COST B21 "Physiological modelling of MR image formation"

Approaches to MRI simulation

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Cyprus, 1 October 2004

Why simulation?

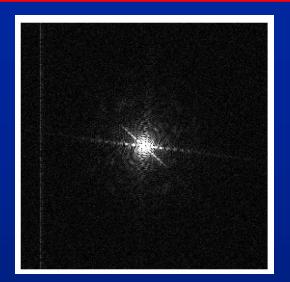
- complement to real, costly MRI measurements
- understanding of the method (varying parameters, sequences,...)
- educational tool in medical and technical environments
- study of artefacts (e.g. their comparison for different sequences)
- development and optimization of sequences
- comparisons of registration and segmentation algorithms

Classification of methods (1)

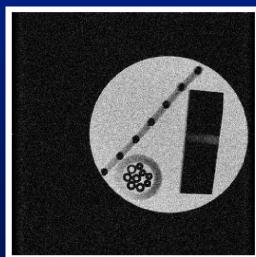
- based on partial or approximate solutions of the Bloch equation with respect to the MR image contrast behaviour
- no account of hardware errors and some physical effects
- differ in the extent of applicability, accuracy, and computation effort
- well-suited mainly for medical and technical education
- Torheim G, Rinck PA. Jones RA, Kvaerness J, 1994, A simulator for teaching MR image contrast behaviour MAGMA 2:515-522
- Rundle D, Kishore S, Seshadri S, Wehrli F, 1990, Magnetic Resonancce Imaging Simulator: A teaching tool for radiology, Journal of Digital Imaging 3(4):226-229

Classification of methods (2)

- based on the k-space formalism
- permits only linear effects to be taken into account
- not able to cover nonlinear gradients or objects distortions due to inhomogeneous main magnetic field
- very fast
- lacks generality







• Petterson J S, Christoffersson J O, Golman K, 1993, MRI simulation using the k-space formalism, Magnetic Resonance Imaging 11:557-568

Classification of methods (3)

- solution of Bloch's equation employing matrix operators
- most general modelling of the imaging process
- decomposing the object into spin attributed with a magnetization vector and local properties like relaxation
- superposition of all spins yields the MR signal
- often results in huge demands on computer memory and speed
- early simulators lacked generality, e.g. because of a limited set of main field inhomogeneity distributions
- Bittoun J, Taquin J, Sauzade M, 1984, A Computer Algorithm for the Simulation of any Nuclear Magnetic Resonance Imaging Method, Magn. Res. Imag. 2:113-120
- Summers R M, Axel L, Israel S, 1986, A Computer Simulation of Nuclear Magnetic Resonance Imaging, Magnetic Resonance in Medicine 3:363-376

Recent projects



MAGNETIC RESONANCE IMAGING

Magnetic Resonance Imaging 22 (2004) 315-328

MRI simulator with object-specific field map calculations¹

Duane A. Yoder^a, Yansong Zhao^b, Cynthia B. Paschal^{b,c,d}, J. Michael Fitzpatrick^{b,d,e,*}

<u>Includes</u>

- object-specific inhomogeneities and static field errors
- chemical-shift
- intravoxel dephasing applied throughout the acquisition protocol

Two parts

- static field perturbation
- signal and image generation

Ignores RF inhomogeneity, eddy currents and gradient nonlinearity.

<u>Inputs</u>

Object definition

Object-specific inhomogeneities

Applied static field errors

Chemical shift values

Description of the acquisition protocols



MR

m

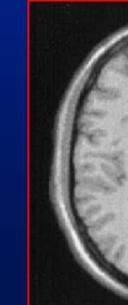


<u>Output</u>

• time sequence of complex numbers organized into kspace;

• image produced by means of a Fourier transform







Theory (1)

The Bloch equation

$$\frac{\mathrm{d} \overrightarrow{M}}{\mathrm{d} t} = \gamma \cdot (\overrightarrow{M} \times \overrightarrow{B}) - \left(\begin{array}{c} M_x / T_2 \\ M_y / T_2 \\ (M_z - M_0) / T_1 \end{array} \right)$$

