

Quantitative Susceptibility Mapping with Magnitude Prior

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Quantitative Susceptibility Mapping (QSM)

- Quantitative Susceptibility Mapping (QSM) aims to quantify tissue magnetic susceptibility with applications such as,
 - ❖ Tissue contrast enhancement¹
 - ❖ Estimation of venous blood oxygenation²
 - ❖ Quantification of tissue iron concentration³

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$$\delta = \mathbf{F}^{-1} \mathbf{D} \mathbf{F} \chi$$

F: Discrete Fourier Transform matrix

D: susceptibility kernel in k -space

$$\delta = \frac{\varphi}{\gamma \cdot \mathbf{TE} \cdot \mathbf{B}_0} : \text{normalized field map}$$

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measured

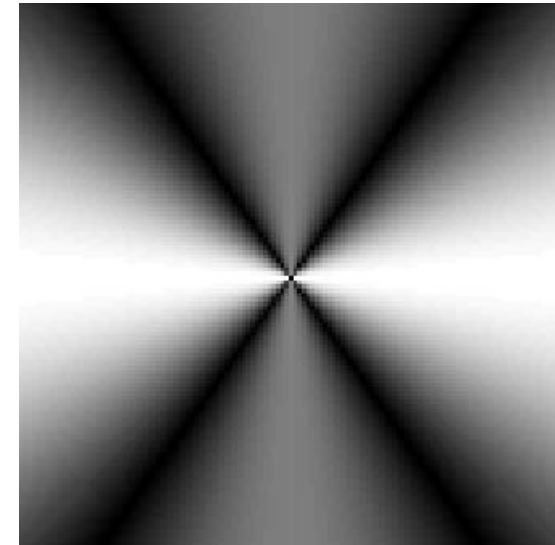
to be estimated

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- Estimation of the susceptibility map χ from the unwrapped phase φ involves solving an inverse problem, $\delta = \mathbf{F}^{-1}\mathbf{D}\mathbf{F}\chi$
- The inversion is made difficult by zeros on a conical surface in susceptibility kernel \mathbf{D}

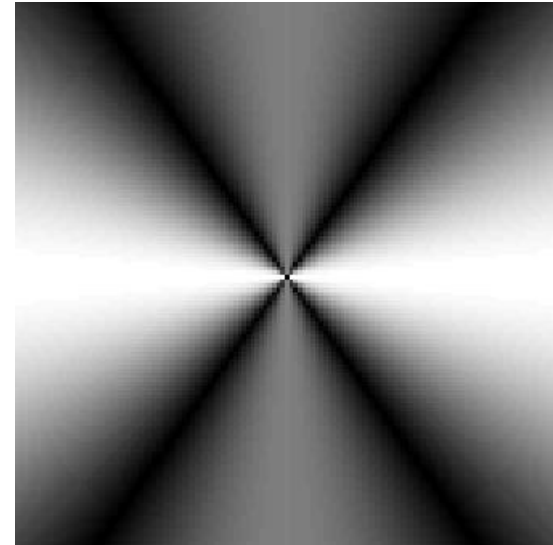
$$\mathbf{D} = \begin{pmatrix} 1 & \mathbf{k}_z^2 \\ 3 & \mathbf{k}^2 \end{pmatrix}$$

$|\mathbf{D}|$



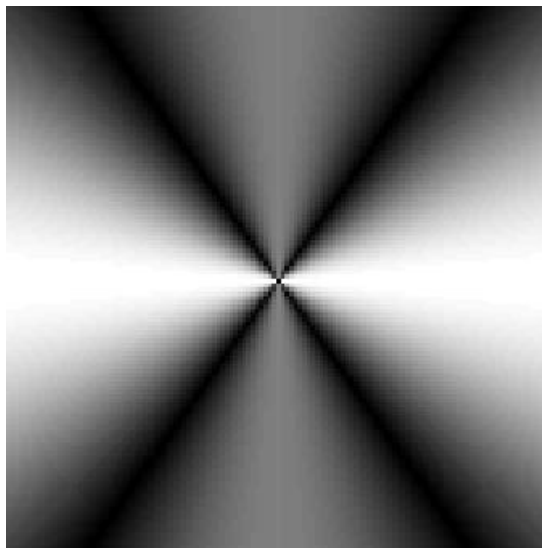
Regularized Inversion for QSM

|D|

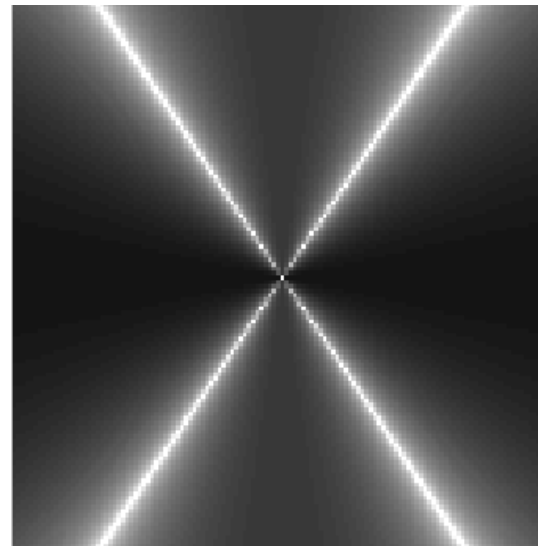


Regularized Inversion for QSM

$|\mathbf{D}|$



$\log|\mathbf{D}^{-1}|$

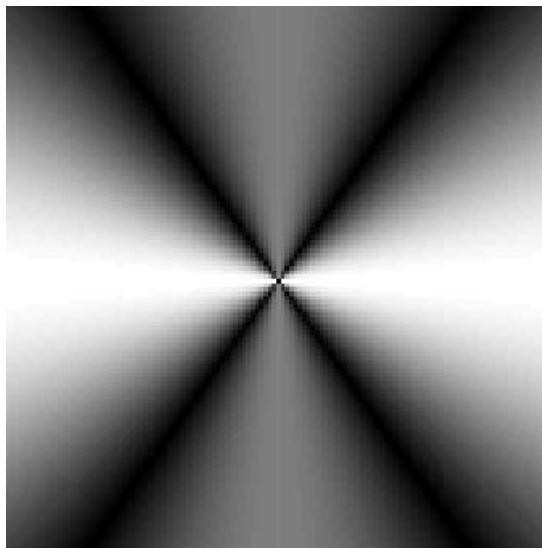


$$\mathbf{F}\mathbf{D}^{-1}\mathbf{F}\delta = \chi$$

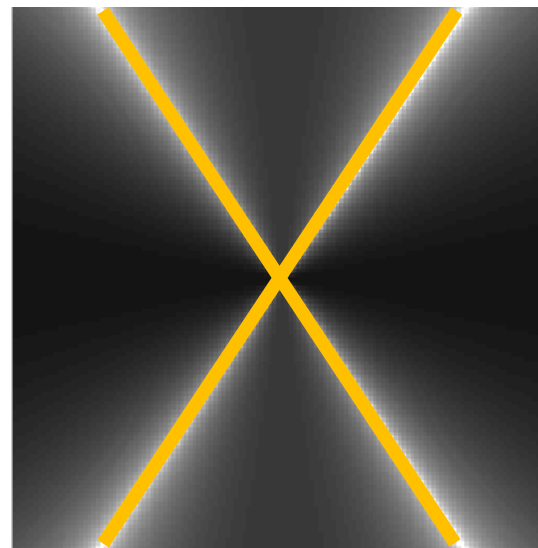
- Solving for χ by convolving with the inverse of \mathbf{D} is not possible, as it diverges along the magic angle

