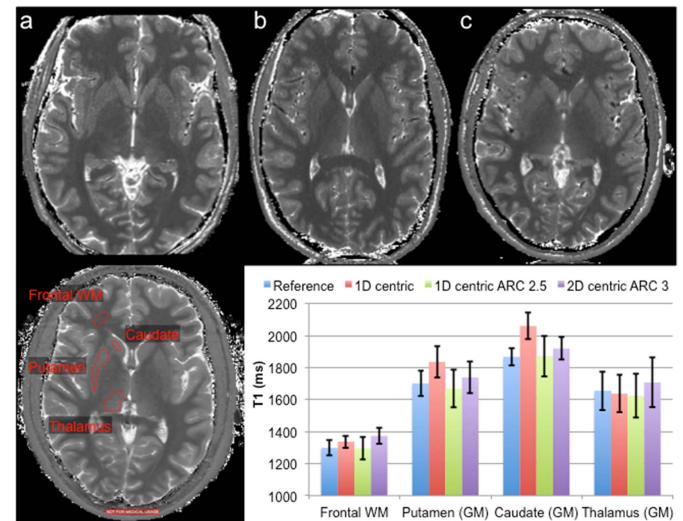
**Figure 1. Axial phantom T<sub>1</sub> maps from IR-FSE (a), and 3-TI-MP with 1D centric ordering (b) and 2D-RFB ordering (c). 3-TI-MP maps are reformatted in the axial plane to show blur. Note the reduced blur and better quality in (c). Upper right: 3-TI-MP T<sub>1</sub> from (c) is plotted vs. the reference T<sub>1</sub>.**



**Discussion and Conclusions:** This experimental demonstration of B<sub>1</sub>-insensitive 3-TI-MP whole-brain T<sub>1</sub> mapping at 7 T, validated against a reference technique in phantoms and *in vivo* human volunteers, demonstrates high accuracy and precision, and holds promise for future research and clinical applications. The proposed 2D-RFB k-space ordering scheme decouples readout train length from the slice dimension, extending the readout train and allowing for 2D acceleration, and reducing spatial blur in the k<sub>z</sub> direction without any use of k-space filtering.

**References:** [1] Deoni, SCL *et al.*, HBM 25:353–359(2005), [2] Bazin, PL *et al.*, OHBM 2013, Seattle, p.3384, [3] Mezer, A *et al.* Nat. Med. doi:10.1038/nm.3390 (2013) [4] Liu, JV *et al.*, NeuroImage 56:1154–1163 (2011) [5] Saranathan, M and Glockner, J, JMIR doi: 10.1002/jmri.24113 (2013). [6] Barral, JK *et al.*, MRM 64:1057-1067 (2010)

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**Figure 2. T<sub>1</sub> maps obtained using variants of 3-TI-MP: fully sampled 1D-centric (a), 1D-centric with ARC 2.5×1 (b), and 2D-RFB with ARC 3×1 (c). Bottom: T<sub>1</sub> values (mean ± std. dev.) in 4 manual ROIs (shown on left). Note the high quality of the 2D-RFB map in (c), and small variation in ROI T<sub>1</sub> values for the variant compared to 1D-centric with ARC 2.5.**