 $\vec{M} = (M_x, M_y, M_z)^T$ - magnetization vector

 M_0 – magnetization at thermal equilibrium

 T_1, T_2 – spin-lattice and spin-spin ralaxations

- flux of the magnetic field

\vec{B} - flux of the magnetic field

$$\vec{B} = \vec{B}_0 + \Delta \vec{B} + (\vec{G}(t) \cdot \vec{r}) \vec{e}_z + \vec{B}_1(t)$$

 $\vec{B}_0 = B_0 \cdot \vec{e}_z$ - main magnetic field $\Delta \vec{B}$ - inhomogeneity $\vec{G}(t)$ - applied gradients

 $\vec{r} = (x, y, z)^T$ - position in space

 $\vec{B}_1(t)$ - magnetic flux of RF pulses

 \vec{e}_z - unit vector in the z direction

Theory (2)

Solution of the Bloch equation

$$\overrightarrow{M}(\overrightarrow{r},t+\Delta t) = \mathbf{R}_{grad} \cdot \mathbf{R}_{inh} \cdot \mathbf{R}_{relax} \cdot \mathbf{R}_{RF} \cdot \overrightarrow{M}(\overrightarrow{r},t) + M_0(\overrightarrow{r}) \left(1 - e^{-\frac{t}{T_1(\overrightarrow{r})}}\right) \overrightarrow{e}_z.$$

 R_{grad} – influence of the applied gradients

 R_{inh}° – main field inhomogeneity

 R_{relax} – effect of relaxation

 R_{RF} – rotating properties of the RF pulse

Implementation

- The core: Matlab 6.1
- Computation of the field for 181 x 217 x 181 1 mm³ voxels phantom, performed on 1,5 GHz Pentium 4 with 256 MB memory takes 19,5 hours

Validation (experimental data)

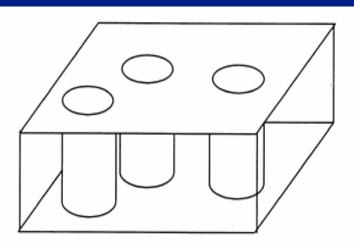


Fig. 4. A diagram of the "sinus" phantom constructed to model air/tissue interfaces. Each cylinder was filled with air with the remainder of the phantom containing distilled water. The width, depth, and height of the phantom are 12 cm, 6.6 cm, and 5.2 cm, respectively.

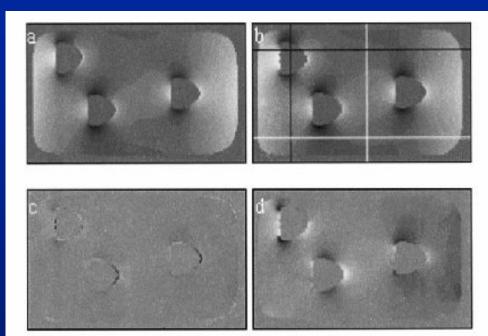


Fig. 7. The perturbed field (a) measured by experiment and (b) calculated by SVC. The range of (a) and (b) is from -2.95 ppm (black) to 5.3 ppm (white). (c) Subtraction image (b) - (a). The range is from -5.3 ppm (black) to 5.3 ppm (white). The average absolute error is 0.16 ppm. Note: There are regions where the experimental and the distorted SVC field map do not intersect (e.g., around the air cylinders) due to the peaks at air/tissue interfaces in the calculated map affecting the amount of distortion when Eq. (28) is applied. Therefore, (d) is included to eliminate these differences. (d) Subtraction image (b) - (a) in intersecting non-air regions. The range is from -1.31 (black) to 1.32 (white) ppm. Note: The lines on (b) indicate the placement of the scan lines in Fig. 8.