Regularized Inversion for QSM

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$\log|\mathbf{D}^{-1}|$

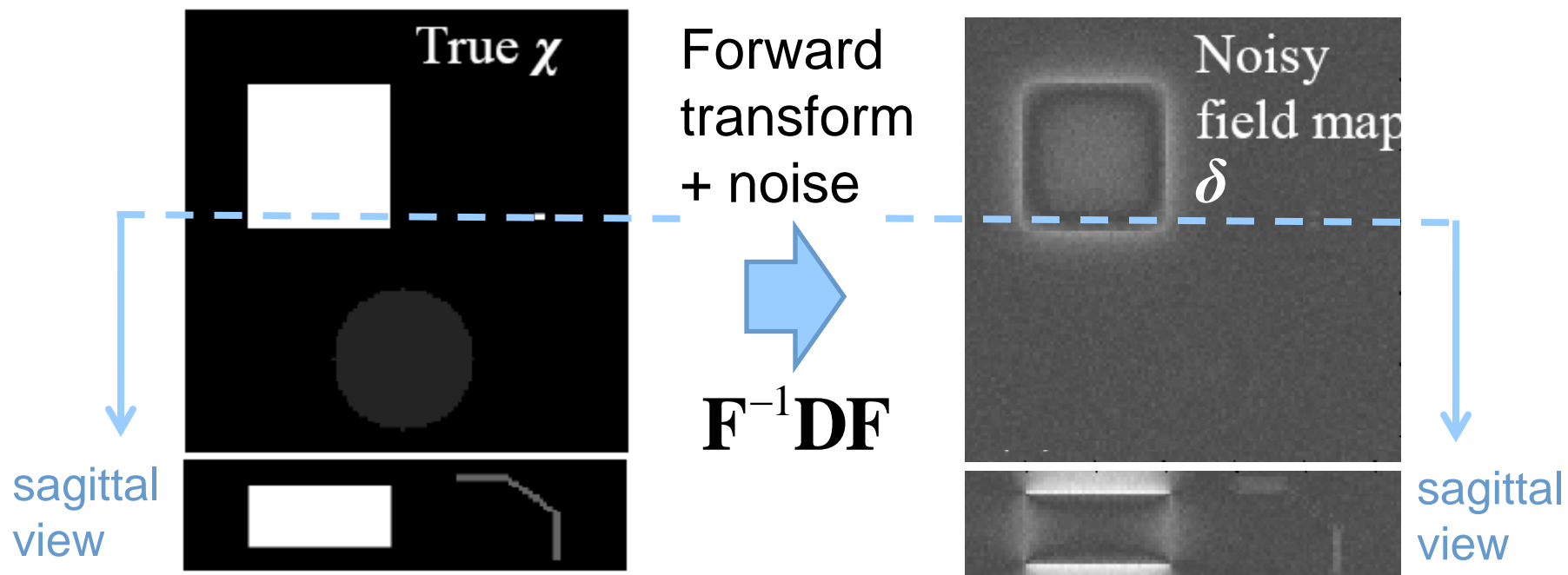


~~$\mathbf{D}^{-1}\mathbf{F}\mathbf{s} = \chi$~~

diverges to ∞

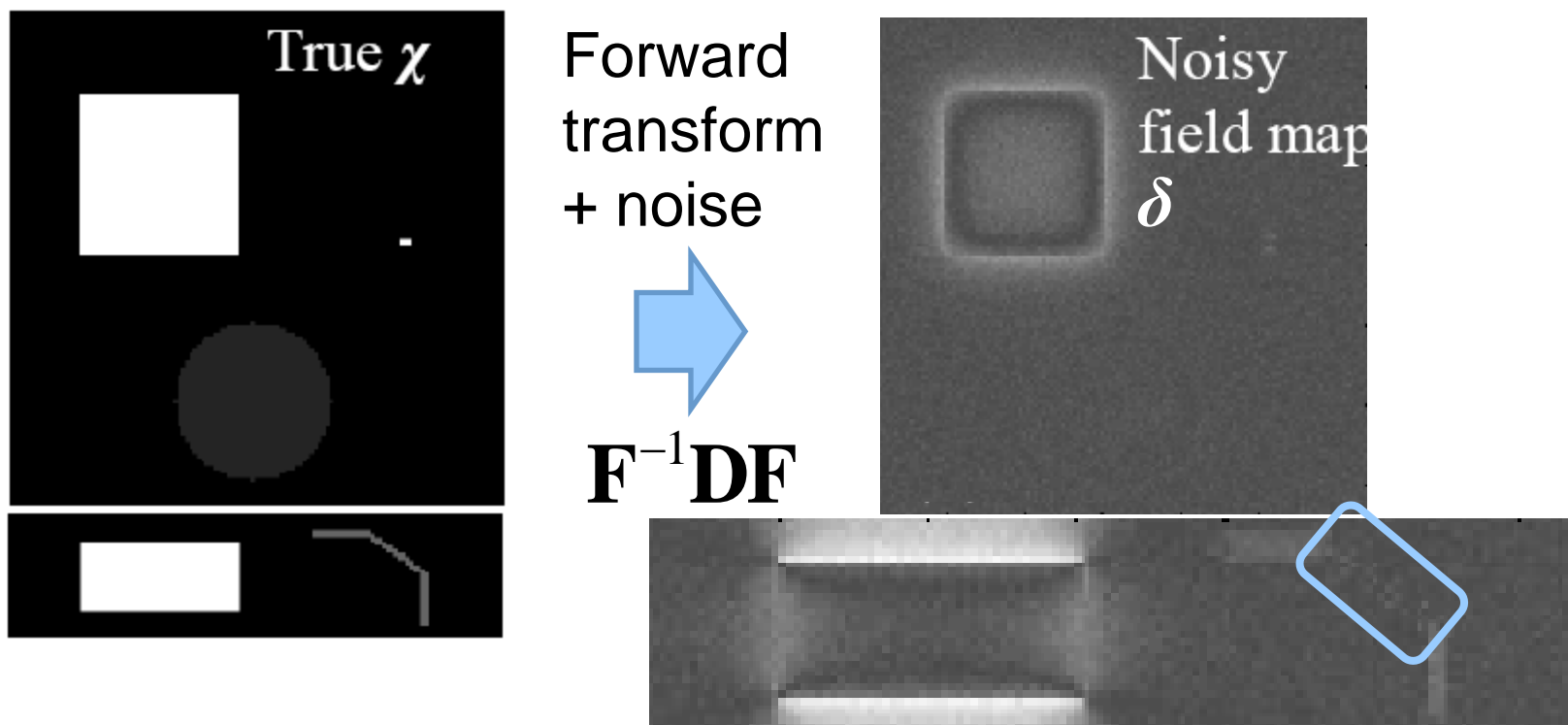
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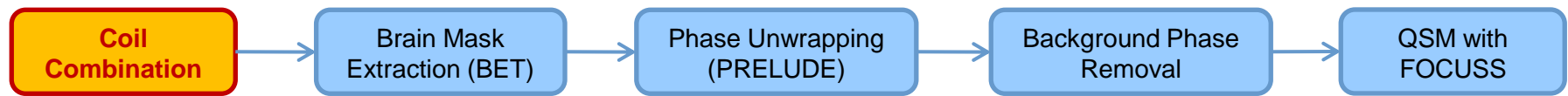


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- Spatial details that have frequency components at the magic angle lose conspicuity in the field map δ

Regularized Inversion for QSM



- Solving for χ by convolving with the inverse of \mathbf{D} is not possible, as it diverges along the magic angle
- Spatial details that have frequency components at the magic angle lose conspicuity in the field map δ
- We propose to use regularization to facilitate the inversion



Phase-aware Coil Combination

- 3D GRE acquisition with phased array coils and body coil
- Normalize each channel image with the body coil

Coil
Combination

Brain Mask
Extraction (BET)

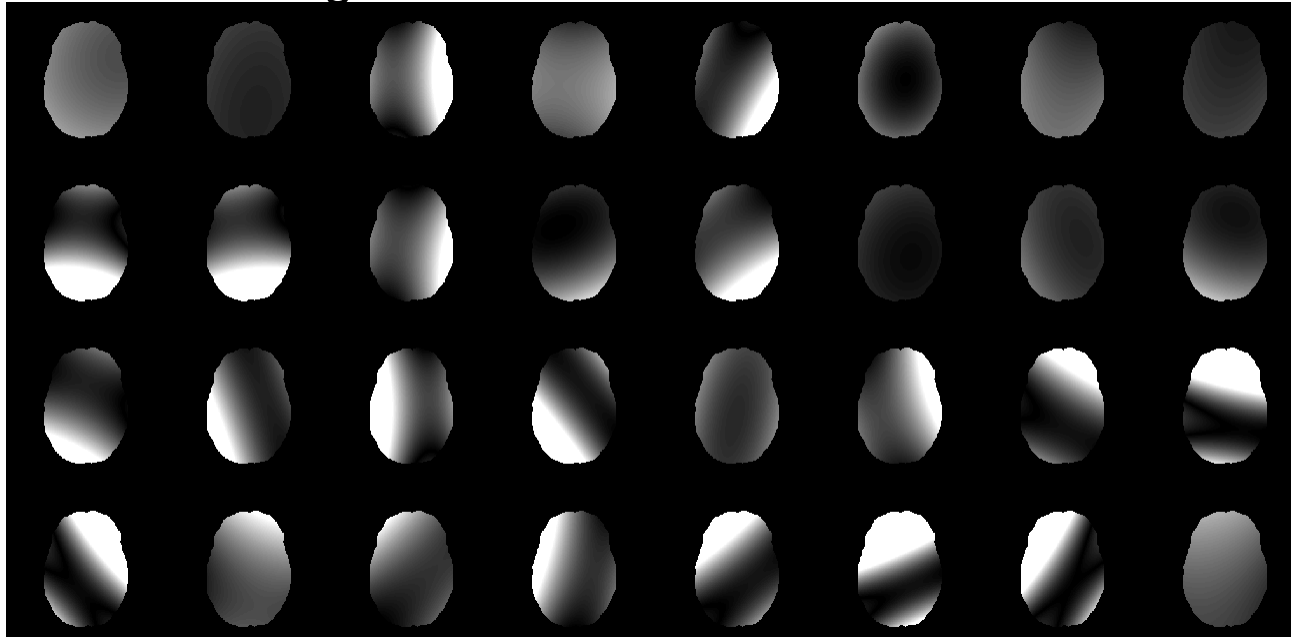
Phase Unwrapping
(PRELUDE)

Background Phase
Removal

QSM with
FOCUSS

Phase-aware Coil Combination

magnitudes of the coil sensitivities



- 3D GRE acquisition with phased array coils and body coil
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- Fit 2nd order polynomials to the magnitude of the normalized images → magnitude of the coil sensitivities

