Lecture #6 In Vivo Water

- Topics
	- Hydration layers
	- Tissue relaxation times
	- Magic angle effects
	- Magnetization Transfer Contrast (MTC)
	- CEST
- Handouts and Reading assignments
	- Mathur-De Vre, R., "The NMR studies of water in biological systems", Prog. Biophys. Molec. Biol. Vo1 35, pp 101-134, 1979.
	- Bydder, M., et al., "The magic angle effect: a source of artifacts, determinant of image contrast, and technique for imaging", JMRI,25:290-300, 2007.
	- Henkelman, RM, et al., "Magnetization transfer in MRI: a review", NMR in Biomedicine, 14:57-64, 2001.
	- van Zijl, P., et al., "Chemical exchange saturation transfer (CEST): What is in a name and what isn't", MRM 65:927-948, 2011.

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Relaxation Recap

- NMR relaxation is due to interactions between nuclear spins and local fluctuating fields arising from…
	- Thermal motion of the lattice
	- Molecular motion
	- Chemical exchange processes
	- Paramagnetic centers
- Effects of these interactions depend on the time scale and nature of the motion.
	- T₁ most sensitive to fluctuations at the Larmor frequency $\omega = \gamma B_0$.
	- T_2 sensitive to fluctuations at the very low frequency $(\omega = 0)$.
	- T₁₀ most sensitive to fluctuations at the Rf frequency $\omega = \gamma B_{1}$.

2 Note, we haven't yet discussed $T_{1\rho}$ or paramagnetic effects.

In Vivo Water

• Relaxation times of in vivo water protons are typically much shorter and diffusion constants much lower than those of pure water.

- A significant fraction of in vivo water is associated with macromolecules in the form of an hydration layer.
- Hydrogen bonding to hydrophylic surfaces results in restricted motion, cross-relaxation, and chemical exchange effects.

Edelman, et al., *Clinical MRI*, W.B. Saunders Co, Phil., 1996.

Tissue Water Models

• The two water pools are in fast exchange leading to relaxation rates being avearges of the rates for the two pools.

- There are also three compartment models which add a tightly bound water "ice-like" pool. This lead to:
	- $-$ Tissue T₁ dominated by total water content and fraction in the hydration layer
	- Tissue T_2 dominated by thickness of hydration layer as well as the size of tightly bound pool (see Fullerton, et al, Mag Res Imag, 1:209-226, 1982).

de Graaf, *In Vivo NMR Spectroscopy*, Wiley, 2002.

Biological Water T_1 s and T_2 s

1.5T

Biological Water T_1 s and T_2 s

"proton density"

" T_1 -weighted"

 T_2 -weighted"

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What do these images tell us about the tissue in this tumor versus normal brain?

Biological Water T_1 **S and** T_2 **S**
Table 2.6. Longitudinal and transverse relaxation times of water in

biological tissues at different magnetic field strengths¹

¹Reported values are average relaxation times from reported literature [241-256].

¹H Brain Metabolite T_1 s and T_2 s

de Graaf, *In Vivo NMR Spectroscopy*, Wiley, 2002.

¹Reported values are average relaxation times from reported literature $[257-272]$.

²Measured on humans.

 3 Measured on rats.

Magic Angle Effects

- Observation: on moderate to short TE sequences, signal intensity of tendons, ligaments, and cartilage depends on tissue orientation with respect to the large B_0 field.
- These highly ordered tissues contain collagen fibers with bound water that is not free to tumble isotropically.
- Dipole interaction among bound water protons is angle dependent.

Collagen fiber during hydration

The Nuclear Dipolar Coupling Hamiltonian

• Hamiltonian

$$
\hat{H}_{dipole} = -\frac{\mu_0 \gamma_I \gamma_S}{2\pi r^3} \hbar \left(\hat{\vec{I}} \cdot \hat{\vec{S}} - \frac{3}{r^2} (\hat{\vec{I}} \cdot \vec{r}) (\hat{\vec{S}} \cdot \vec{r}) \right)
$$
 where \vec{r} vector from spin *I* to spin *S*

