

NMR Relaxation Time Measurements of ^{129}Xe Dissolved in Tissue and Blood: Toward MR Imaging of Laser-Polarized ^{129}Xe in Tissue

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Nuclear magnetic resonance relaxation measurements of ^{129}Xe in tissue homogenates and blood were performed to explore the feasibility of magnetic resonance (MR) imaging of laser-polarized (LP) ^{129}Xe in tissues. Laser polarization of ^{129}Xe by optical pumping and spin exchange increases the signal obtained from ^{129}Xe to a sufficient level that, after inhalation of LP ^{129}Xe , it may be detected by MR throughout the body. Image signal-to-noise ratio (SNR) will depend on longitudinal (T1) and transverse (T2) relaxation times of ^{129}Xe in tissues and blood.

^{129}Xe T1 and T2 were measured at 9.4 T and 10 degrees C in rat whole blood and homogenates of rat liver, brain, kidney, and lung at varying oxygenation levels. T1 values ranged from 4.4 +/- 0.4 to 22 +/- 2 s. T2 values ranged from 1.4 +/- 1.0 to 7.00 +/- 0.09 ms. Oxygenation dependence of the relaxation times varied between tissues. The observed oxygen dependence suggests relaxation due to paramagnetic deoxyhemoglobin dominates that due to free oxygen in tissues with sufficiently high fractions of blood. In addition, T1 and T2 of ^{129}Xe in separated blood plasma were measured at both 9.4 and 1.89 T to explore the dependence on field strength. T1 of ^{129}Xe in plasma decreased from 46 +/- 8 to 20 +/- 3 s and T2 increased from 11 +/- 2 to 127 +/- 8 ms when the field strength decreased from 9.4 to 1.89 T.

When an inhalation model of gas delivery to tissue is used, the expected intrinsic SNR for LP ^{129}Xe in gray matter at 9.4 T is approximately 3% of the conventional proton intrinsic SNR at 1.5 T. Short ^{129}Xe T2 values (at 9.4 T) and the finite amount of magnetization available for imaging LP ^{129}Xe further reduce the expected image SNR compared to conventional proton imaging. Several common pulse sequences were modeled to predict their relative effectiveness for imaging LP ^{129}Xe at both 9.4 and 1.89 T. Each modeled sequence predicted higher SNR at 1.89 than at 9.4 T.