## Tau<sub>b</sub>: A Metabolic Neuroimaging Biomarker

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Introduction: Capillary lumen water is "well-mixed" (1). During passage through a typical  $^{1}\text{H}_{2}\text{O}$  MRI voxel [(1 mm) $^{3}$ ],  $\text{H}_{2}\text{O}$  molecules exchange into and out of blood cells, and collide with the capillary wall, many times. The escape probability [equilibrium extravasation] is the inverse of the mean  $\text{H}_{2}\text{O}$  molecule lifetime in a blood capillary ( $\tau_{b}$ ).  $\tau_{b} = 0.5 \cdot r \cdot (P_{W})^{-1}$ ; with r and  $P_{W}$  the voxel *mean* capillary radius and water permeability coefficient, respectively (2). With sufficient compartmentalized paramagnetic contrast agent concentration, [CA], the  $^{1}\text{H}_{2}\text{O}$  inversion recovery (IR) is non-mono-exponential and compartmental  $\text{H}_{2}\text{O}$  lifetime ( $\tau$ ) values can be determined (2,3). But it is likely impossible to reach the required blood [CA] *in vivo* (4). However, even with mono-exponential IR, if [CA] is *varied*,  $\tau$  values can be measured (2,4). Originally, blood [CA] was deliberately varied (animal model) to obtain  $\tau_{b}$  (5). Then it was shown that, in the human brain, a single CA bolus, (Dynamic-Contrast-Enhanced) DCE-MRI, is sufficient because [CA] increases and decreases during the bolus passage: CA does not extravasate in the normal brain (6). But the approved monomeric Gd(III) chelate CAs do extravasate in brain tumors. Here, we use a superparamagnetic iron oxide (SPIO) nanoparticle CA that can be bolus injected (7). During the first pass, and indeed the first hour, it does not extravasate even from tumor capillaries. We use the 2-site-exchange (2SX) shutter-speed paradigm (SSP) DCE-MRI analysis (6).

Methods: DCE-MRI quantitative  $T_1$  (q $T_1$ ) data were acquired on a Siemens 3T Tim Trio instrument with a multislice 2D GRE-EPI based sequence with a (128)<sup>2</sup> image matrix, resulting in nominal (2 mm)<sup>3</sup> isotropic resolution (8). Points were obtained in the CA steady-state before, and after IV injections of Ferumoxytol [Fe-tol; MW = 750 kDa] (7), fractionated into 3 doses [1:2:4] totaling 4 mg(Fe)/kg. The q $T_1$  acquisitions were initiated 120 s after each Fe-tol injection, during the plasma CA steady-state period; *i.e.*, when plasma [CA] was uniform and constant: [CA] was larger after each successive injection. This was a multi-day study, and a similar protocol using the momomeric GdHPDO3A CA, was conducted 24 hr prior to the Fe-tol MRI session. Six *glioblastoma multiforme* [GBM] subjects consented. Data are also reported from a separate 7T study of 2M/4F healthy controls using GdHPDO3A.

Results: Fig. 1 displays sample data from a 52 y M GBM subject. In the center is an axial qT<sub>1</sub>-w image obtained 30 min after GdHPDO3A injection. The CA-enhancing tumor is clearly visible at the lower left. Inset are data (crosses) obtained from 4 representative ROIs [frontal WM, thalamus, putamen, and tumor; ellipses] during Fe-tol injections of this subject. Each plot shows the R<sub>1b</sub>-dependence of R<sub>1t</sub>. [R<sub>1</sub> ≡ T<sub>1</sub><sup>-1</sup>: R<sub>1b</sub> is taken at 4 times from a sagittal sinus ROI, each R<sub>1t</sub> at the same 4 respective times from the ROI indicated]. If the tracer pharmacokinetic paradigm [τ<sub>b</sub> → 0] obtained, the R<sub>1t</sub>,R<sub>1b</sub> plot would be linear. None are, and all are well fitted by 2SX SSP expression (4,6) curves with τ<sub>b</sub>, v<sub>b</sub>, and R<sub>1exv</sub> varied [v<sub>b</sub> is the blood volume fraction, R<sub>1exv</sub> the pre-exchange extravascular <sup>1</sup>H<sub>2</sub>O R<sub>1</sub>]. The population- and ROI-averaged v<sub>b</sub> and τ<sub>b</sub> values are given in the **Table** [NA, normal-appearing]. For comparison, the normal WM and GM healthy control biomarkers are

| Table 1 Human Brain                   | ROI SSP Values |                    |
|---------------------------------------|----------------|--------------------|
|                                       | V <sub>b</sub> | τ <sub>b</sub> (s) |
| Healthy Controls (n = 6) <sup>a</sup> |                |                    |
| NWM                                   | 0.014          | 0.345              |
| NGM                                   | 0.031          | 0.403              |
| Glioblastoma (n = 6) <sup>b</sup>     | •              | •                  |
| NA-frontal WM                         | 0.008          | 0.44               |
| NA-thalamus                           | 0.017          | 0.38               |
| NA-putamen                            | 0.012          | 0.43               |
| tumor                                 | 0.046          | 5.55               |