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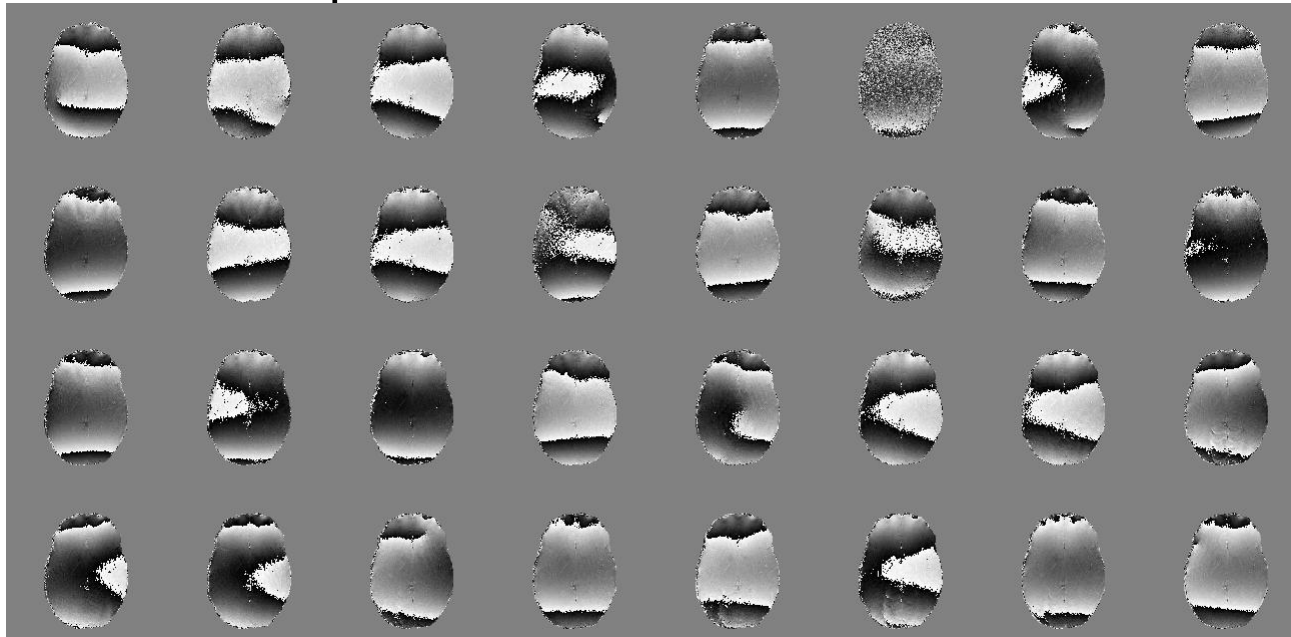
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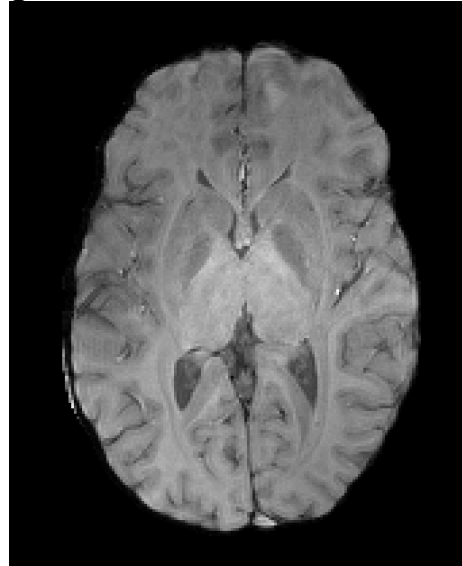


- 3D GRE acquisition with phased array coils and body coil
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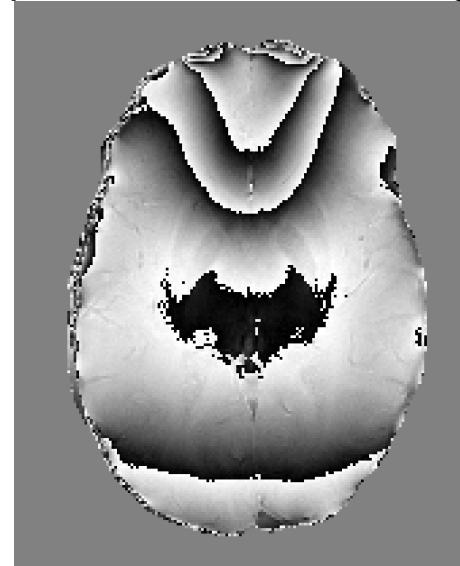


Phase-aware Coil Combination

magnitude of combined image



phase of combined image

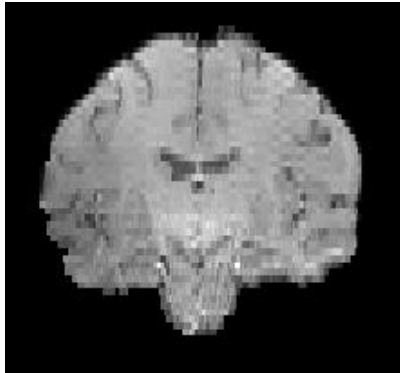


- 3D GRE acquisition with phased array coils and body coil
- Normalize each channel image with the body coil
- Fit 2nd order polynomials to the magnitude of the normalized images → magnitude of the coil sensitivities
- Phase of the normalized images → phase of the coil sensitivities
- Final image is obtained by least-squares coil combination



Brain Mask Extraction & Phase Unwrapping

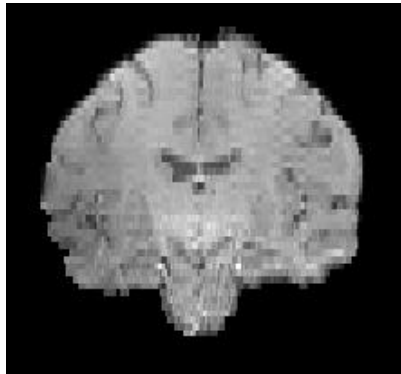
- Brain mask was generated with the FSL Brain Extraction Tool¹



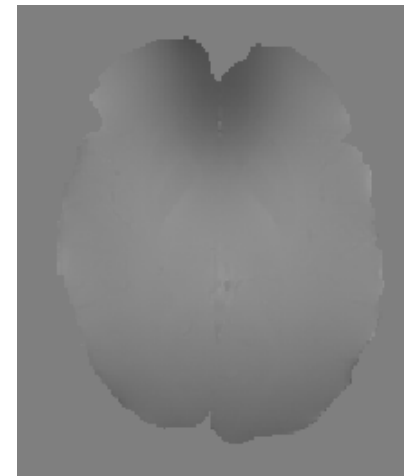
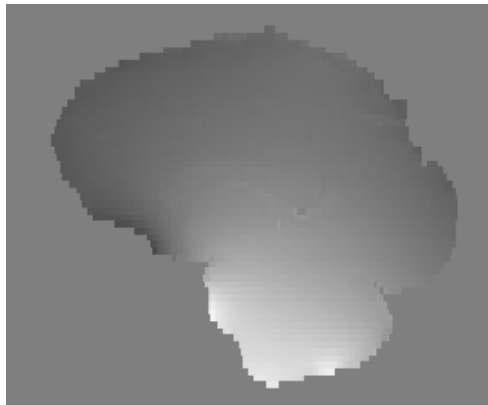
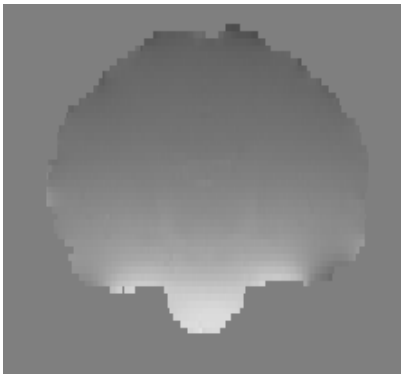


Brain Mask Extraction & Phase Unwrapping

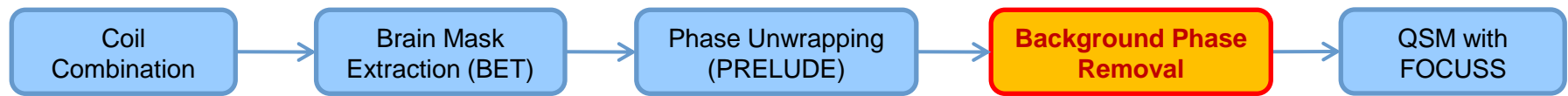
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- Phase unwrapping was done with the FSL PRELUDE²

