## Monitoring of T2 with application of diffusion gradients to remove microcirculation contributions to signal for optimisation of diffusion protocols and generation of flow-free T2 maps

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Introduction: The bi-exponential IVIM model used in diffusion-weighted magnetic resonance imaging (DW-MRI) separates self diffusion from a second, faster, pseudo-diffusion component that is often equated to perfusion or flow (1); with applied diffusion gradient b-values of 200 s/mm<sup>2</sup> or higher, this flow component is considered essentially absent from the image. Maki (2) describes the varying types of microcirculation that may be present within an imaging voxel, and that diffusion

gradients induce either phase shift or phase dispersion in various forms of microcirculation leading to signal attenuation. Commonly in DW-MRI, diffusion gradient parameters  $\delta$  and  $\Delta$  are calculated by the scanner and vary with TE, but explicitly controlling these parameters allows further investigation of complex diffusion phenomena (3). Using explicit  $\delta$  and  $\Delta$  control to acquire DW-MRI at varying TE, we investigate the effect of applied diffusion gradients to T<sub>2</sub> measurements. Where T<sub>2</sub> is often considered in MRI as a simple exponential decay, there is evidence it reflects a more complex picture regarding tissue structure (4). For highly vascular tissue (vascular fraction f), where blood has a known  $T_{2blood}$ , flow desensitization with applied diffusion gradient gives the potential to generate  $T_2$  maps that eliminate the flow component and more accurately reflect the intrinsic tissue T<sub>2</sub>:

## $S=S_0(f.exp(-bD^*).exp(-TE/T_{2blood}) + (1-f).exp(-bD).exp(-TE/T_{2tissue}))$

where D and D\* are the apparent diffusion coefficients of the tissue (random) and microcirculatory (pseudo-random) components, and a sufficiently large b-value is used to reduce the first term to zero; the decay curve then reduces to a single exponential, where the (1-f).exp(-bD) scaling factor is constant and can be absorbed into S<sub>0</sub>. Incomplete removal of the flow term would necessitate fitting the full five parameters (f, D, D\*, T<sub>2blood</sub>, T<sub>2tissue</sub>) and may best be accomplished in tandem with IVIM diffusion modeling; conversely, monitoring changes in T<sub>2</sub> with increasing applied b-value has the potential to unambiguously identify the minimum threshold beyond which microcirculatory components can be considered absent, which may aid in modeling diffusion data.

Method: Five volunteers were recruited and consented. Coronal free-breathing DW-MRI images of the abdomen were acquired on a MAGNETOM Avanto 1.5T scanner (Siemens AG, Healthcare Sector, Erlangen, Germany). Three echo times were acquired (71.2 - 130 ms), with either four (0, 50, 100, 200 s/mm<sup>2</sup>, n=2) or two (0, 200 s/mm<sup>2</sup>, n=3) b-values, with multiple averages (6 to 8), TR 4500 ms,  $\delta$  17.6 ms and  $\Delta$  30 ms. A diffusion phantom with multiple physiological T<sub>2</sub> compartments (no flow component) was also scanned with matched b-values and TEs. ROIs were drawn for different regions in the phantom, for each kidney, and homogenous tissue regions in the liver and spleen; for each b-value series of TE the data were fitted with a mono-exponential decay model using in-house software (ADEPT, ICR, UK). Results are given as mean  $\pm$  s.d., with significance from comparison using 2-tailed paired t-test set at 0.05.

Results: Single-compartment phantom data shows that the presence of the diffusion gradient has no effect on  $T_2$ estimates over a relevant range of b-value and TE (figure 1) as expected, whereas the mono-exponential model applied to liver and spleen (figure 2a for ladder plots of liver) shows a systematic variation in observed  $T_2$  in the presence of the  $b=200 \text{ s/mm}^2$  gradient pulses, decreasing from  $46.0 \pm 4.9$  to  $38.3 \pm 6.2$  ms in the liver (p=0.024), and from  $85.8 \pm 13.8$ 

to 78.9  $\pm$  11.3 ms in the spleen (p=0.037). Figure 3 shows an example map of  $\Delta T_2$ , showing structure visible within the liver, kidney, and spleen. In the kidney ROIs, exclusion of an apparent outlier gives a significant decrease in  $T_2$  from 103.0 ± 12.5 to 94.8 ± 10.2 ms (p<0.001). In the subset where acquisitions at b=50 and 100 s/mm<sup>2</sup> were acquired, the decrease observed occurs predominantly for b=0 to 50 s/mm<sup>2</sup> (figure 2b for spleen and liver, n=2), leveling off from b=100 to 200 s/mm<sup>2</sup>.

Discussion: The observed T<sub>2</sub> changes for mono-exponential T<sub>2</sub> fit with applied diffusion gradients are consistent with the removal of fast microcirculatory components of the voxel and their contribution to the overall T<sub>2</sub> decay curve. The necessarily increased TE to allow for the gradient pulses does not preclude sufficient sampling of the (now simplified) signal decay curve, whereas attempting to resolve components of the T<sub>2</sub> decay curve from a standard multi-echo acquisition remains problematic when the composition of tissue voxels is unknown and/or complex. We demonstrate the potential use of diffusion gradients to produce microcirculatoryfree  $T_2$  maps, as well as suggesting that an applied b-value of 200 s/mm<sup>2</sup> appears large enough to

remove the pseudo-diffusion component modeled in IVIM DWI. Limitations of the current protocol are the sparse sampling of TE and the long echo times used for  $T_2$  estimation in abdominal organs; further exploration of the TE/b-value space will allow a more confident assignment of an optimal protocol. Vascular properties that vary on a short timescale, particularly in highly vascular organs or heterogeneous tumours, may confound the use of  $T_2$  as a biomarker; the ability to modulate or remove the vascular component of the observed  $T_2$  may provide a more robust measure of tissue water environment.

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Acknowledgements: This work was funded by CR-UK grant number C7273. We also acknowledge the support received for the CRUK and EPSRC Cancer Imaging Centre in association with the MRC and Department of Health (England) (grants C1060/A10334 and C16412/A6269) and NHS funding to the NIHR Biomedical Research Centre.

Figure 1:  $T_{\rm 2}\ of$  simple diffusion phantom (inset) is





Figure 3: Subtraction map for T<sub>2</sub>(b0) - T<sub>2</sub>(b200), showing structure within abdominal organs



Proc. Intl. Soc. Mag. Reson. Med. 21 (2013)