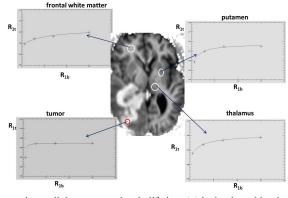
also listed. In the GBM brain, the  $v_b$  values are generally smaller than normal, except in the tumor. The  $\tau_b$  values in the NA-GBM brain are rather normal, and most definitely non-zero. However, in tumor tissue  $\tau_b$  is increased by an order-of-magnitude. Is this remarkable increase because r is increased,  $P_{W}'$  is decreased, or both? In separate analyses, we have found - for cells in suspension, and in breast tumors and in myocardium  $in\ vivo$  - the

M-putamen | 0.012 | 0.43 | 0.44 | 0.55 | 0.046 | 5.55 | 0.046 | 5.55 | 0.046 | 0.55 | 0.046 | 0.55 | 0.046 | 0.55 | 0.046 | 0.55 | 0.046 | 0.55 | 0.046 | 0.55 | 0.046 | 0.55 | 0.046 | 0.55 | 0.046 | 0.55 | 0.046 | 0.55 | 0.046 | 0.55 | 0.046 | 0.55 | 0.046 | 0.046 | 0.55 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.046 | 0.0

tonimant  $P_W$  (active) component in the MS brain, where  $\tau_b$  is also increased: 1.) PMRSI shows PCI and 2.) <sup>23</sup>NaMRSI shows tissue Na content [likely Na<sub>i</sub>] is increased [~28%] in MS brain (9), and 3.)  $\tau_b$  is significantly increased [almost doubled] in MS lesions (unpub.). It is well-known that MS lesion capillaries are CA-permeable, and those in GBM tumor tissue even more so (7). There is little doubt this is due to widened para(endothelial)cellular pores. Thus, paracellular water extravasation [a  $P_W$ (passive) pathway] must also increase. But this would make  $\tau_b$  decrease. The conclusion is inescapable that the large  $\tau_b$  increase observed must be dominated by a large  $P_W$ (active) decrease in the GBM tumor.

**Discussion:** The mean intracellular water molecule lifetime  $[\tau_i]$  is dominated by active trans-membrane water cycling [10<sup>12</sup> H<sub>2</sub>O molecules/s/cell] (3). This process accompanies the active trans-membrane osmolyte cycling paced by the driving cell membrane ATPase transporter. For mammalian cells, this is the Na<sup>+</sup>/K<sup>+</sup>ATPase [NKA] pump, possibly the most important enzyme in biology. The  $\tau_i$  magnitude is sensitive to the ATP<sub>i</sub> and  $K_0$ substrates of, and to specific inhibitors of, the driving ATPase transporters (3,10,11). Fig. 2 is a cartoon of the "gliovascular unit" (12). Endothelial [gray], neuroglial [pink], and neuronal [blue] cells, in proximity with synaptic dimensions, act in exquisite metabolic synchrony (12,13). Active trans-membrane water cycling processes are indicated with stars. A cascade of these could constitute the Pw'(active) pathway [2d]. Possible Pw'(passive) pathways, simple transcellular H<sub>2</sub>O diffusion across cell membranes [2b], transcellular transport through membrane aquaporin H<sub>2</sub>O channels and/or transcellular leakage through membrane transporters [2c] are also shown. The paracellular H<sub>2</sub>O pathway is shown as [2a], but this caries a miniscule flux compared with transcellular pathways. Our results strongly suggest that the  $P_W$ (active) pathway [2d] dominates the  $\tau_b$ magnitude. If this changes in parallel with changes in "Magistretti Metabolic Mechanism" [2e,f,g] fluxes (12,13),  $\tau_b$  provides a high-resolution MRI biomarker sensitive to changes in cerebral neuronal activity. The lower the NKA activity, the larger the  $\tau_b$  value. If this is true, cells in the GBM tumor have extremely slow NKA activity. Grant Support: NIH: RO1-NS040801; UO1-CA154602; R44 CA180425.

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T<sub>b</sub>

H<sub>O</sub>

**Figure 2.** Gliovascular unit water exchange mechanisms determining mean water molecule lifetimes in blood ( $\tau_b$ , beige), interstitium ( $\tau_o$ , aqua), and endothelial ( $\tau_i$ ', gray), neuroglial ( $\tau_i$ , pink), and neuronal ( $\tau_i$ '', blue) cell spaces. The equilibrium paracellular (a), simple diffusion (b), facilitated transcellular (c), and active water cycling (d, stars) pathways are indicated, as are "Magistretti steps" (e.f.g).