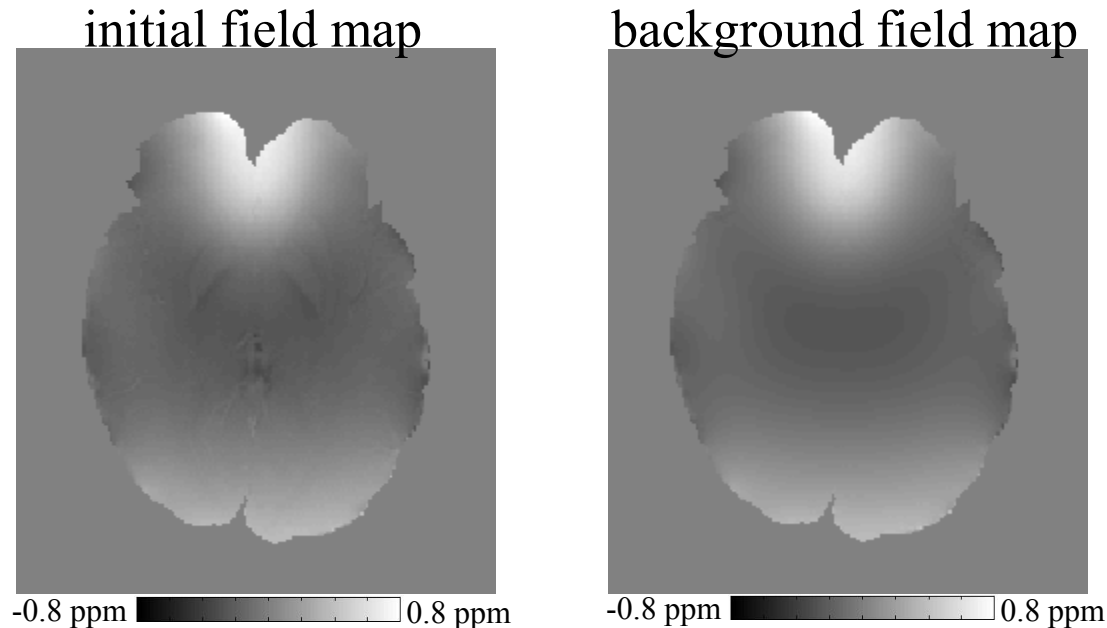


¹ Smith SM, Hum. Brain Mapp. 2002 ² Jenkinson M, MRM 2003



Background Phase Removal

- The background phase was estimated with the Effective Dipole Fitting method¹

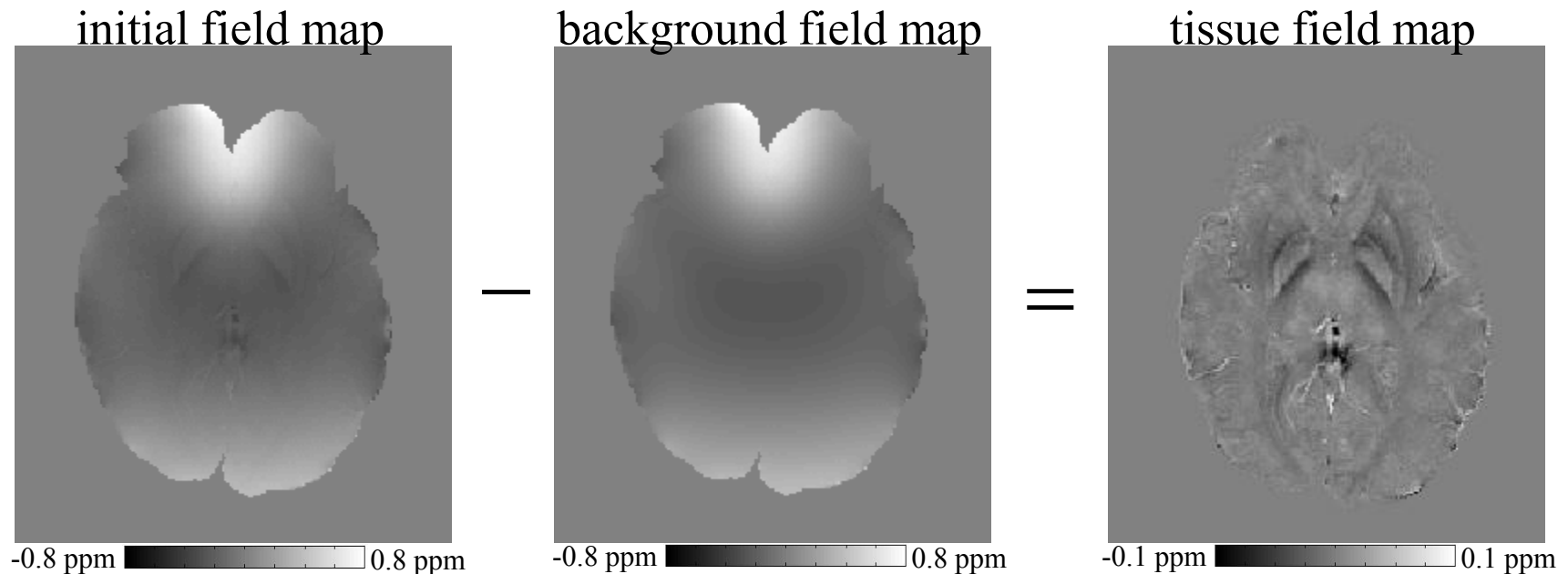


¹ Liu T *et al.*, ISMRM 2010

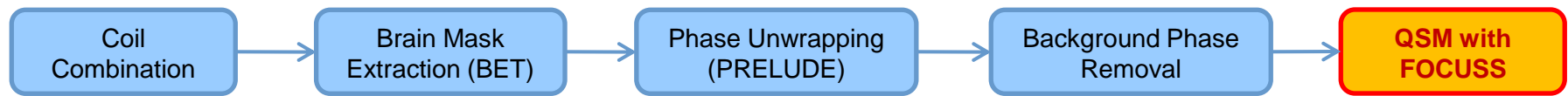


Background Phase Removal

- The background phase was estimated with the Effective Dipole Fitting method¹
- Subtracting the estimated background from the initial field map gives the tissue field map



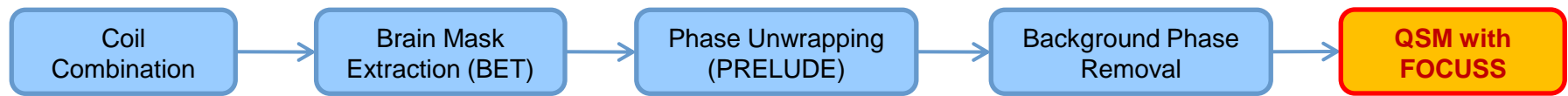
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FOCUSS-QSM with magnitude prior

- The tissue field map δ is related to the susceptibility distribution χ via

$$\delta = \mathbf{F}^{-1} \mathbf{D} \mathbf{F} \chi$$



FOCUSS-QSM with magnitude prior

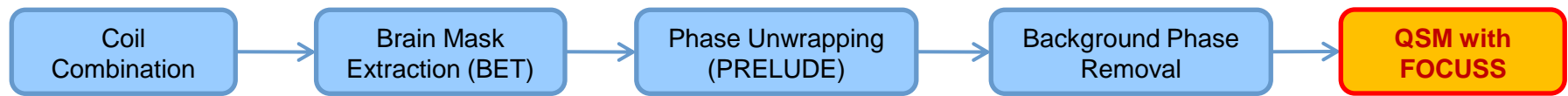
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- Multiplying both sides with $\mathbf{V}_x \mathbf{F}$

$$\mathbf{V}_x \mathbf{F} \delta = \mathbf{V}_x \mathbf{D} \mathbf{F} \chi$$

where \mathbf{V}_x is a diagonal matrix with $\mathbf{V}_x(\omega, \omega) = (1 - e^{-2\pi j \omega / n})$



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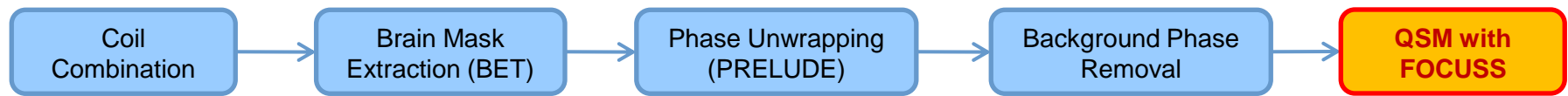
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- This corresponds to taking the spatial gradient along the x axis

$$\mathbf{F}(\partial_x \delta) = \mathbf{D} \mathbf{F}(\partial_x \chi)$$



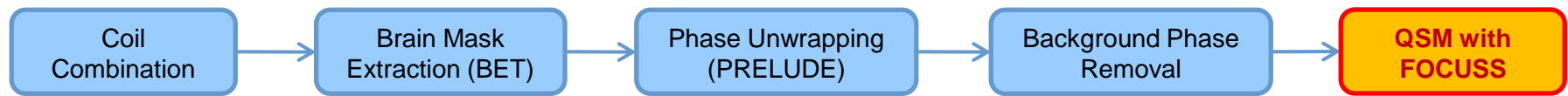
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$$\mathbf{F}(\partial_x \delta) = \mathbf{DF}(\partial_x \chi)$$

- We solve for $\partial_x \chi$ with the FOCUSS algorithm¹ at k^{th} iteration,

$$\mathbf{W}_k = \text{diag}\left(|\partial_x \chi_{k-1}|^{1/2}\right)$$



FOCUSS-QSM with magnitude prior

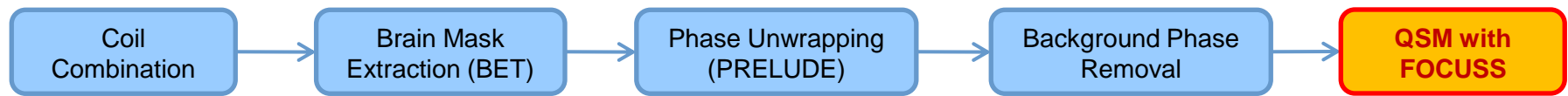
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$$\mathbf{q}_k = \underset{\mathbf{q}}{\text{argmin}} \left\| \mathbf{F}(\partial_x \delta) - \mathbf{DFW}_k \mathbf{q} \right\|_2^2 + \lambda \|\mathbf{q}\|_2^2$$



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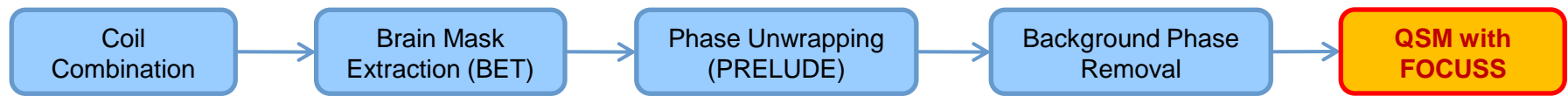
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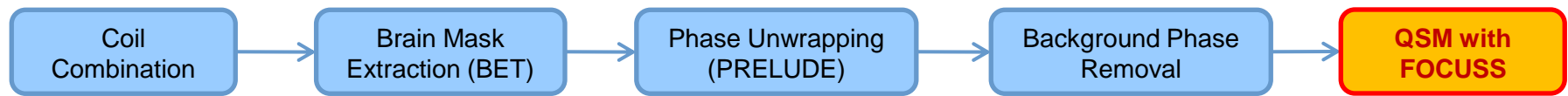
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$$\partial_x \chi_k = \mathbf{W}_k \mathbf{q}_k$$



FOCUSS-QSM with magnitude prior

- We expect the susceptibility distribution to share similar spatial gradients as the magnitude image.



