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Introduction: In spectroscopic imaging (chemical shift imaging CSI), in addition to the two (or three) spatial dimensions, a spectral dimension is acquired. In contrast to normal imaging, in CSI all spatial dimensions are encoded using phase gradients requiring long experiment times. For this reason especially 3D CSI imaging is an appropriate candidate for acceleration by the parallel imaging concept. In previous work, CSI acceleration using the SENSE reconstruction algorithm has been presented [1]. However, significant artifacts could be induced by SENSE, due to the low-resolution nature of spectroscopic imaging [2]. Therefore, in this work we focused on parallel CSI acquisition using the GRAPPA [3] algorithm, which does not have this limitation. Furthermore, the flexibility of the GRAPPA approach allows one to derive the reconstruction weights either from a few extra acquired CSI data in the center of k-space or from a fast fully-encoded low-resolution reference pre-scan [4].

Methods: 2D phantom and 3D *in vivo* CSI head experiments were performed on a 1.5T clinical scanner (Avanto, Siemens, Erlangen, Germany) equipped with a 12 channel head array. The 2D CSI phantom was made up of 3 bottles containing Na-Acetate and Li-Lactate in different compartments. The acquisition parameters were: FOV=160x160mm², slice thickness 15mm, matrix=32x32, TR=1500ms, TE=135ms, bw=1kHz with 1024 spectral data points. In addition, a fast, fully-encoded low-resolution FLASH pre-scan experiment of the phantom was performed, which served as a coil calibration reference for the GRAPPA reconstruction.

The 3D head *in-vivo* acquisition parameters were FOV=200x160x120mm³, matrix=16x16x8, TR=1070ms, TE=135ms, bw=1.6kHz with 1024 spectral points.

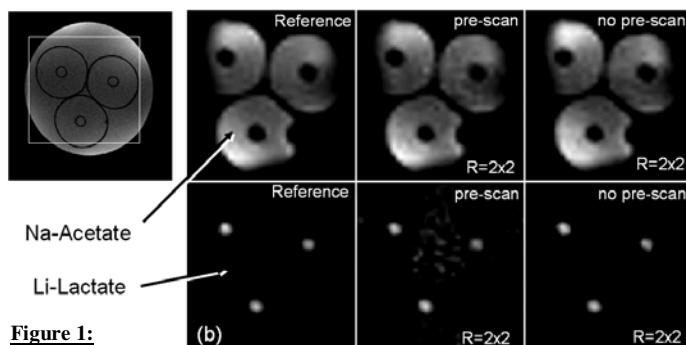


Figure 1:

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