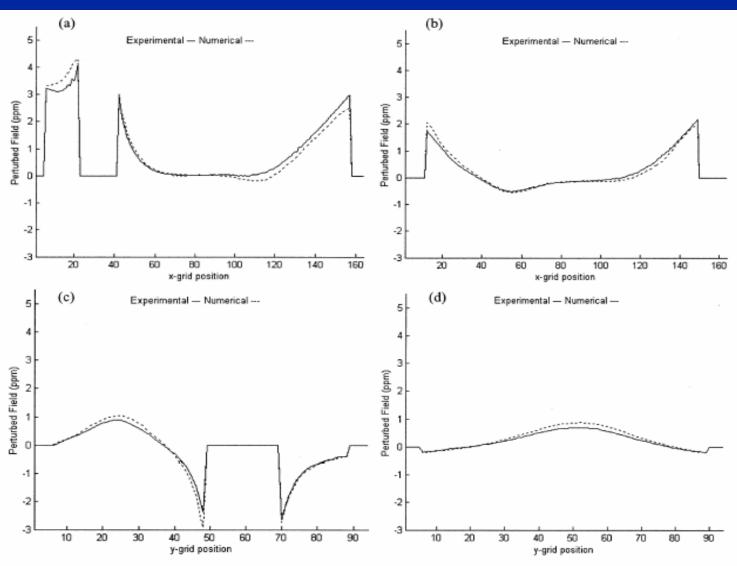


Fig. 8. The perturbation in the field along different scan lines of the sinus phantom. The experimental and SVC solutions are represented by solid and dashed lines, respectively. (a) Perturbed field along the black horizontal line in Fig. 7(b). The max absolute error is 0.7 ppm. (b) Perturbed field along the white horizontal line in Fig. 7(b). The max absolute error is 0.32 ppm. (c) Perturbed field along the black vertical line in Fig. 7(b). The max absolute error is 0.55 ppm. (d) Perturbed error along the white vertical line in Fig. 7(b). The max absolute error is 0.20 ppm.

Validation (known analytic solution)

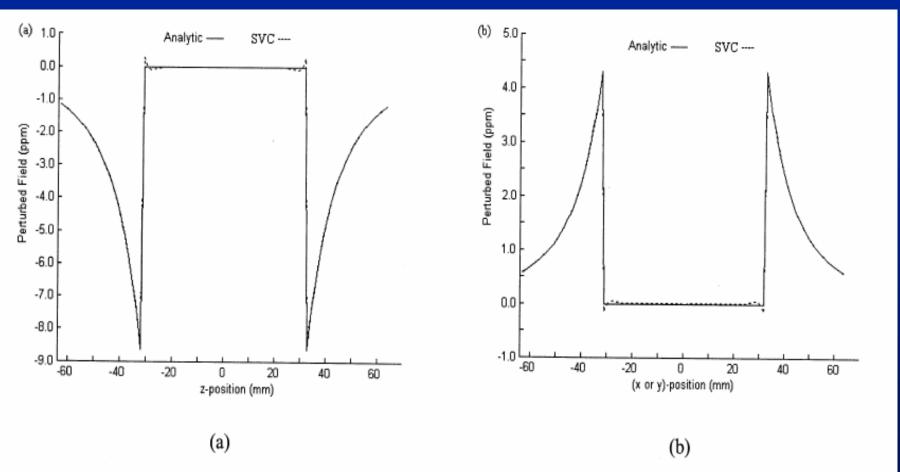


Fig. 1. The perturbation of the field along two orthogonal scan lines through the center of a sphere of water, $\chi = -9.05 \times 10^{-6}$, of radius 32 mm in a 1.5 T static field. Position 0 represents the center of the sphere. The analytic and SVC solutions are represented by solid and dashed lines, respectively. (a) The perturbed field along the z axis. The maximum error is 0.15 ppm.

Part 2 - signal and image generation

<u>Object</u> <u>definition</u>

• every voxel's T₁, T₂, PD, frequency offset values

Simulator

- executing the pulse sequence
- solving the Bloch equations to update each magnetization vector
- storing data (signal readout)

Event sequencer

- RF pulse (start time, flip angle, duration of the pulse, axis about which to rotate)
- gradients (start time, duration, amplitude)
- readout (start time, number of samples, sampling frequency, repetition time)