FOCUSS-QSM with magnitude prior

- We expect the susceptibility distribution to share similar spatial gradients as the magnitude image.
- To impose this prior, we modify the update equations as,

$$\mathbf{W}_{prior} = \text{diag}\left(|\partial_x m|^{1/2}\right), \quad m: \text{magnitude image}$$

at k^{th} iteration,

$$\mathbf{W}_k = \text{diag}\left(|\partial_x \chi_{k-1}|^{1/2}\right)$$

$$\mathbf{q}_k = \underset{\mathbf{q}}{\text{argmin}} \left\| \mathbf{F}(\partial_x \delta) - \mathbf{DF} \mathbf{W}_{prior} \mathbf{W}_k \mathbf{q} \right\|_2^2 + \lambda \|\mathbf{q}\|_2^2$$

$$\partial_x \chi_k = \mathbf{W}_{prior} \mathbf{W}_k \mathbf{q}_k$$

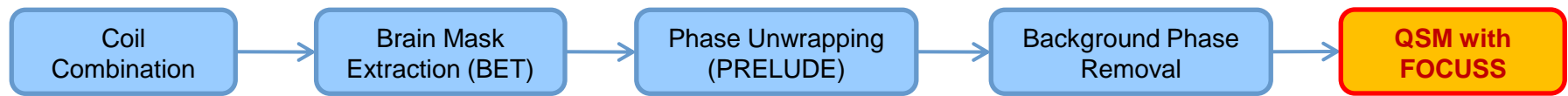


FOCUSS-QSM with magnitude prior

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- Expressed in terms of $\partial_x \chi$,

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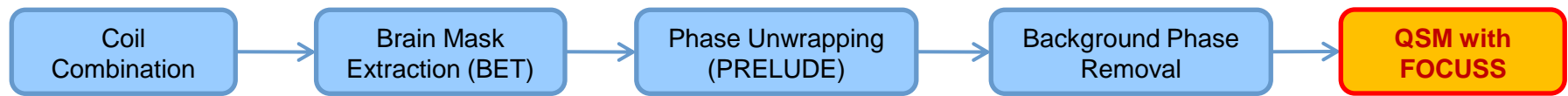
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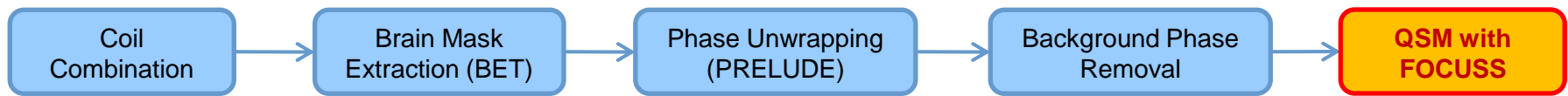
if $\partial_x \mathbf{m}_i$ is small, $\mathbf{W}_{prior}^{-1}(i,i)$ will be large and penalize $\partial_x \chi_i$ more



FOCUSS-QSM with magnitude prior

- After estimating the spatial gradients along x , y and z axes, the susceptibility distribution that matches these is found by solving a least squares problem,

$$\chi = \operatorname{argmin}_{\theta} \sum_{r=x,y,z} \|\partial_r \theta - \partial_r \chi\|_2^2 + \beta \cdot \|\delta - \mathbf{F}^{-1} \mathbf{D} \mathbf{F} \theta\|_2^2$$

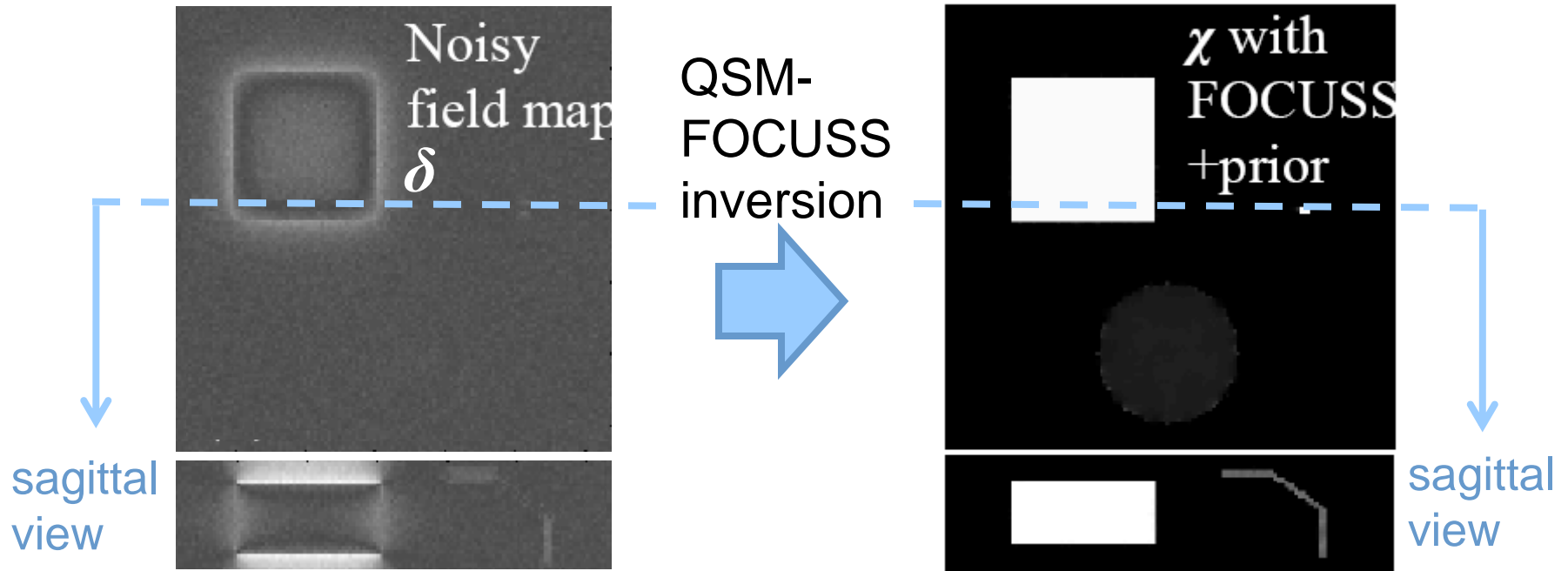


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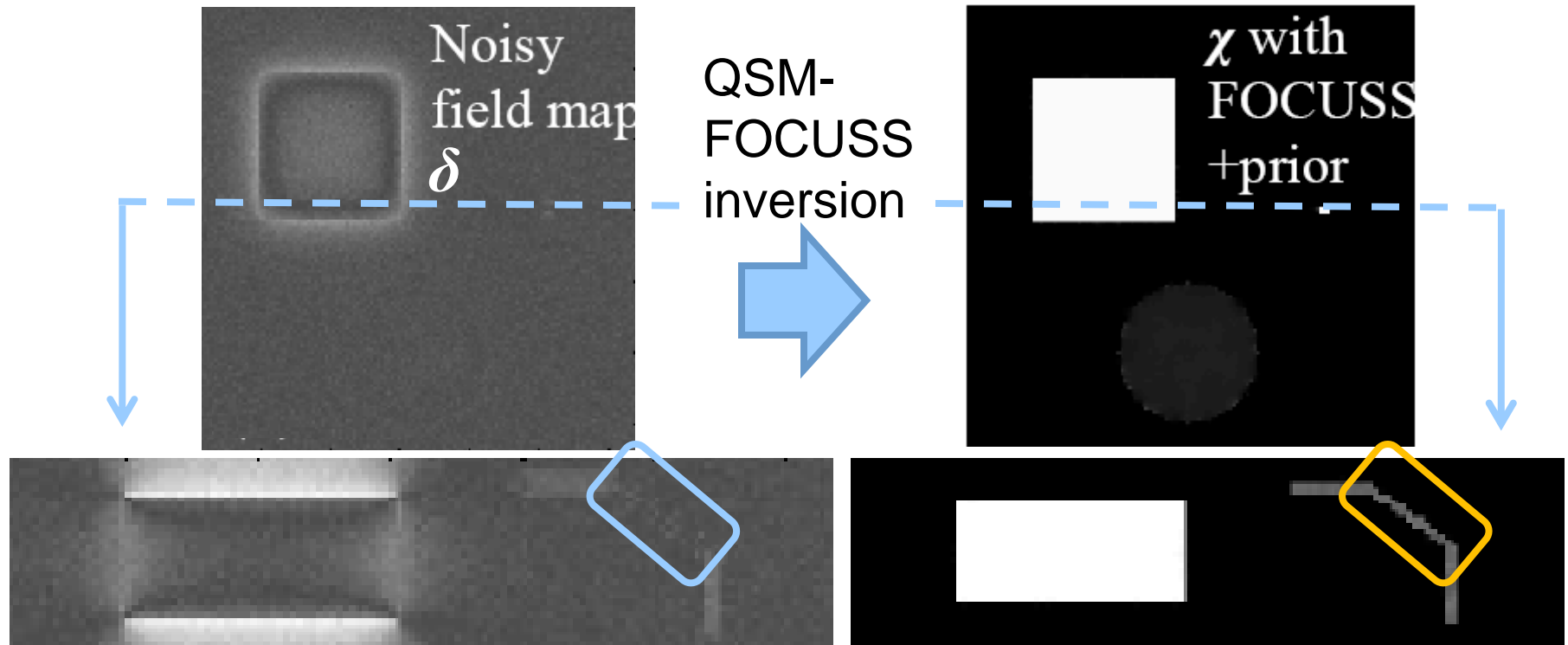
$$\chi = \operatorname{argmin}_{\theta} \underbrace{\sum_{r=x,y,z} \|\partial_r \theta - \partial_r \chi\|_2^2}_{\text{matching gradients}} + \beta \cdot \underbrace{\|\delta - \mathbf{F}^{-1} \mathbf{D} \mathbf{F} \theta\|_2^2}_{\text{data consistency}}$$

