

Bone Mineral and Matrix Densities Measured by Solid-State ¹H and ³¹P MRI

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Introduction: Osteoporosis and osteomalacia are distinct bone disorders that both manifest as reduced apparent bone mineral density (BMD), and are indistinguishable by in vivo densitometric techniques. Osteoporosis involves both bone matrix and mineral loss, while osteomalacia involves mineral deficiency only. The differentiating factor is bone mineralization [1]: mineral per volume of bone matrix. MRI of bone is made difficult by unfavorable magnetic relaxation properties, but advances in solid-state ³¹P and ¹H MRI have led to the possibility of quantitative measurement of bone mineral ³¹P and collagen-bound water ¹H densities, respectively [2-4]. The objective of this work was to quantify relative bone mineral ³¹P and matrix-bound water ¹H densities on a clinical MRI scanner to infer bone mineralization in specimens from human donors.

Theory: Water ¹H NMR signal in bone at 3T occurs in two compartments: long T₂ > 1 ms, corresponding to free water in pores ('pore water'); and short T₂ ~ 300-400 μs, corresponding to motionally restricted water bound to bone matrix collagen ('bound water') [5]. Bone matrix density is proportional to bound water and inversely proportional to pore water [6]. Adiabatic RF pulses saturate bound water protons (longitudinal magnetization M_z ~ 0) while partially inverting pore water protons (M_z < 0). After an inversion time delay (TI), pore water M_z will be nulled (M_z ~ 0) and bound water M_z will have recovered (M_z > 0) due to partial longitudinal (T₁) recovery. Imaging can then be performed, yielding an image predominantly of bound water.

Methods:

Specimens: Tibial cortical bone specimens, 36 mm long, were taken from 16 human donors (27-97 y/o, NDRI). Donors with bone-demineralizing disorders were excluded. Each bone was scanned with suitable ³¹P and ¹H reference intensity phantoms.

Hardware: MRI scanning (¹H at 3T, ³¹P at 7T) was performed on Magnetom whole-body MRI scanners (Siemens, Erlangen, Germany). A custom RF coil was used (7T ³¹P: 120.3 MHz, 3T ¹H: 123.3 MHz). Three fiducial markers of triethyl phosphate (C₂H₅O₄P) were affixed to the coil for registration of an RF field map, acquired on a water phantom at 3T.

MRI: ³¹P relaxation times were measured in each specimen. Each bone was imaged using ³¹P zero echo time imaging with pointwise encoding time reduction with radial acquisition [7] (³¹P ZTE, res=2 mm³ isotropic, flip angle=5°, gradient=36.7 mT/m, TR=20 ms, NEX=100, 5000 half-projections, PETRA radius=5 points, scan time 3h3m) and ¹H Single Adiabatic Inversion Recovery Rapid ZTE-PETRA (¹H SIR-rZTE, res=1 mm³ isotropic, gradient=18.3 mT/m, TR=300 ms, inversion pulse BW=5 kHz, duration=5 ms, TI=100 ms, 7 readouts/excitation, flip angle=24-40°, 10000 half-projections, PETRA radius=4 points, scan time 26m45s).

Density Quantification: Bones, reference intensities, and fiducial markers were masked and thresholded. Images were corrected for transmit and receive B₁ inhomogeneity and differences in relaxation times between bone and reference. Relative ³¹P and bound water ¹H densities in bone were expressed relative to the reference intensity using Eq. 1, where f_{xy} and f_z describe the response of transverse and longitudinal magnetization to a square pulse [8].

pQCT: A 2D image was acquired at the center of each bone with 0.4x0.4x2.3-mm³ resolution using a Stratec XCT 2000 scanner (Orthometrix, White Plains, NY). BMD was quantified using the scanner software's in vivo tibial measurement protocol.

μCT: A 3D image of each bone was acquired with 9-μm³ isotropic resolution using a Bruker μCT scanner (Bruker, Kontich, Belgium), segmented, and binarized. Porosity was quantified as the ratio of pore volume to total volume, both excluding the endosteal cavity.

Analysis: Correlations were examined by regression calculated using least squares.

