

# Characterization of myelin damage in multiple sclerosis using myelin water imaging: Insight from simulations on a four pool model of white matter

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## Introduction:

Magnetic resonance imaging is a very sensitive technique for the study of white matter (WM) damage in multiple sclerosis (MS), but suffers from a general lack of specificity to the underlying pathology. Quantitative imaging techniques that provide putative *in vivo* measures related to myelin, such as myelin water imaging [1], have been developed and shown to correlate strongly with demyelination [2]. To help further the understanding of WM, a four-pool model has been proposed and characterized [3], and its properties have recently been measured in bovine WM [4]. While perhaps overly complicated for routine *in vivo* imaging applications, this model can provide insight into the impact of changes in the four-pool complex on observations made using established methods for quantitative imaging. We have studied the effect of modeling tissue pathology on the resulting  $T_2$  spectrum by simulating CPMG measurements using a four-pool model, analysed using a multi-component  $T_2$  model.

## Methods:

A modified four-pool model of bovine white matter, illustrated in Figure 1, was constructed from an existing model [4], with changes to the treatment of the solid pool [5]. The non-myelin pool is used here to describe the intra- and extra-cellular component observed in myelin water MRI experiments. This model was used in simulations of a 32-echo CPMG sequence with echo-spacing ( $\tau_{180}$ ) of 10 ms, a repetition time (TR) of 3 s, and composite refocusing pulses, identical to that which is used in our imaging experiments. The saturation effect of the refocusing pulses on the semi-solid pools was approximated using a simple model for pulsed saturation adapted for on-resonance pulses [6]. The resulting decay curves were analyzed using non-linear least-squares (NNLS) using a 120-component  $T_2$  model with logarithmically-spaced values and no regularization [7]. To simulate pathological situations observed in MS, the model parameters were modified in three ways: *i*) “pure demyelination” (absolute decrease of myelin and myelin water), *ii*) “pure edema” (increase in non-myelin water), and *iii*) “combined demyelination and edema” (decrease of myelin and myelin water with concurrent increase in non-myelin water). Since enlargement of compartmental spaces can result in increases in the  $T_2$  of the water in that compartment [8], the models with “edema” were also simulated with modest increases (up to 20%) of  $T_{2, \text{nmw}}$  in addition to increases in the non-myelin water content. Furthermore, myelin breakdown has the potential to cause increases in the diffusion-mediated movement of water between the myelin and non-myelin compartments: the impact of a 20% change in the rate was also evaluated in the models with “demyelination”. We evaluated the subsequent impact on spectral metrics such as the total observed signal ( $S_{\text{total}}$  = integral of the entire spectrum), the myelin water fraction (MWF = observed signal with  $T_2 < 50\text{ms}$  / total observed signal), and the average  $T_2$  of the spectrum ( $\langle T_2 \rangle$ ) and each peak (short:  $\langle T_2 \rangle_{\text{short}}$ , intermediate:  $\langle T_2 \rangle_{\text{interm}}$ ).

## Results:

The saturation effects of the refocusing pulses in these simulations were negligible, confirming the relative insensitivity of the CPMG signal to semi-solid tissue component saturation. The observations for the models of pathology are summarized in Table 1. The total observed spectrum signal  $S_{\text{total}}$  varied directly with total input magnetization in all cases, and the average  $T_2$  of the short peak ( $\langle T_2 \rangle_{\text{interm}} = 12$  ms) did not change in any of the cases; both are omitted from Table 1.

## Discussion:

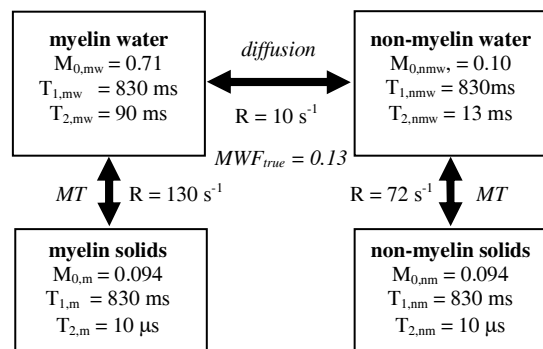
We have modeled the effect of pathology in simulations of CPMG experiments using a four-pool model of WM, providing insight on the analysis of  $T_2$  spectra from a forward-problem perspective, assuming that bovine WM can be taken as a reasonable model for human WM. Our simulations reflect the negligible semi-solid saturation effect of composite refocusing pulses widely used in CPMG imaging sequences. All pathological models resulted in effective decreases in the observed MWF. However, changes in the log-averaged  $T_2$  values of the spectrum ( $\langle T_2 \rangle$ ) and of the intermediate peak ( $\langle T_2 \rangle_{\text{interm}}$ ) differed for each of the models: the “pure demyelination” model resulted in increases in both, while the values remained stable for the “pure edema” model. The “combined” model showed a mitigated increase in the  $\langle T_2 \rangle$  and  $\langle T_2 \rangle_{\text{interm}}$ ; interestingly, the associated change in  $\langle T_2 \rangle$  for this model is mostly mediated by the myelin water, such that a large change in  $\langle T_2 \rangle$  still reflects true demyelination while slight changes in  $\langle T_2 \rangle$  may indicate edema. Increases in the non-myelin water  $T_2$  ( $T_{2, \text{nmw}}$ ) acted as a confounding factor to these distinctions, making the “edema” model resemble the “demyelination” model. The slight increase in the exchange between myelin and non-myelin water compartments results in over-estimation of the MWF decrease. Our results highlight the changes observed in the multi-component  $T_2$  spectrum under specific pathology models, and set objectives for the precision required to identify such pathological effects in MRI experiments.

**Table 1. Summary of observations for the pure demyelination, pure edema, and combined models.**

Model	Baseline estimate	“Pure demyelination” $\Delta M_{0, \text{mw}} = -20$ to $-100$ %	“Non-myelin edema” $\Delta M_{0, \text{cw}} = +10$ %, $+20$ %	“Demyelination with edema” $\Delta M_{0, \text{mw}} = -20$ to $-80$ % $\Delta M_{0, \text{cw}} = +5$ % to $+20$ %
$\langle T_2 \rangle$	68 ms	Increase of 5-22 ms	increase of 1-2 ms	increase up to 17 ms with MWF decrease increase of 1-2 ms with “edema”
$\langle T_2 \rangle_{\text{interm}}$	84 ms	increase of 1.5-7 ms	no effect ( $<0.5$ ms)	no effect ( $<0.5$ ms)
MWF	0.106	decreases of 18%, 47% and 100% correlated with input myelin water	decreases of 10% and 18% with increasing edema	decreases of 8-80 % correlated strongly with input myelin water, weakly with edema
Effect of $T_{2, \text{nmw}}$ increase with “edema”	(n/a)	(no effect)	corresponding increase in $\langle T_2 \rangle$ and $\langle T_2 \rangle_{\text{interm}}$	increase in $\langle T_2 \rangle$ and $\langle T_2 \rangle_{\text{interm}}$ now dominated by change in $T_{2, \text{nmw}}$
Effect of 20% increase in exchange with “demyelination”	(n/a)	decreases of 22%, 49% and 100% correlated with input myelin water	(no effect)	slight over-estimation of decrease in MWF

## References

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**Figure 1. Illustration of the four-pool model, with pool parameters. Exchange is indicated by the normalized exchange rate between each pair of pools. (MT = magnetization transfer)**