QSM result: FOCUSS-QSM with magnitude prior



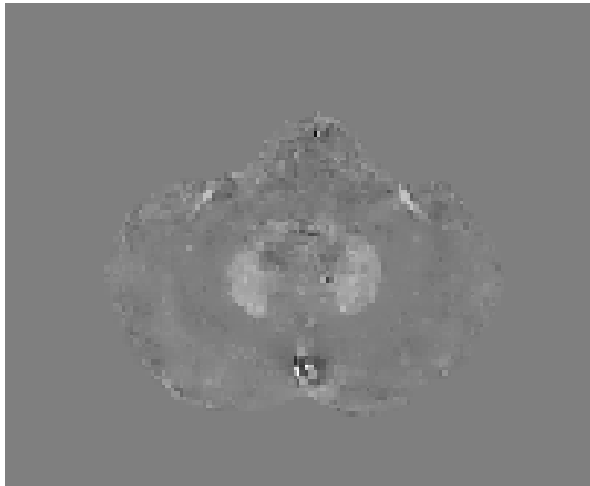
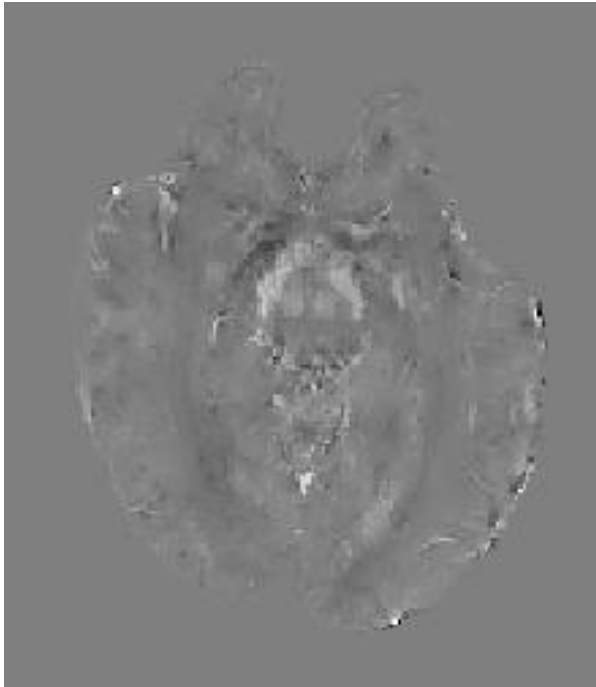
- Starting from the noisy field map δ , FOCUSS-QSM with magnitude prior yielded a susceptibility map with 1.3 % RMSE relative to true χ .

QSM result: FOCUSS-QSM with magnitude prior

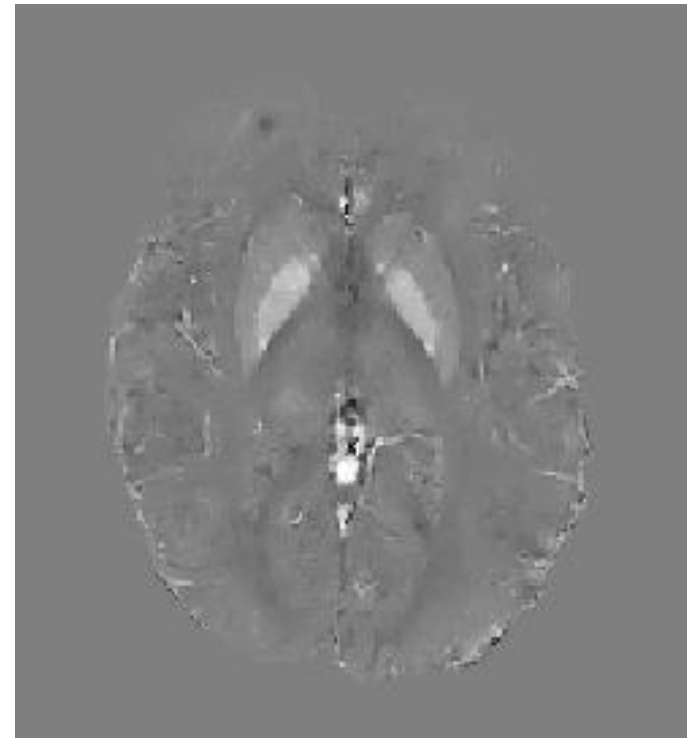


- The reconstructed susceptibility map managed to recover the vessel at the magic angle, which was virtually lost in the field map.

In vivo QSM result: FOCUSS-QSM with magnitude prior

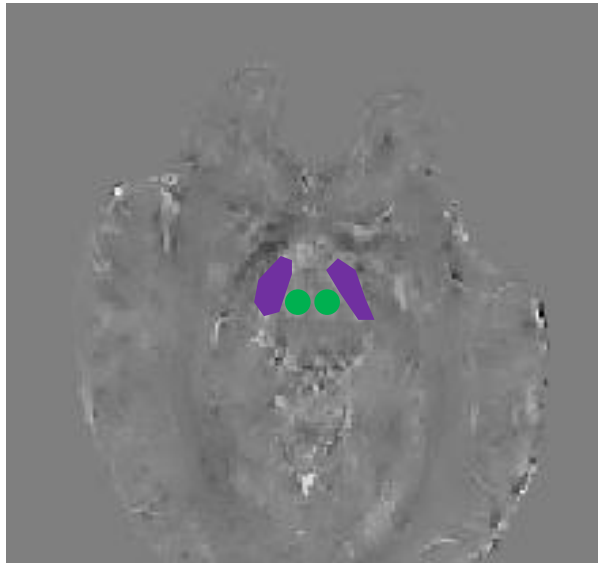


- ❖ 3D GRE acquisition at 3T
- ❖ 32 channel receive array
- ❖ $0.94 \times 0.94 \times 2.5 \text{ mm}^3$ resolution
- ❖ TE: 20 ms



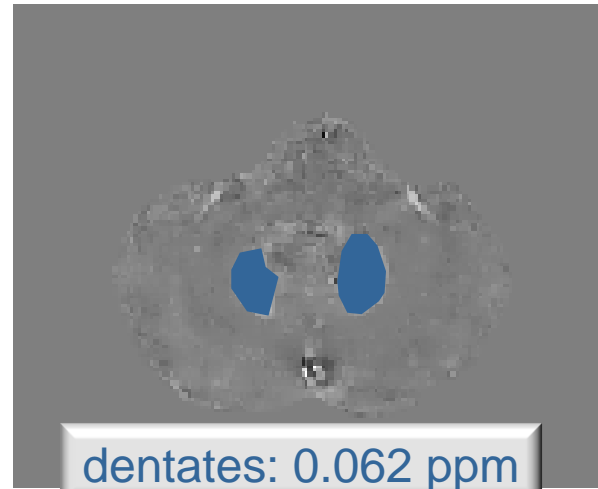
-0.3 ppm  0.3 ppm

In vivo QSM result: FOCUSS-QSM with magnitude prior



subs. nigra: 0.105 ppm

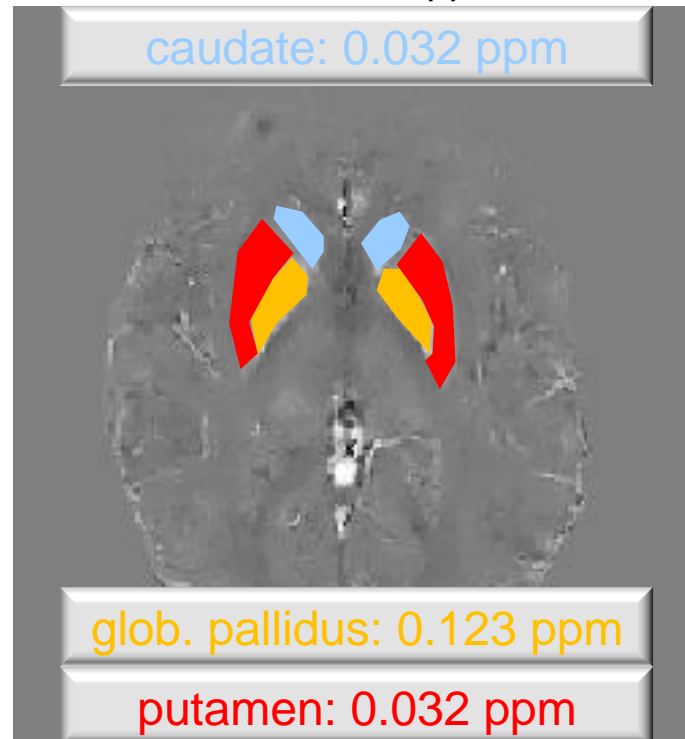
red nuclei: 0.045 ppm



dentates: 0.062 ppm

| Structure | $\Delta\chi$ [ppm] |
|------------------|--------------------|
| Globus Pallidus | 12.3 |
| Substantia Nigra | 10.5 |
| Dentate | 6.2 |
| Red Nucleus | 4.5 |
| Putamen | 3.2 |
| Caudate | 2.3 |