Results: Maps of relative ³¹P and bound water ¹H densities are shown in Fig. 1. Both densities correlate negatively with age and porosity, and positively with pQCT density. The relative densities correlate with each other (R² = 0.59, p < 0.001), and their ratio, which represents bone mineralization, is not correlated with age, pQCT density, or porosity. R² values of other correlations are shown in Table 1.

Discussion and Conclusions: MRI and pQCT cannot resolve individual pores, so an age-related increase in porosity manifests as a decrease in apparent density as long as the mineralization is invariant. Bone mineralization can be inferred from pQCT BMD by dividing by (1-porosity); this was uncorrelated with age, indicating full mineralization. Consistent with this observation, both MRI-based densities correlate with pQCT density and porosity. Further studies in demineralized bones will test the ability of this MRI-based method to distinguish osteoporotic- from osteomalacic-type disruptions of bone homeostasis.

References: [1] Meunier PJ, et al. Bone 1997;21:373-7. [2] Ackerman JL, et al. MRM 1992;25:1-11. [3] Anumula S, et al. Bone 2010;46:1391-9. [4] Robson MD, et al. MRM 2004;51:888-92. [5] Horch RA, et al. MRM 2010;64:680-7. [6] Ong HH, et al. JBMR 2012;27(12):2573-81. [7] Grodzki, et al. MRM 2012;67:510-8. [8] Sussman MS, et al. MRM 1998;40:890-9.

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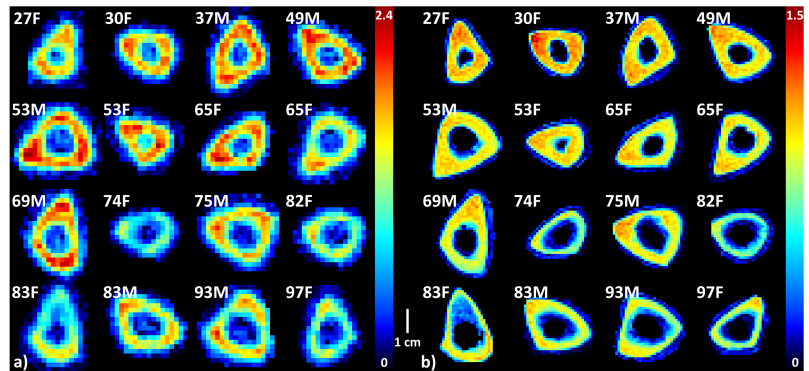


Fig. 1: (a) Relative ³¹P and (b) bound water ¹H density maps. Age and gender are indicated above each bone.

	Age	Porosity	pQCT	pQCT/BVF	³¹ P T ₁	r ³¹ PD	r ³¹ PD/BVF	rBWD	rBWD/BVF
RDR	2.5x10 ⁻³	9.9x10 ⁻⁷	0.09	0.18	0.33*	0.50**	0.62***	8.6x10 ⁻³	0.02
rBWD/BVF	0.59***	0.40*	0.41*	0.04	0.21	0.59**	0.51**	0.94***	
rBWD	0.70***	0.63***	0.50**	0.01	0.19	0.59**	0.45**		
r ³¹ PD/BVF	0.26*	0.15	0.36*	0.17	0.47**	0.96***			
r ³¹ PD	0.39*	0.32*	0.46**	0.11	0.45**				
³¹ P T ₁	0.02	0.09	0.51**	0.45**					
pQCT/BVF	4.2x10 ⁻³	0.02	0.39*						
pQCT	0.29*	0.48**							
Porosity	0.54**								

Table 1: Correlation matrix of R² values (*p<0.05, **p<0.005, ***p<0.0005). RDR, relative density ratio; rBWD, relative bound water density; r³¹PD, relative ³¹P density; BVF, bone volume fraction = 1-porosity.

$$S(\vec{r}) \propto \rho(\vec{r}) \frac{1 - \exp\left(-\frac{TR}{T_1}\right)}{1 - f_z(\vec{r}) \exp\left(-\frac{TR}{T_1}\right)} f_{xy}(\vec{r}) \exp\left(-\frac{TE}{T_2}\right) \hat{B}_1(\vec{r})$$

Equation 1: Steady-state signal equation for ZTE. Terms f_{xy} and f_z depend on T₂ and RF pulse duration and amplitude. The signal equation is more complex for the multiple-readout SIR-rZTE sequence.