A surface crusher coil for human cardiac phosphorus (³¹P) MR spectroscopic imaging study at 7 tesla

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Introduction: ³¹P-MRS provides direct insights into myocardial energy supply (ATP, ADP, phosphocreatine (PCr) and inorganic phosphate)¹. An initial study demonstrated that 7T cardiac ³¹P-MRS has 2.8x greater SNR and higher quantification precision than at 3T². Additional RF pulses for outer volume suppression are often inserted into main MR spectroscopy pulse sequence to suppress potential contaminating signals (e.g. overlying skeletal muscle may contaminate myocardial ³¹P-MR spectra). However, the translation of these approaches to suppressing contamination at 7T is particularly challenged by increased RF heating of tissue at 7T. This incurs extended repetition times (TR) for MRS(I) at 7T and in some sequences prevent to achieve complete saturation with the remaining SAR. Chen and Ackerman³ introduced the surface spoiling coil in 1990: a concept that was recently further developed⁴ for lipid suppression in human brain ¹H-CSI. In this work, we introduce the first crusher coil for cardiac ³¹P-MRS at 7T. This allows us to saturate more efficiently skeletal muscle signal whilst removing the RF heating associated with RF saturation bands.

Methods: Data were acquired with a Siemens 7T scanner. Localization used a 10cm ¹H Tx/Rx RF coil (Rapid Biomedical) to acquire CINE FLASH images. ³¹P-MR spectra were acquired with a custom 10cm ³¹P Tx/Rx loop. The magnetic field generated by the crusher coil was simulated using Matlab (Mathworks) and crusher coil design was optimized. A capacitor continuously charged by a power supply unit was used to drive the current pulse in the crusher coil (100µs duration). Spoiling was timed to coincide with the existing phase encoding gradients. A 2D-CSI experiment (TR=1s, TE= 2.3 ms, slice thickness=20 mm, matrix size = 180×180 mm², res = 12×12) was performed on a two-compartment phantom with the ³¹P-RF coil and the crusher coil placed above it. In vivo 3D-CSI was then performed with the crusher coil (TR=1s, TE=2.3ms, matrix size = $240 \times 240 \times 200$ mm³,

resolution = $16 \times 16 \times 8$, 10 averages, acquisition weighted, TA = 28 min). The mean PCr skeletal muscle was calculated over voxels placed above the interventricular septum (n = 5). The residual signal obtained with crusher coil was compared with BISTRO saturation⁵ in phantom and *in vivo*. The spectral analysis was performed using a custom Matlab fitting program that includes the AMARES fitting⁶. The SAR level was reduced from $96\pm1\%$ to $16\pm1\%$ when using the crusher coil (without BSITRO pulses).



Figure 1: (A) Simulated residual signal in function of the height above the crusher coil. The residual signal was calculated for different I_{spoil}×T_{spoil}: 0.75 (blue), 1.5 (red), 2.25 (cyan), 3 (purple), 4.5 (green) and 6 A-ms (orange).



Figure 2: 2D CSI over a transverse slice with 5 saturation protocols: (i) BISTRO saturation bands (thickness = 30mm), (ii) crusher spoiling, (iii) BISTRO and crusher spoiling and (iv) crusher spoiling.

would otherwise be SAR-prohibitive e.g. adiabatic excitation for absolute quantitation, ¹H-³¹P NOE enhancement or saturation-transfer pulses for future clinical studies at 7T, without compromising the skeletal muscle suppression.

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myocardium. The flexibility offered

by using the crusher coil allows us to

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Figure 3: ³¹P-MR spectra acquired in the skeletal muscle and in the interventricular septum with different saturation protocols: no saturation (blue spectra), BISTRO saturation band (red spectra) and crusher coil (black spectra, Ispoil×Tspoil=0.9 Ams). Inset: Overlap of CSI grid on CINE FLASH images showing voxel whose spectra are plotted. Yellow strip corresponds to BISTRO saturation band. Voxel 7 was used for AMARES fitting.

References and Acknowledgements: [1] Bottomley PA, MRM, 1985 [2] Rodgers, CP, MRM, 2013 [3] Chen and Ackermann, NMR in BioMed, 1990 [4] Boer, MRM 2014 [5] Luo, MRM 2001 [6] Vanhamme JMR 1997. Funded by the Wellcome Trust and the Royal Society [098436/Z/12/Z]