Validation – part 2

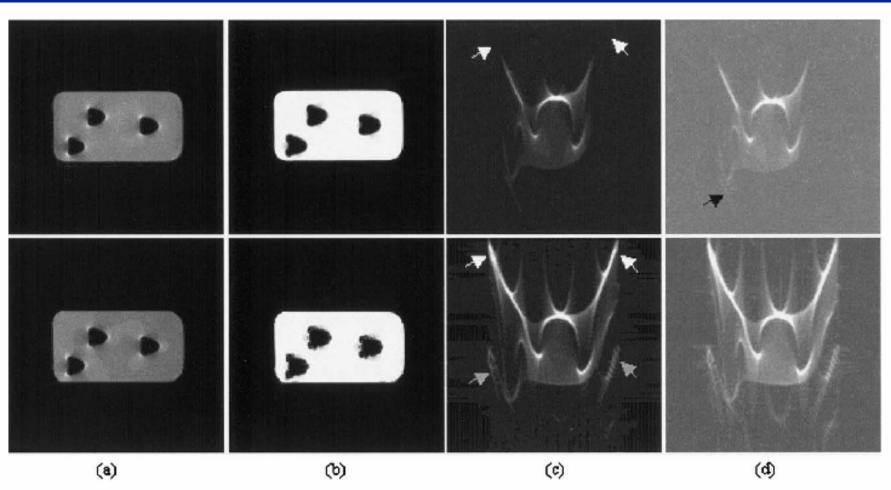
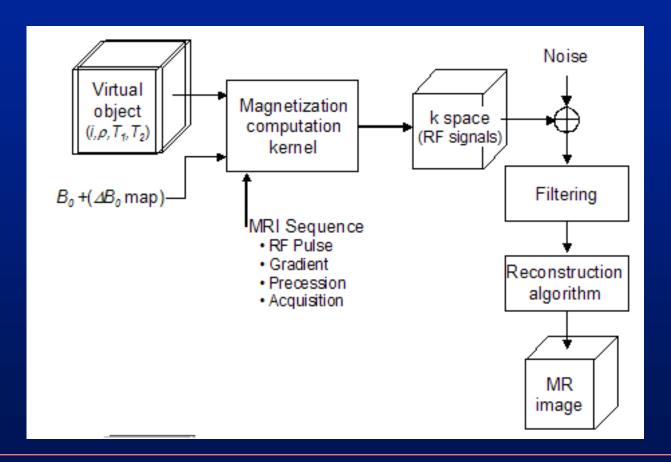


Fig. 10. Comparison of experimental (top row) and simulated (bottom row) images of sinus phantom. The frequency encoding direction is from left to right and the FOV = 20 cm for all images. The arrows highlight various features that are discussed in Section 3.2. (a) SE. Imaging parameters: TE = 20 ms, TR = 2000 ms, $256 \times 256 \text{ imaging matrix}$. (b) GE. Imaging parameters: TE = 10 ms, TR = 800 ms, $256 \times 256 \text{ imaging matrix}$. (c) SE EPI. Imaging parameters: TE = 180 ms, TR = 800 ms

Implementation - part 2

- PC platform, MS Visual C ++ 6.0
- GUI dependent on Visual C++ objects, but the core can be compiled on other architectures
- computationally expensive

SIMRI - interactive 3D MRI simulator

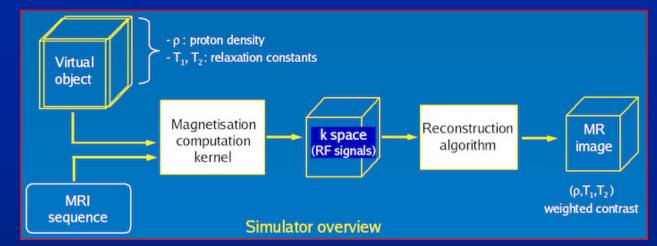


Benoit-Cattin H., Belaroussi B., Bellet F., Odet C, 2004, A versatile and interactive 3D MRI simulator, Journal of Magnetic Resonance

SIMRI

- based on Bloch equations
- T2* effect
- chemical shift artifact incl. off-resonance
- static field inhomogeneity
- susceptibility variation within an object

- spin echo
- gradient echo
- 1D, 2D, 3D images

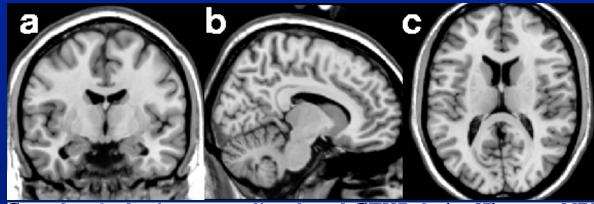


- C programming language
- MRI programming: "high level" C functions with simple programming interface (Python)
- magnetization kernel: parrallel computing (PC grid architecture)
- 1D interactive interface to illustrate magnetization vector motion and MRI contrast

Benoit-Cattin H., Belaroussi B., Bellet F., Odet C, 2004, A versatile and interactive 3D MRI simulator, Journal of Magnetic Resonance

"Montreal" phantom

3-D high-resolution, anatomically accurate human brain phantom



Coronal, sagittal and transverse slices through CJH27, the (n=27) average MRI volume used to build the phantom.

