

Investigation of Relationship between Free-Water T1 and Age in Human Cortical Bone Employing Short-TE 1H-MRI at 1.5T

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Target Audience

Researchers, scientists, clinicians and students who work in the field of quantifying cortical bone using MRI techniques

Purpose

Less than thirty percent of human cortical bone volume is composed of water which plays a pivotal role in its mechanical competence, from which only twenty percent appears as free water, occupying larger pores such as Haversian and Lacuno-canalicular systems [1]. The volume of cortical pores increases by aging and some bone related diseases such as Osteoporosis, leading to increase free water content; therefore, quantification of free water is a reliable measure to assess cortical bone porosity. Commercially available short-TE (STE) pulse sequences in clinics with the echo-time (TE) in the range of 0.5-1msec are shown to be appropriate candidates to acquire enough signal from protons residing in large pores of human cortical bone, leading to successful quantification of free water T₁ values [2]. In the present work, we investigated relationship between STE-based cortical bone T₁ values and age, studies on a group of healthy volunteers at 1.5 T.

Materials and Methods

Subjects: Eight normal volunteers, 3 males and 5 females (20-57yrs with the mean age of 37.4yrs), were incorporated into this study.

Image Acquisition: Mid-tibia images were acquired using STE pulse sequence with two different TR values on a 1.5T MR scanner (Siemens, Magnetom Avanto 18 channel) to implement previously proposed dual-TR technique for cortical bone T₁ quantification *in-vivo* [2, 3]. The imaging parameters are selected to be: TR₁/TR₂/TE = 20/60/1.19msec, field-of-view (FOV) = 267×267mm², spatial resolution = 0.8×0.8mm², slice thickness = 5mm, flip angle = 20°, total scan time of about 20 minutes, using an 8-channel Tx/Rx knee coil (an example is shown in Fig. 1).

T₁-Quantification: Steps of quantification are as follows: (1) manual segmentation of the whole cortical bone at each of the two images with different TRs; (2) computation of the ratio value (r), as in Eq. 1, by dividing the mean signal intensities of the segmented cortical bone acquired from long-TR (TR₂) and short-TR (TR₁) images, respectively; (3) calculation of cortical bone T₁-value at each imaging slice by solving Eq. 1 using nonlinear solver in MATLAB 7.14 (The MathWorks) [2];

$$r = \frac{1 - \exp(-TR_1/T_1)}{1 - f_2 \exp(-TR_1/T_1)} / \frac{1 - \exp(-TR_2/T_1)}{1 - f_2 \exp(-TR_2/T_1)} \quad \text{Eq. 1}$$

and (4) calculation of the average T₁-values for each subject and from ten different slices. As quantification of T₁-values are very sensitive to f₂ – a parameter which characterizes the longitudinal magnetization as a function of pulse duration to the tissue T₂^{*} (τ/T₂^{*}) [4] – it must be carefully determined based on Bloch equation simulation employing T₂^{*} value of the cortical bone extracted from the literature at 1.5T, and parameters of the actual excitation pulse such as pulse shape and flip angle.

Evaluation of signal-to-noise ratio (SNR): SNR values, computed by dividing the mean signal intensities from segmented cortical bone in high-SNR (long-TR) images to the mean signal intensities from a region-of-interest (ROI) placed in the background noise, were in acceptable range for all slices (SNR>12). Steps to quantify T₁ and SNR values were shown in Fig. 3.

Results

Table 1 shows results for quantitative measurement of T₁-values in eight healthy volunteers using STE pulse sequence. Measurements were performed for both genders, resulting in the mean T₁-values of about 202.81msec for human cortical pore (free) water at 1.5T. Such T₁ quantity has been reported in the range of 380-775msec and 200-400msec at 4.7T [5] and 3T, respectively, showing rationale results achieved with the STE pulse sequence at 1.5T. T₁-values are strongly correlated with age as shown in Fig. 2 (R²=0.75, p<0.0001).

Discussions and Conclusions

Results suggest successful application of STE-MRI for accurate quantification of cortical bone T₁-values, with the advantages of total scan-time of about half of ultra-short TE's (UTE) pulse sequences, widespread clinical availability and cost-effective procedure, meaning that STE sequences can be utilized as proper alternatives in quantifying cortical bone parameters *in-vivo* [2]. Also this suggests that quantification of pore (free) water T₁ using STE is a reliable measure of cortical bone deterioration with age. Furthermore, our results follow the well-known theory describing cortical bone relaxivity as a function of its geometrical characteristics, 1/T₁ ∝ (S/V) in which S and V are surface area and volume of the pore, respectively, meaning as surface-to-volume ratio decreases for larger cortical porosities due to aging, we see consistent increase in the T₁-values (R²=0.75, p<0.0001) [3].

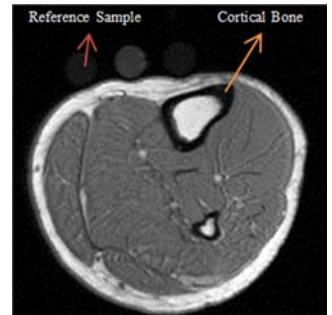


Fig. 1 A sample image of the mid-tibia acquired by the STE pulse sequence

Table 1. Quantitative measurement of T₁ and SNR in 8 normal subjects

Subject	Age	Gender	T ₁	SNR
1	20	F	120.95	14.12
2	28	F	162.92	13.60
3	29	F	183.47	14.10
4	34	F	220.37	17.52
5	38	M	233.46	18.25
6	46	M	204.65	13.44
7	47	F	206.20	12.95
8	57	M	290.50	15.83

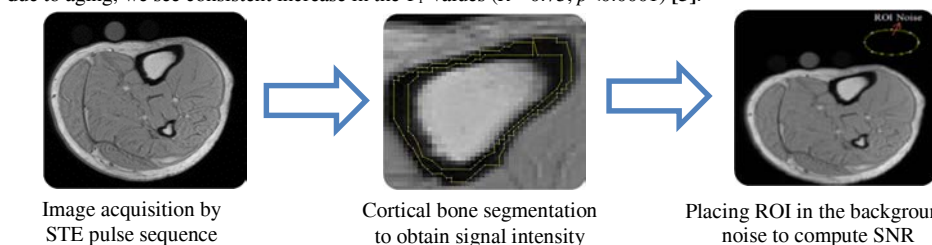


Fig. 3 Steps to acquire signal intensities from manually segmented cortical bone and background noise for T₁/SNR quantification purposes.

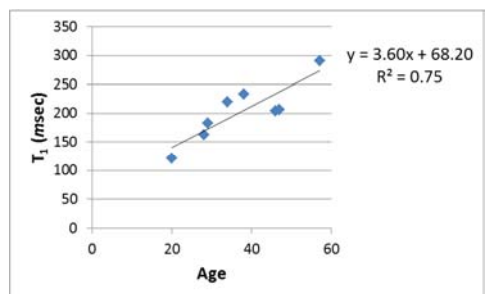


Fig. 2 Correlation between T₁ and Age in 8 healthy volunteers using STE pulse sequence (p<0.0001)

References: [1] Du J. *ISMRM 21* (2013) [2] Akbari A. *et al, ESMRMB 30* (2013) [3] Saligheh Rad H. *et al, NMR Biomed*, 23: 1-11 (2011) [4] Sussman M. *et al, MRM*, 40:890-899 (1998) [5] Horch RA. *Thesis* (2011)

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