x 0.01 ppm, relative to χ_{CSF}



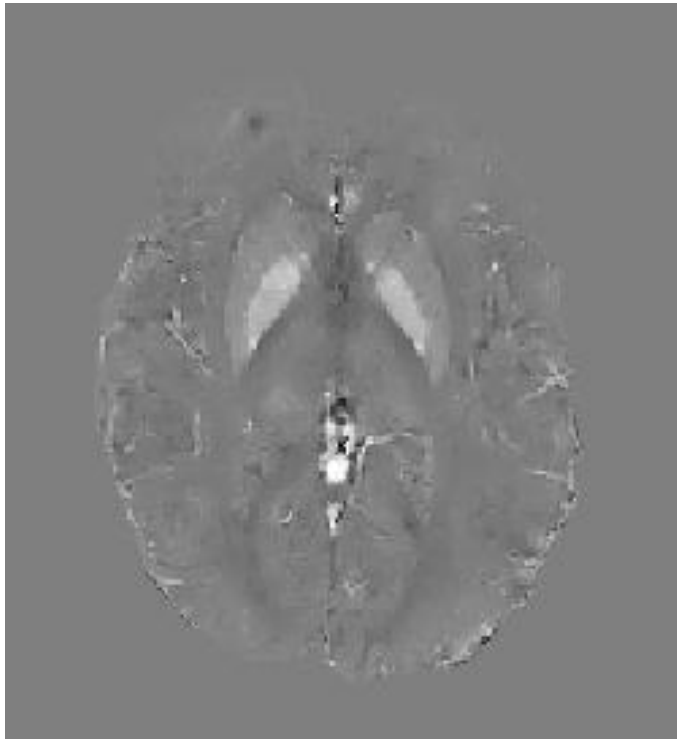
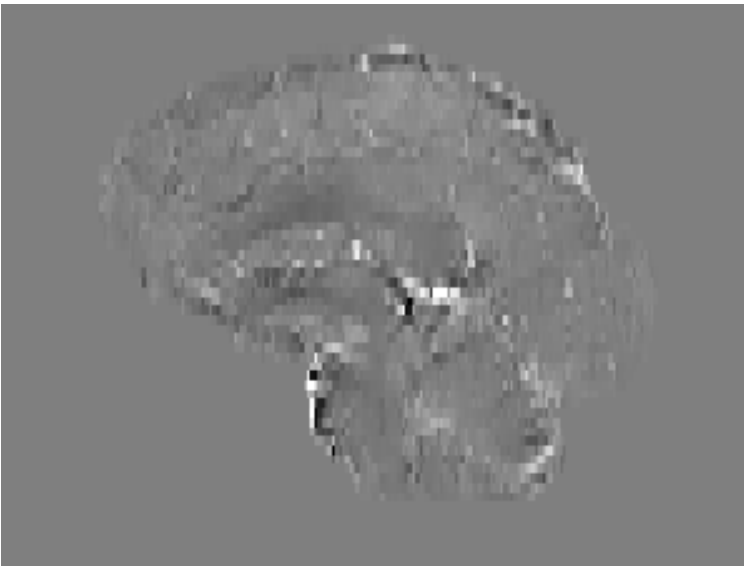
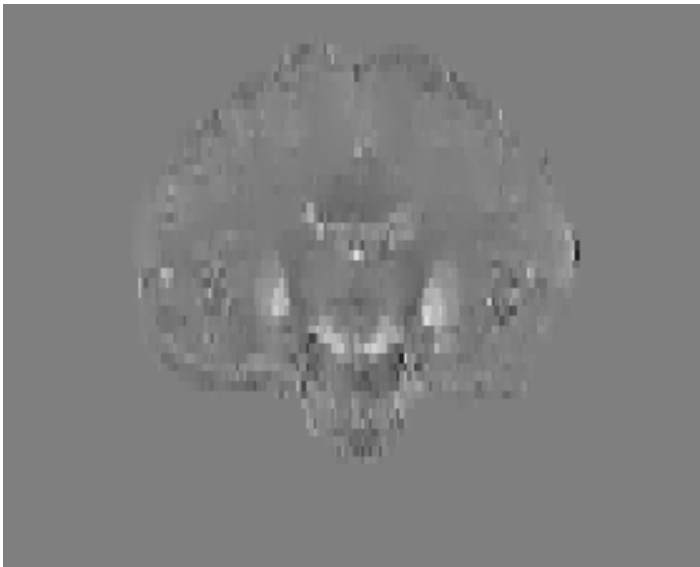
caudate: 0.032 ppm

glob. pallidus: 0.123 ppm

putamen: 0.032 ppm

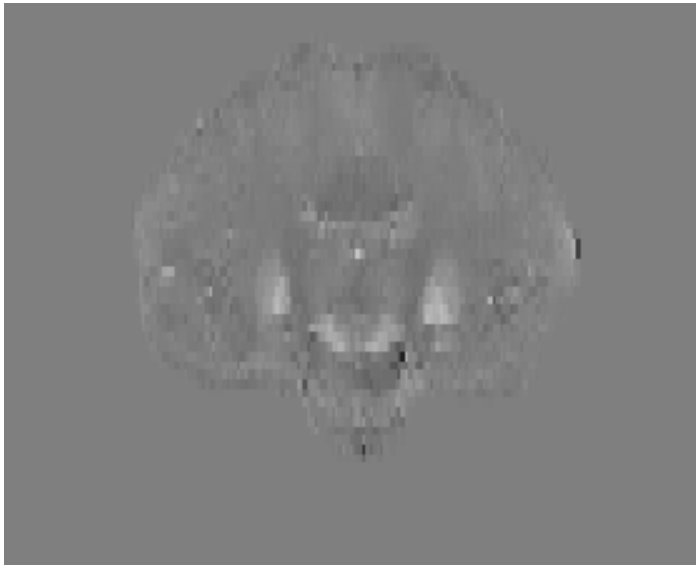


In vivo QSM result: FOCUSS-QSM with magnitude prior

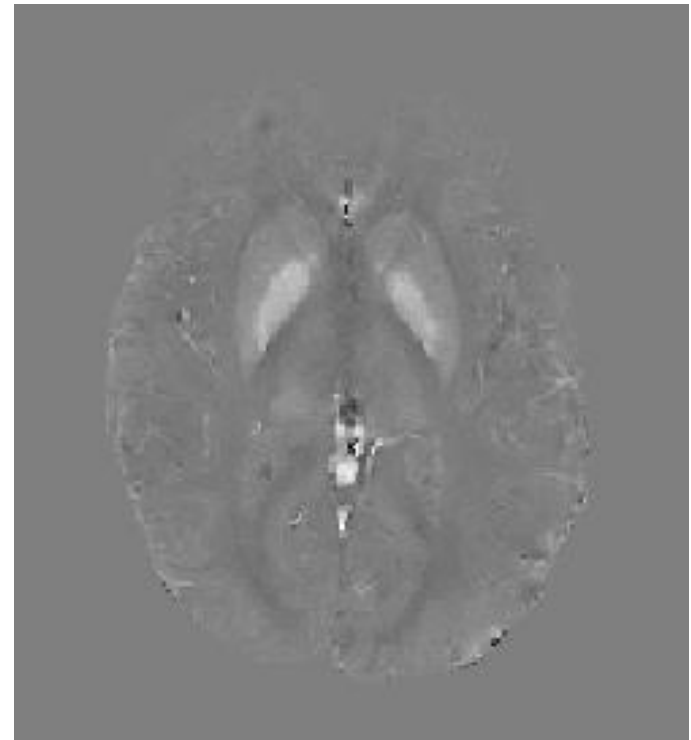
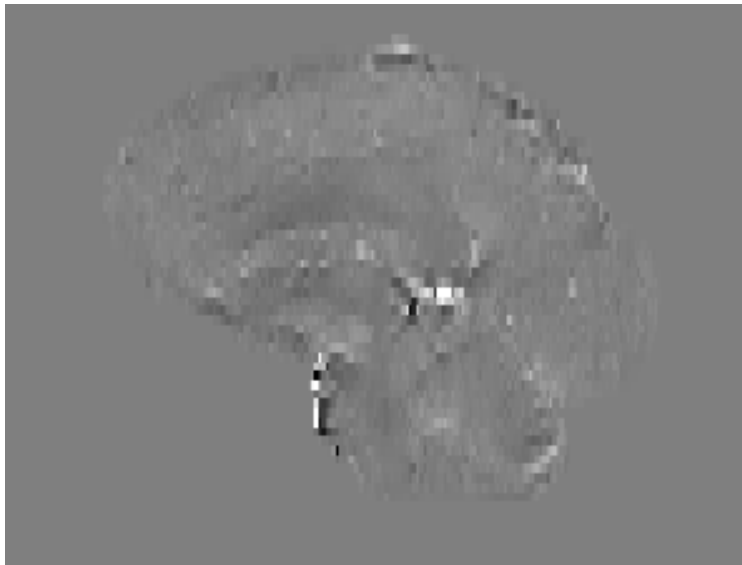


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In vivo QSM result: FOCUSS-QSM with a prior

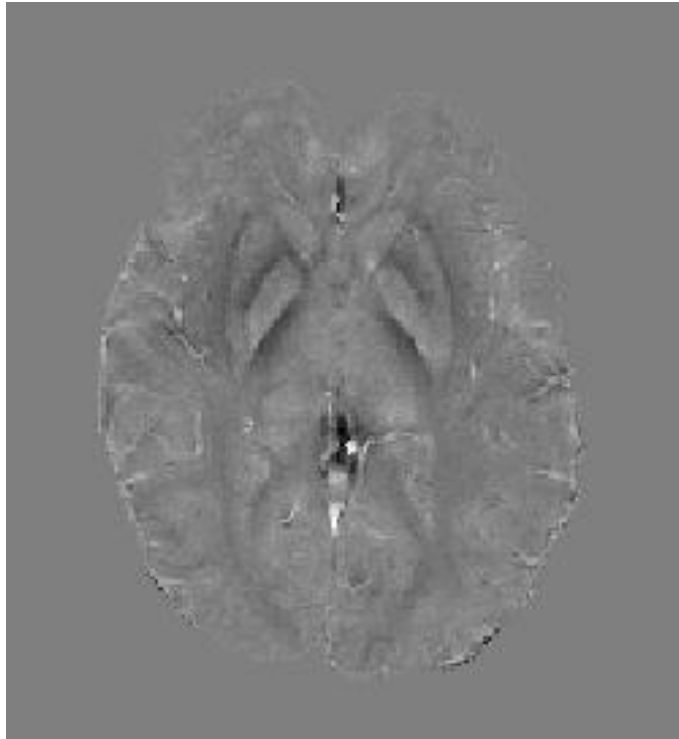
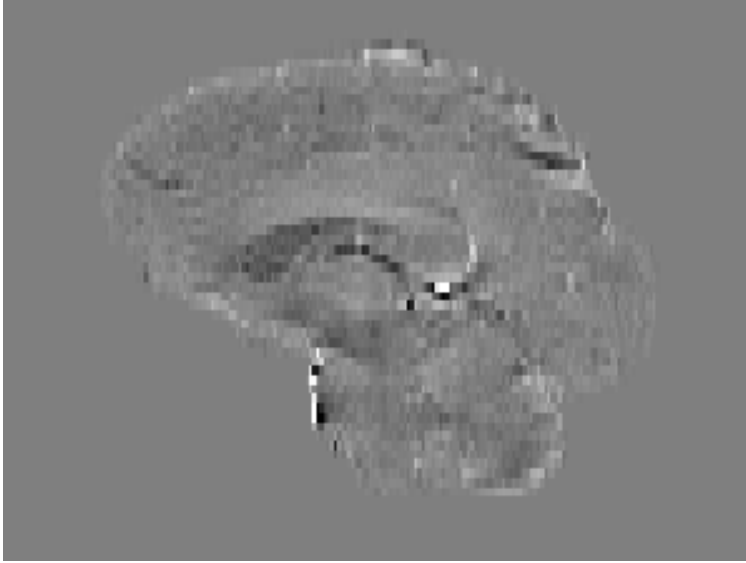
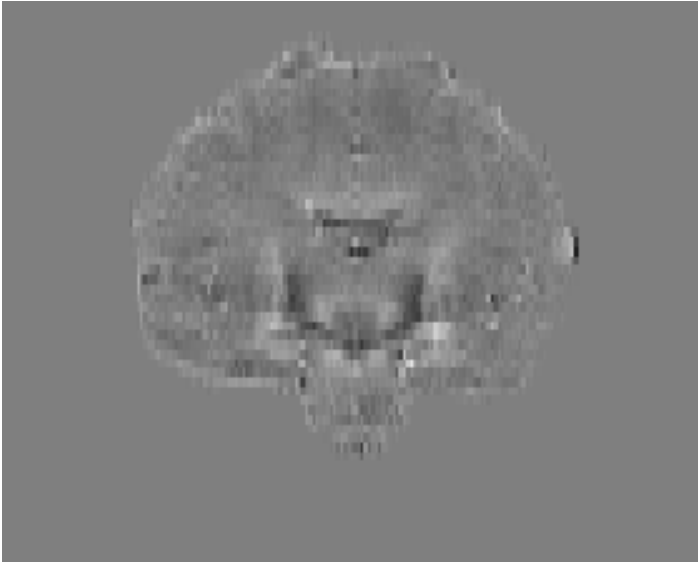


