

WORKSHOP PRESENTATION

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Saturation recovery allows T_1 mapping in the human heart at 7T with a commercial MRI scanner

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Background

Myocardial T_1 mapping at 1.5T and 3T distinguishes powerfully between normal and diseased tissue with focal and diffuse pathology. We recently reported the first human myocardial T_1 s at 7T using the ShMOLLI +IE inversion recovery sequence. Yet even using a unique 7T scanner with 16kW RF output, perfect magnetization inversion was impossible. We now introduce a saturation recovery method to enable myocardial T_1 mapping with standard commercial 7T MRI scanners.

Methods

The saturation recovery single-shot acquisition (SASHA, Chow, 2013) sequence was modified for 7T by using: an optimised train of 4xHS8 pulses to saturate and 10 FLASH readouts with saturation delays (T_S): non-saturated, 100, 200, 300, 400, 500, 600, 650ms, 1hb + 100ms and 1hb + 700ms in a 12 heart-beat breath-hold. Data were acquired with a Siemens 7T MRI scanner (with 8kW RF), an 8-element cardiac coil and ECG gating. Signals were fitted pixelwise to “s (T_S) = A - B exp(- T_S / T_1)”.

10 healthy subjects (males, 22-45yrs, 70-84kg) were recruited according to ethics regulations. For each subject, coil tuning, B_0 -shims, B_1 -shims and the central frequency were optimised over the left ventricle. Then 7T SASHA native (i.e. non-contrast) T_1 mapping was performed in SA and HLA views.

In three subjects, post-contrast T_1 maps were acquired ~5min after 2 peripheral bolus injections of Dotarem with a power injector (Accutron MR, MEDTRON).

Results

“7T SASHA” T_1 s were validated against IR-SE reference T_1 s: values agreed to within 6% for readout flip angles $\leq 25^\circ$ (Fig. 1).

In-vivo, the native 7T SASHA T_1 s in the interventricular septum were 1939 ± 73 ms. Native T_1 s in the LV blood pool showed strong artefacts, likely due to blood flow. The post-contrast T_1 s were 999, 1107 and 1674ms in myocardium and 472, 567 and 966ms in blood for

