

# Simulating MRI Signal Contrast Using Spatially Varying $T_1$ and $T_2$ Relaxation Times via the Bloch Equations

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## 1 Introduction

Magnetic Resonance Imaging (MRI) is a powerful, non-invasive imaging modality whose contrast is governed by intrinsic tissue parameters: the longitudinal relaxation time ( $T_1$ ) and the transverse relaxation time ( $T_2$ ) [1]. The response of tissues to radiofrequency excitation and the relaxation dynamics are described by the Bloch equations [2], which capture the time evolution of the net magnetization vector under specific imaging parameters such as repetition time ( $T_R$ ) and echo time ( $T_E$ ).

The steady-state solution of the Bloch equations for a spin-echo sequence provides a formula for the observed signal intensity based on  $T_1$ ,  $T_2$ , and other sequence parameters [3]. This project leverages this steady-state solution to explore how intra-regional variations in  $T_1$  and  $T_2$  (e.g., within a lesion) affect the final signal intensity and perceived image contrast.

While many educational simulations treat tissue regions as homogeneous in relaxation behavior, real biological tissue—particularly pathological tissue—often exhibits heterogeneity in relaxation parameters due to perfusion, necrosis, edema, or other microstructural effects [4]. Simulating these variations can provide insight into how MRI encodes subtle differences in tissue composition.

## 2 Objectives

The primary objective of this study is to simulate and analyze MRI signal contrast arising from spatially varying  $T_1$  and  $T_2$  relaxation times within a tissue slice using the Bloch equations. To achieve this, the study specifically aims to:

1. Implement a computational model for MRI signal intensity using the steady-state solution of the Bloch equations under spin-echo conditions;
2. Encode spatial variation in relaxation parameters ( $T_1$ ,  $T_2$ ) within a lesion region embedded in homogeneous background tissue; and
3. Generate and visualize synthetic contrast maps to assess the effect of intra-lesion gradients on signal intensity.

## 3 Methodology

### 3.1 Overview

This project simulates the MRI signal produced by a tissue slice with spatially varying relaxation times under a spin-echo pulse sequence. A 2D matrix (phantom) will be used to represent a 25-pixel cross-sectional slice, where the background tissue is homogeneous gray matter and a circular lesion is embedded at the center. While the background will have fixed longitudinal and transverse relaxation values ( $T_1$ ,  $T_2$ ), the lesion region will exhibit radial gradients in both parameters, simulating intra-lesion heterogeneity.

The net magnetization vector  $\mathbf{M}(t)$  evolves according to the Bloch equations:

$$\frac{d\mathbf{M}}{dt} = \gamma \mathbf{M} \times \mathbf{B} - \begin{bmatrix} M_x/T_2 \\ M_y/T_2 \\ (M_z - M_0)/T_1 \end{bmatrix} \quad (1)$$

where  $\gamma$  is the gyromagnetic ratio,  $\mathbf{B}$  is the effective magnetic field, and  $M_0$  is the equilibrium magnetization. For this simulation, we consider the steady-state signal under a spin-echo imaging sequence.

### 3.2 Signal Equation (Spin-Echo)

In a conventional spin-echo (SE) imaging sequence with repeated radiofrequency (RF) excitations, the observable signal intensity at each voxel  $(x, y)$  can be modeled. After an initial transient period, the longitudinal magnetization reaches a dynamic equilibrium. If the repetition time (TR) is sufficiently long for the transverse magnetization to decay to zero before the next RF pulse, the system is said to be in a steady-state incoherent (SSI) condition.

Under these circumstances, the steady-state transverse magnetization, which is directly proportional to the MRI signal, is given by:

$$S(x, y) = \rho(x, y) \left(1 - e^{-TR/T_1(x, y)}\right) e^{-TE/T_2(x, y)} \quad (2)$$

where:

- $S(x, y)$  is the MRI signal intensity,
- $\rho(x, y)$  is the proton density (set to 1.0 throughout),
- $TR = 2000$  ms, repetition time,
- $TE = 100$  ms, echo time,
- $T_1(x, y)$  and  $T_2(x, y)$  are the voxelwise relaxation times.

### 3.3 Phantom Design

We construct a  $128 \times 128$  voxel matrix representing a tissue slice. A circular lesion of radius 25 pixels is placed at the center. The following parameter values are assigned:

- *Background (gray matter)*:  $T_1 = 920$  ms;  $T_2 = 100$  ms
- *Lesion Center (core)*:  $T_1 = 1000$  ms;  $T_2 = 90$  ms
- *Lesion Edge*:  $T_1 = 1400$  ms;  $T_2 = 130$  ms

The lesion's  $T_1$  and  $T_2$  values vary radially from center to edge:

$$T_1(x, y) = T_{1, \text{core}} + (T_{1, \text{edge}} - T_{1, \text{core}}) \cdot \frac{r(x, y)}{r_{\text{max}}} \quad (3)$$

$$T_2(x, y) = T_{2, \text{core}} + (T_{2, \text{edge}} - T_{2, \text{core}}) \cdot \frac{r(x, y)}{r_{\text{max}}} \quad (4)$$

where  $r(x, y)$  is the Euclidean distance from the lesion center and  $r_{\text{max}}$  is the lesion radius.

### 3.4 Simulation Platform

All simulations will be implemented in Python using NumPy for numerical array operations and Matplotlib for image visualization. The outputs will include:

- A 2D  $T_1$  relaxation map;
- A 2D  $T_2$  relaxation map; and
- A 2D signal intensity map computed from the spin-echo Bloch equation.

## References

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