• Secular approximation:

$$
\hat{H}_{dipole} = d\left(3\hat{I}_z\hat{S}_z - \hat{I}\cdot\hat{S}\right) \text{ where } d = -\frac{\mu_0\gamma_1\gamma_S}{4\pi r^3}\hbar\left(3\cos^2\Theta_{IS} - 1\right)
$$
\ndipole coupling
\n
$$
\text{angle between B}0\nconstant\nspins I and S
$$

- With isotropic tumbling, the time average of $\hat{H}_{\text{dipole}} = 0$
- $-$ With non-isotropic tumbling, $\hat{H}_{\text{dipole}}(t) \neq 0$

Magic Angle Spinning

- A common technique used in solid-state NMR is to artificially spin the sample in order to average-out dipolar coupling effects.
- Residual dipolar coupling effects disappear is the sample is spun at an angle of $3\cos^2\theta_0 - 1 = 0$ ($\theta_0 = 54.7^\circ$) relative to B₀

• Magic angle spinning is also use to analyze tissue biopsy samples.

Collagen-bound water

• No spinning allowed for in vivo studies, but we do have restricted tumbling. 2.5 والمستوات والمتواطن والمسارين والمتواطن والمتواطن والمتواطن

Figure 3. (a) Parameters associated with the magnetic field generated by a classic magnetic dipole μ at the origin. The gray lines represent the local direction of the dipolar field generated by μ . (b) Dipolar interaction between two protons (μ_1 and μ_2) in a water molecule that is bound to a collagen fiber (not to scale). Each proton dipole generates a local dipolar field as shown in (a). Each proton experiences a small contribution of magnetic field from its (many) neighbor protons.

Figure 4. The $(3\cos^2\theta - 1)$ factor in the equation for nuclear dipolar interaction. (The arrows identify the discrete sampling points used in the microscopic MRI experiments described in text.⁴⁰) (Reprinted from Xia Y. Relaxation anisotropy in cartilage by NMR microscopy (μ MRI) at 14 μ m resolution. Magn Reson Med, Copyright © 1998, John Wiley & Sons, Inc. Reprinted by permission of Wiley-Liss, a subsidiary of John Wiley & Sons, Inc.)

> Xia, et al. Investigative Radiology, Volume 35, Number 10, 602–621 (2000)

Example: Tendon Imaging • $T₂$ of tendons is strongly dependent on the angular

orientation with respect to B_0 : magic angle = 54.7^o

Bydder , et al. JMRI, 25:290–300 (2007)

Cross Relaxation in Vivo

Consider the following general pulse sequence:

• If no interactions between saturated and observed components, we get familiar results.

– e.g. fat suppression, water suppression

• What happens if the saturated and observed components interact?

Magnetization Transfer in Tissue

- Selectively saturate short- T_2 pool (bound protons)
- Magnetization exchanged between saturated bound protons and unsaturated mobile protons

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What about T_1 ?

Why do macromolecules have a short T_2 ? • Observe reduced magnetization of longer T_2 (mobile) water protons

Dipolar coupling leads to NOE effect

S spin on small mobile molecule S spin on large immobile molecule η NOE =1+ ^γ *^S* γ *I W*² −*W*⁰ *W*⁰ + 2*WI* + *W*² \$ % & ' () Here we're dealing with slowly tumbling macromolecules. So NOE is negative.

Magnetization Transfer Contrast (MTC)

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MTC Imaging

- White matter: lots of macromolecules (primarily myelin)
- Gray matter: less macromolecules
- Blood: very few macromolecules

Conventional Imaging

MR Angiography

no MTC with MTC

MTC MRI

• Interestingly, the equations are very similar if MTC is based on a dipole-dipole interaction (cross relaxation) or a chemical exchange effect.

Chemical Exchange Staturation Transfer (CEST)

- Unique image contrast can be generated for spin systems in slow or slow-intermediate chemical exchange.
- The basis idea is to selectively saturate spins in one chemical environment, which are then exchanged into a second environment that can be readily imaged.

- Image the water with and without Rf saturation
- Need exchange slow enough to have two distinct peaks, but fast enough to allow magnetization transfer before T_1 recovery.

