## **Ouantitative Susceptibility Map Reconstruction with Magnitude Prior**

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INTRODUCTION: Quantitative Susceptibility Mapping (QSM) aims to quantify tissue magnetic susceptibility with applications such as tissue contrast enhancement [1], venous blood oxygenation [2], and iron quantification [3]. The magnetic susceptibility  $\chi$  maps to the observed phase shift via a well-understood transform; hence  $\chi$  inversion benefits from additional regularization [4]. Here we propose enhanced regularization for  $\chi$  inversion by incorporation of magnitude priors. Since the data acquisition step for QSM yields both phase and magnitude data, the inverse problem can be better conditioned if the magnitude is incorporated as a prior. By encoding spatial priors derived from a magnitude image into an  $\ell_1$  regularization scheme via the Focal Underdetermined System Solver (FOCUSS) algorithm [5], we report high quality QSM on a numerical phantom with four-fold improvement in RMSE when magnitude priors were applied. We also demonstrate the application of the method on in-vivo data at 7T. THEORY: The system of linear equations  $\delta$ =F<sup>-1</sup>DF $\chi$  defines our ill-posed deconvolution task where  $\delta$ = $\varphi$ /( $\gamma$ -TE- $\theta$ <sub>0</sub>) is the normalized field map, D is the susceptibility kernel in k-space, F is the Fourier transform operator,  $\chi$  is the susceptibility vector,  $\varphi$  is the unwrapped phase,  $\gamma$  is the gyromagnetic ratio, TE is the echo time and  $\theta$ <sub>0</sub> is the main field strength. We assume that  $\chi$  shares tissue contrast boundaries with the magnitude image, and is therefore expected to have similarly sparse spatial gradients as the magnitude; this prior knowledge can be imposed on the reconstruction via the regularized FOCUSS algorithm [6]. Letting  $\partial_x \chi$  denote the spatial gradient along x, the k-th step of the iterative algorithm is as follows: set  $W_{k+1}$ -diag( $|\partial_x \chi_k|^{0.5}$ ), solve  $q_{k+1}$ -argminq || $V_x$ -F $\delta$ -DF $W_{prior}$ -W<sub>k+1</sub>q||q||2 +  $\chi$ -q||2, and update  $\partial_x \chi_{k+1}$ -w<sub>prior</sub>-W<sub>k+1</sub>q<sub>k+1</sub>. Here,  $V_x$  is a diagonal matrix that acts as gradient operator in k-space due to  $V_x$ ( $\alpha$ ,  $\alpha$ )

three compartments: a rectangular prism ( $\chi$ =1ppm), a cylinder ( $\chi$ =0.047ppm) simulating graywhite matter susceptibility difference [1], and a 2-pixel wide vessel ( $\chi$ =0.4ppm). The vessel has three segments; i) along  $B_0$  (z-direction), ii) in-plane part, and iii) a 35° slanted segment, which is perpendicular to the magic angle of 55° and therefore poses the most challenging inversion geometry. We also created a magnitude image with shared boundaries, but with different compartment intensities. Starting from the true susceptibility, we forward simulated the field map by convolution and corrupted it with complex valued Gaussian noise so that the noisy field map had 17.9% normalized root mean squared error (RMSE) relative to the noise-free case. We tested the FOCUSS algorithm without a prior (by setting  $W_{prior} = I$ ) and with magnitude prior and used an optimal  $\lambda$  setting ( $\lambda$ =10<sup>-2</sup> without prior and  $\lambda$ =10<sup>-3</sup> with prior) to reconstruct the susceptibility from the noisy field map. Second, we tested the FOCUSS on in-vivo data. At 7T, a 3D GRE sequence was used to acquire axial images with 0.33 mm in-plane resolution, 1.0 mm slice thickness and FOV of 192×168×64 mm<sup>3</sup> for a TE of 10 ms on a young (26 years, female), healthy subject. After high-pass filtering the phase with a Hanning filter of size 64×64, the susceptibility distribution was reconstructed from the field map using the FOCUSS algorithm  $(\lambda=10^{-5})$  by using the magnitude as a prior. We report susceptibility differences  $\Delta \chi = \chi_{vessel}$  $\chi_{tissue}$  for a selected vessel and  $\Delta \chi = \chi_{putamen} - \chi_{tissue}$  for the putamen by manually generating interior and surrounding tissue masks for averaging.

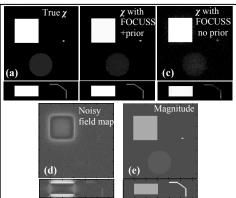


Fig.1 (a) Axial and sagittal views of true χ (b) QSM using FOCUSS with magnitude yielded 1.3% RMSE (c) FOCUSS without a prior returned 5.2% RMSE (d) Noisy field map (e) Magnitude image

**RESULTS:** For the numerical phantom, susceptibility reconstruction without a prior resulted in 5.2% RMSE whereas using the magnitude prior yielded 1.3% error (Fig. 1). We note that the slanted segment of the vessel is almost invisible in the field map due to the ill-posed kernel, but FOCUSS with magnitude prior successfully recovered this segment. Fig. 2 depicts 7T QSM results obtained after taking maximum intensity

projection (MIP) over 4 slices that contain the vessel of interest. We computed  $\chi_{vessel}$  by taking the mean of the MIP image inside the vessel ROI (Fig. 2b) and  $\chi_{tissue}$  by taking the mean susceptibility of pixels inside the tissue

ROI across 4 slices. In this case,  $\Delta \chi$  was estimated to be 0.34ppm. Fig. 3 presents QSM results for the putamen obtained by taking the average susceptibility over 6 slices. After computing average  $\chi_{putamen}$  and  $\chi_{tissue}$  inside the ROIs in Fig. 3b-c,  $\Delta \chi$  was estimated to be 0.011ppm. Our result falls within the range of putamen susceptibility values (0 to 0.054ppm) reported in [3] for subjects between 20 and 30 years.

**CONCLUSION:** By making use of magnitude information to add spatial priors to  $\ell_1$  regularization, we demonstrate high quality QSM on numerical and in-vivo data. In addition to estimating venous oxygenation, the algorithm can be used for quantification of susceptibility inside iron-rich brain structures.

REFERENCES:[1] Duyn JH et al., PNAS 2007;104(28):11796-11801 [2] Fan AP et al., ISMRM 2010;693 [3] Liu T et al., ISMRM 2010;4364 [4] Liu J et al., ISMRM 2010;4996 [5] Gorodnitsky IF et al., IEEE T Signal Proces 1997;45(3):600-616 [6] Cotter SF et al., IEEE T Signal Proces 2005;53(7):2477-2488

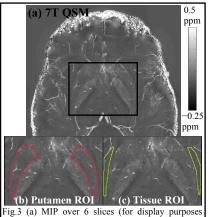
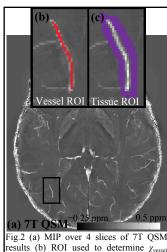


Fig.3 (a) MIP over 6 slices (for display purposes only) of 7T QSM results (b) ROI for  $\chi_{putamen}$  (red) (c) Tissue ROI (green)



(red) (c) Tissue ROI (purple)

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