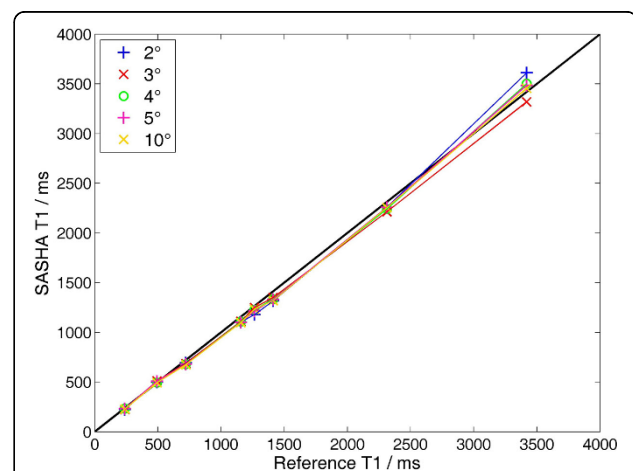
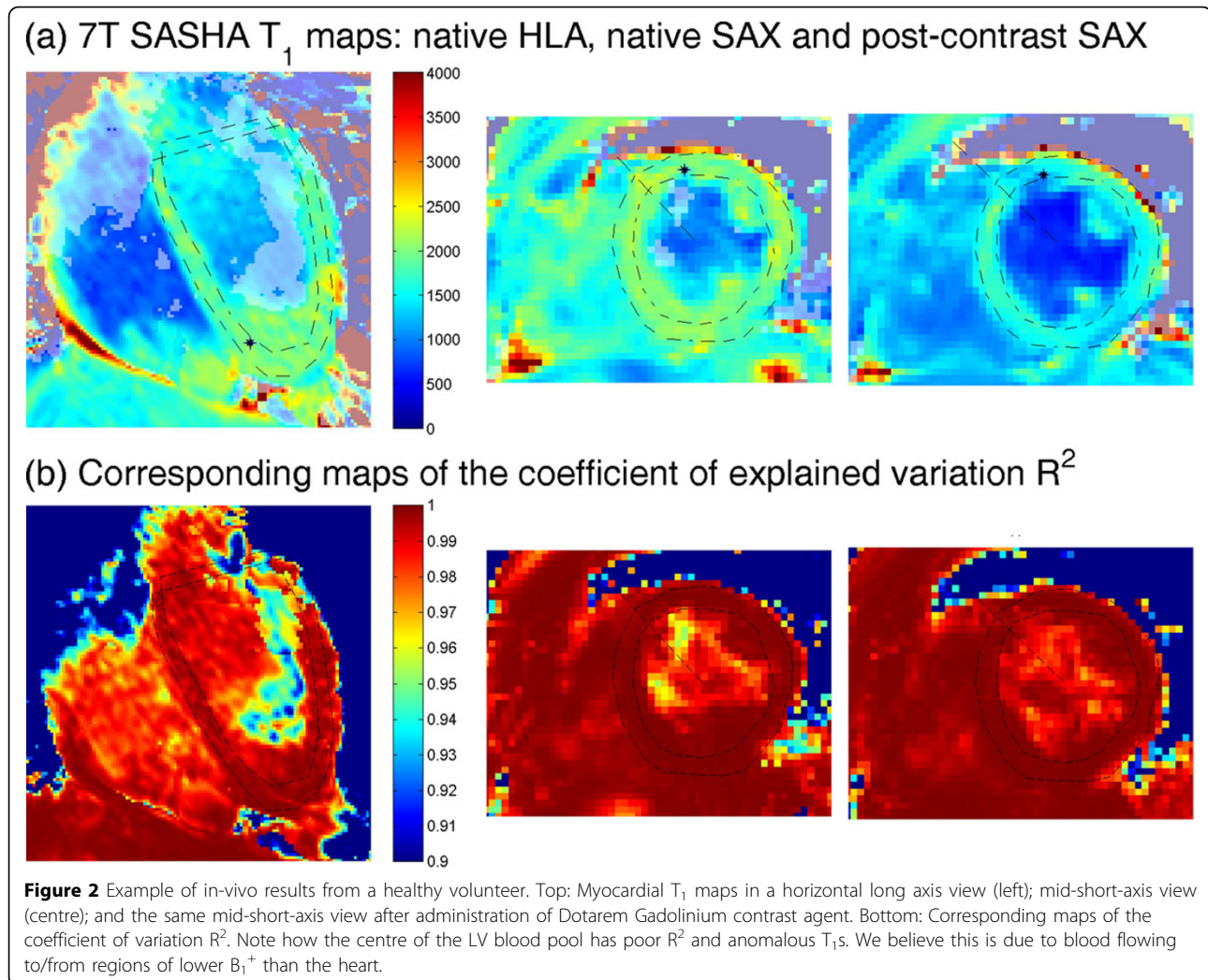


Figure 1 Phantom validation of 7T SASHA sequence T_1 s against inversion-recovery-spin-echo (IR-SE) reference values in a phantom comprising tubes of NiCl_2 -doped agar and carrageenan. Reference data were acquired in a 32-channel head coil (Nova Medical) to ensure sufficient B_1^+ for reliable inversion across the phantom. 7T SASHA data were acquired using the 8-element cardiac coil. For FLASH readout flip angles $< 20^\circ$, the 7T SASHA T_1 s are within 6% of the IR-SE reference T_1 s. Spin echo (SE) reference T_2 s were 50-300ms in this phantom.



Dotarem doses of 2x 50, 50 & 25 and 2x 13.5 $\mu\text{mol/kg}$ in the three subjects respectively. Post-contrast T_1 maps were acquired too soon after bolus infusion to permit calculation of extra-cellular volumes.

These myocardial T_1 s agree with our ShMOLLI+IE finding of a myocardial $T_1 = 1925 \pm 48$ ms. However, with ShMOLLI+IE we had to use a 4-parameter model-based fitting procedure to correct for imperfect inversion, read-out induced saturation and spin history effects. In contrast, with 7T SASHA, it is now possible to achieve comparable T_1 s using a simple 3-parameter fit (on the scanner). Note that these considerations at 7T are different to the well-known differences between MOLLI and SASHA T_1 s at 1.5 and 3T caused by imperfect inversion, T_2 relaxation, and magnetization transfer.

Conclusions

Saturation recovery allows T_1 mapping in the human heart using a commercial 7T MRI scanner. T_1 s from 7T

SASHA with 3-parameter fitting and ShMOLLI+IE with 4-parameter fitting are comparable in normal volunteers at 7T. Our findings hold promise for wider clinical applications of T_1 mapping at ultra-high fields.

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