Chemical Exchange Staturation Transfer (CEST)

• Longitudinal magnetization with chemical exchange...

$$
A_{\overline{k}_{BA}}^{\underline{k}_{AB}}B \implies \frac{dM_z^A}{dt} = \frac{M_z^{A,0} - M_z^A(t)}{T_{1A}} - k_{AB} M_z^A(t) + k_{BA} M_z^B(t)
$$

$$
\frac{dM_z^B}{dt} = \frac{M_z^{B,0} - M_z^B(t)}{T_{1B}} + k_{AB} M_z^B(t) - k_{BA} M_z^A(t)
$$

- Assume slow, slow-intermediate exchange.
- Then, if we can selectively saturate component B with sufficient RF irradiation such that

$$
M_z^B = M_z^{B,0} \frac{1 + (\omega_0 - \omega)^2 (T_2^B)^2}{1 + \omega_1^2 T_2^B T_1^B + (\omega_0 - \omega)^2 (T_2^B)^2} \approx 0,
$$

 M_z^\prime $_{\textrm{\tiny{Z}}}^{A}\left(\infty\right)$ \bm{M}_z' $\frac{(\infty)}{A,0} = \frac{1}{1+k}$ $1 + k_{AB}T_1$ then the new equilibrium for the A component is $\frac{M_z(\omega)}{M^{A,0}} = \frac{1}{1 + kT^A}$

CEST

What does the Z-spectrum from the previously described MT effect look like?

Van Zijl, et al., MRM 65:927–948 (2011)

saturation efficiency, and the amplitude and duration of saturation pulse.

Exploitable Exchange Pathways

ATOM (PROTON) EXCHANGE

small molecules diaCEST some paraCEST macromolecular supraCEST glycoCEST gagCEST multiple molecules **APT**

For this lecture, we'll just focus on **a** and leave **b** and **c** for when we discuss contrast agents.

MOLECULAR EXCHANGE paraCEST

Ln(III)-OH₂ complexes $Ln(III)-XH_n$ complexes where X represents any coordinated molecule

Concentrations needed for ~5% CEST effect with clinical feasible B_1 strengths

COMPARTMENTAL EXCHANGE lipoCEST

CEST examples

- Amide Proton (-NH) Transfer (APT)
	- Chemical shift \sim 3.5 ppm below water
	- Very slow exchange rate $(\sim 30 \text{ s}^{-1})$ and relatively high concentrations
	- Easy to saturate and hence suitable for 3 T and higher
	- Strong pH dependence on exchange rate
	- Applications: imaging of changes in protein content and pH (e.g tumors)
- Hydroxyl (-OH) CEST
	- Chemical shifts ~1 ppm below water: glucose, glycogen, mI, GAG
	- Moderate exchange rate $(\sim 500 1500 \text{ s}^{-1})$ \rightarrow relatively high-power saturation needed
	- Small $\Delta\omega$ with respect to water \rightarrow need for high fields (\geq 7), preclinical models
	- Applications: glucose metabolism (glucoCEST, glycoCEST), cartilage (gagCEST)
- Amine (-NH2) CEST (free amino acids, proteins, peptides)
	- Chemical shifts ~3 ppm below water: e.g. glutamate (gluCEST), creatine
	- Faster exchange rate (\sim 2000-6000 s⁻¹) \rightarrow high transfer efficiency, but high power
	- need for high fields $(\geq 7 \text{ T})$, preclinical models
	- Applications: imaging of protease activity in tumors, pH, glutamate

Some CEST Images

Glioblastoma

Sakata, et al., Journal of Neurooncol, 2015

Multiple Sclerosis

Fig 5. Results of CEST MRI at 7 T on healthy control and MS patient. (A) Z-spectra arising from healthy white matter, MS patient white matter, and MS lesion, solid lines, left y-axis. CEST asymmetry is also shown, dashed lines, right y-axis. (B) Anatomical image of healthy subject with the calculated APT asymmetry map shown in panel (C). (D) Anatomical image of MS patient with calculated APT asymmetry map found in panel (E).

Fig 6. Results from GlycoCEST of skeletal muscle at 7 T. (A) T₁-weighted anatomical image, (B) reference image for glycogen resonance (1.0 ppm), (C) normalized image for glycogen resonance (-1.0 ppm), (D) shift map calculated from polynomial fit with color scale in Hz, and (E) asymmetry map for glycogen (1.0 ppm).

Dula, et al., Journal of Neuroimaging Vol 23 No 4 October 2013

Muscle glycogen

Next Lecture: Redfield theory I