Construction

- preprocessing intensity nonuniformity reduction
- automatic classification of preprocessed MRI volume
- manual correction of the classified data (%)

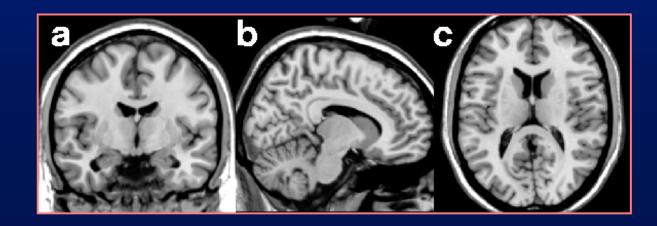
Collins D.L., et al., 1998, Design and Construction of a Realistic Digital Brain Phantom, IEEE Transactions on Medical Imaging, 17:3

C.A. Cocosco C. A., et al., 1997, BrainWeb: Online Interface to a 3D MRI Simulated Brain Database, NeuroImage, vol.5, no.4, part 2/4, S425.

"Montreal" phantom

Phantom characteristics

- high resolution (1 mm isotropic voxels) low-noise data set
- 27 scans (T1-w GE TR/TE/FA 18ms/10ms/30°) of the same individual in stereotaxic space
- subsampled and intensity averaged
- the volume contains 181 x 217 x 181 voxels, covers the whole brain
- exhibits fine anatomical details normally obscured by noise in single image



Tissue volume definition

ID	Description
GM	grey matter within the brain parenchyma
WM	white matter within the brain parenchyma
CSF	cerebro spinal fluid surrounding the brain and within the ventricles
GL	layer of glial tissue lining the ventricles
M+S	muscle and skin
OTH	other tissue
FAT	fatty tissue
SKN	mostly skin
SKL	skull (does not include sinuses)
AIR	air outside head and within sinuses



BrainWeb:

Online Interface to a 3D MRI Simulated Brain Database



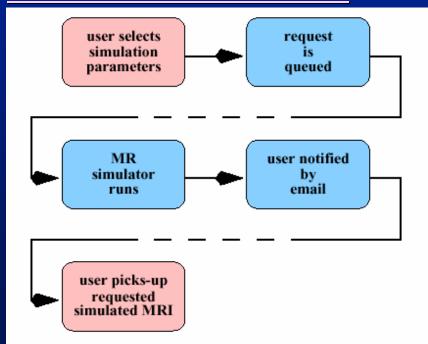
McConnell Brain Imaging Centre, Montréal Neurological Institute, McGill University, Montréal, Canada



Application examples

- study the performance of anatomical brain mapping techniques (relation to different MR acquisition parameters)
- validation of quantitative analyses for neuropathology (e.g. MS lesion quantification, medical pattern recognition and image processing techniques)

Custom MRI simulation



http://www.bic.mni.mcgill.ca/brainweb/

On-line interface to Simulated Brain Database

Click on an image to navigate, or use the buttons below



Modality: (you can choose one of the following pulse sequences)

T1.◆

T2 🗣

PD⊕

Slice thickness: (in-plane pixel size is always 1x1mm)

1mm◆	3mm⊕	5mm⇔	7mm◆	9mm⊕

Noise:

0%♦ 1%♦ 3%♦ 5%♦ 7%♦ 9%♦

Intensity non-uniformity ("RF"):

0%

20%

40%

40%

[Reset form] [View] [Download]

	selectable values		
	normal;		
	MS lesion mild, moderate, severe		
mess	1 - 10 mm		
nique	SE, IR, SFLASH, CEFAST, FISP,		
	FLASH, DSE_EARLY, DSE_LATE		
time	(any) ms		
time	(any) ms		
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ce thickness)	1 - 10 mm		
	0 – 10 %		
	0-60 %		

Questions

- What kind of simulator is needed for COST B21?
- Can MRI <u>texture</u> be simulated?
- Can we share simulators?