

## T<sub>1</sub> Mapping: Methods and Challenges

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**Purpose:** In this educational abstract, we provide an overview of the main T<sub>1</sub> mapping methods and we outline the challenges in performing quantitative T<sub>1</sub> measurement. We describe the gold standard (Inversion Recovery), as well as two widely used alternative methods (Look-Locker and Variable Flip Angle) that speed up the scanning and fitting procedures at the expense of accuracy and precision. The e-poster will include sample T<sub>1</sub> maps of phantoms and in-vivo human brains acquired with each of the above methods, and it will provide a list of useful T<sub>1</sub> mapping references.

### Outline of Content:

**Inversion Recovery (IR) T<sub>1</sub> Mapping:** This gold-standard method for T<sub>1</sub> mapping [1,2] consists of inverting the longitudinal magnetization and sampling the MR signal at several points (T<sub>1n</sub>) along its exponential recovery with a time constant T<sub>1</sub>. The IR pulse sequence is repeated N times, each time applying the same (typically adiabatic) inversion pulse, followed by different waiting times (T<sub>1n</sub>), and an imaging module that can be either spin echo (SE) or gradient echo (GRE). TR must be on the order of the longest measured T<sub>1</sub> to achieve sufficient magnetization recovery. The general equation used for the fitting procedure is given by:  $S_n = a + be^{-T_{1n}/T_1}$ , where a and b are complex-valued parameters and T<sub>1n</sub> is the inversion recovery time of the n<sup>th</sup> IR scan [3]. For precise and accurate measurement, it is recommended to perform at least four scans with TIs that span the range of expected T<sub>1</sub> values [3]. The gold-standard method does not assume a perfect inversion pulse, but it requires temperature monitoring as T<sub>1</sub> values change with temperature [4]. Additional simplifications can be made if TR>>T<sub>1</sub>, or by assuming specific values for  $\theta$  and  $\alpha$  (e.g., 180° and 90°, respectively).

**Look-Locker (LL) T<sub>1</sub> Mapping:** The Look-Locker sequence is similar to the gold standard scan in that it prepares the magnetization with an inversion pulse, but instead of a single sample of the recovery curve per TR it applies a train of N low flip angle pulses spread across the TR with spacing  $\tau$  [5]. The signal after the n<sup>th</sup> sampling pulse is given by:  $S_n = \beta (1 - DR e^{-n\tau/T_1^*})$  where

$$\beta = \frac{M_0(1 - e^{-\tau/T_1})}{(1 - \cos \alpha e^{-\tau/T_1}) \sin \alpha}, \quad DR = -\left(\frac{\cos \alpha (1 - [\cos \alpha e^{-\tau/T_1}]^{N-1})}{1 + \cos \alpha [\cos \alpha e^{-\tau/T_1}]^{N-1}} + 1\right) \quad \text{and} \quad T_1^* = \frac{\tau}{\tau/T_1 - \ln(\cos \alpha)}.$$

This model is sensitive to field inhomogeneity because it assumes perfect RF pulses of negligible duration and no lag between the RF pulse and the readout. The sensitivity to  $\alpha$  reduces as  $\tau/T_1$  increases, so spreading the sample points across TR improves accuracy.

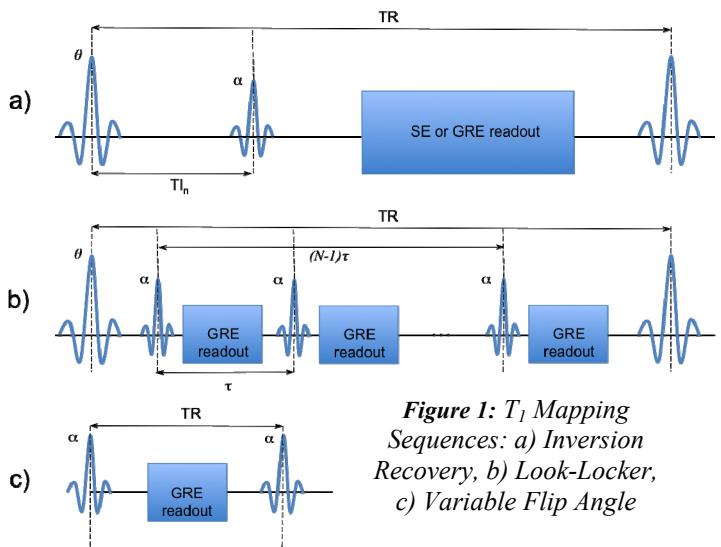
**Variable Flip Angle (VFA) T<sub>1</sub> Mapping:** This method can be used to acquire 3D T<sub>1</sub> maps in clinically feasible times [6, 7]. It utilizes two or more spoiled gradient-echo scans with varying flip angles. The equation describing the signal behavior in a spoiled gradient echo sequence is:  $S_n = \frac{PD(1 - e^{-TR/T_1}) \sin \alpha_n}{1 - \cos \alpha_n e^{-TR/T_1}}$ .

This equation assumes TR>T<sub>2</sub>\* and perfect RF spoiling. Additional noise assumptions can reduce the fitting routine to a weighted least-squares procedure [8]. As is the case for the two previous methods, the VFA method should not assume perfect knowledge of the flip angle  $\alpha$ . To account for B<sub>1</sub> inhomogeneities, a field map can be acquired along with the T<sub>1</sub> mapping scans.

**Summary:** We have outlined the basic pulse sequences and models for accurately mapping the T<sub>1</sub> relaxation time. Attention should be paid to the assumptions underlying any model simplifications, and it is always recommended to check a new method against the gold standard using simulations [9].

### References:

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- [7] Deoni et al., Magn Reson Med 53(1): 237-241 (2004)
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- [9] [mrsrl.stanford.edu/~jbarral/t1map.html](http://mrsrl.stanford.edu/~jbarral/t1map.html)



**Figure 1: T<sub>1</sub> Mapping Sequences:** a) Inversion Recovery, b) Look-Locker, c) Variable Flip Angle