Vessels are less apparent without the magnitude prior

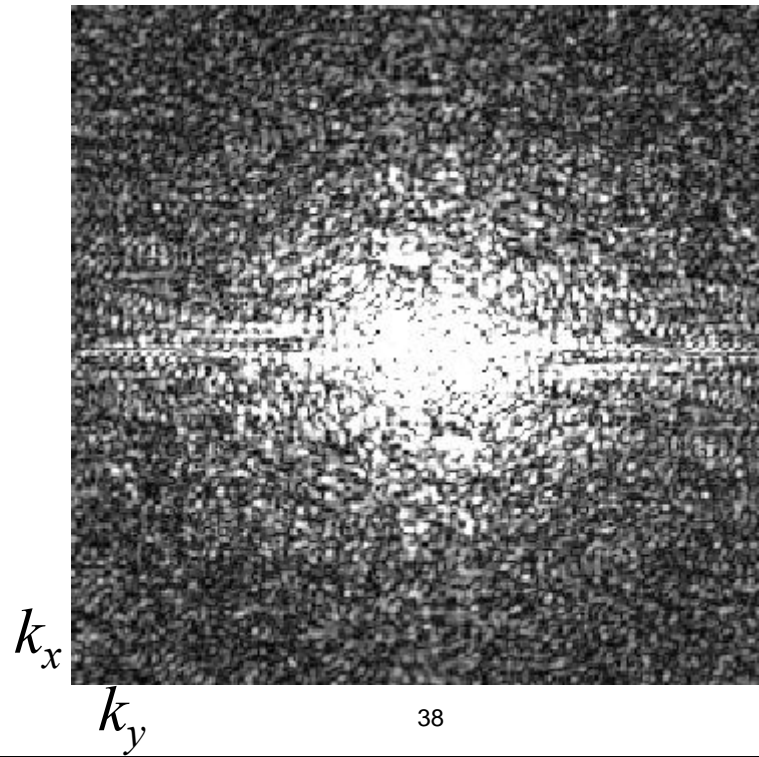
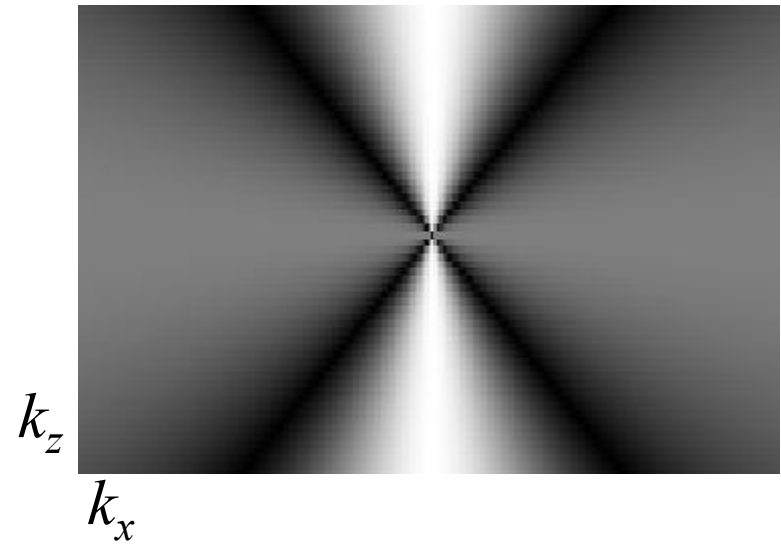
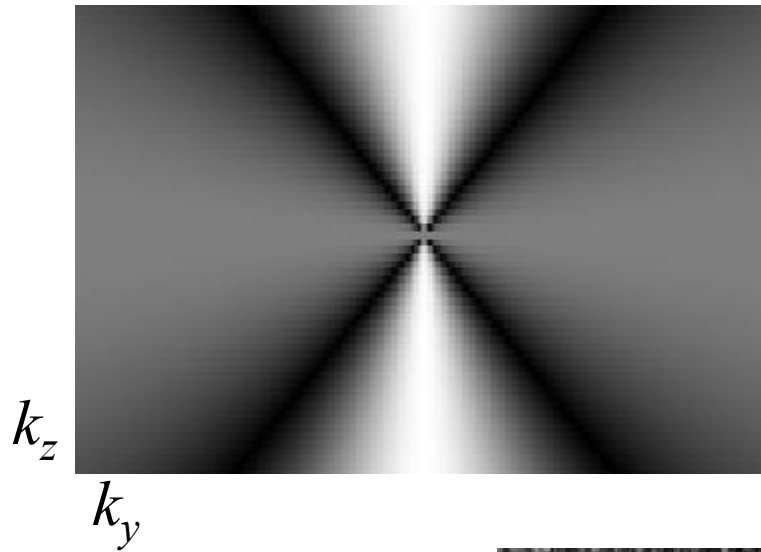


-0.3 ppm  0.3 ppm

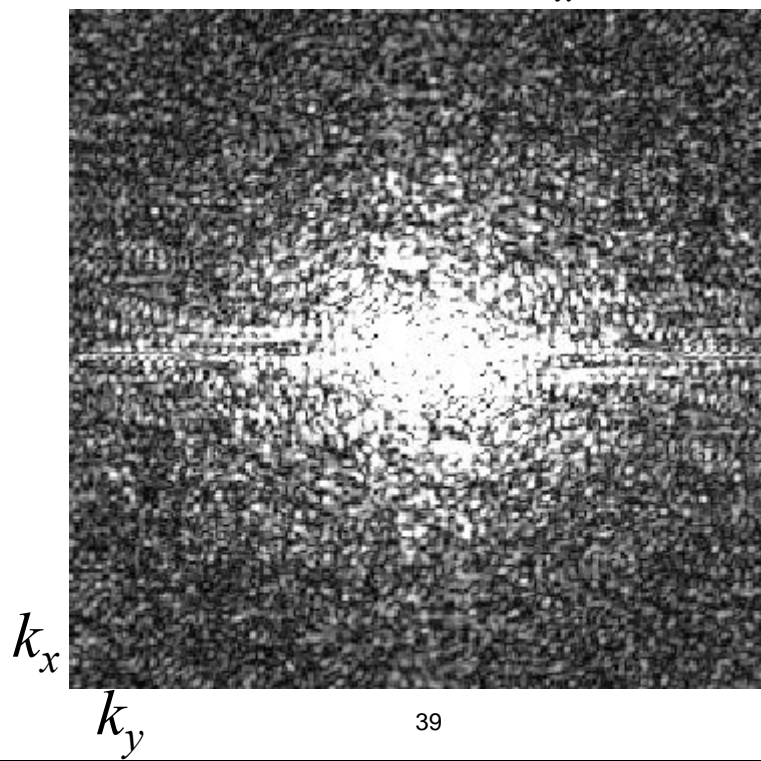
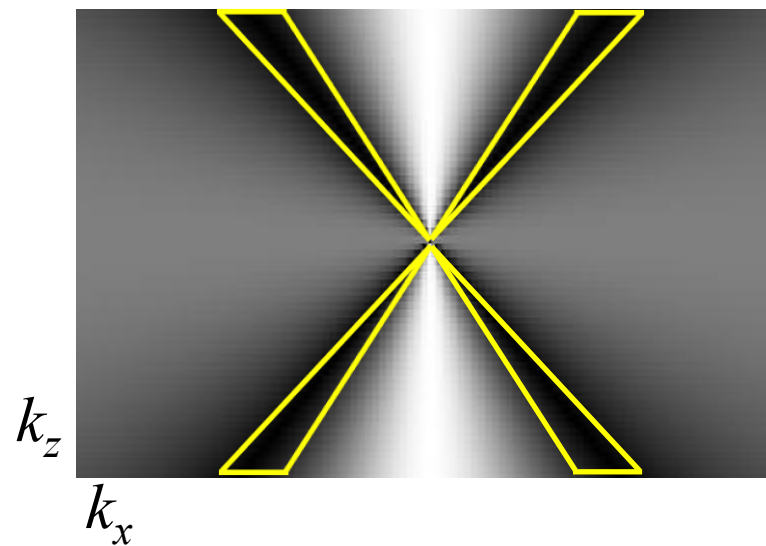
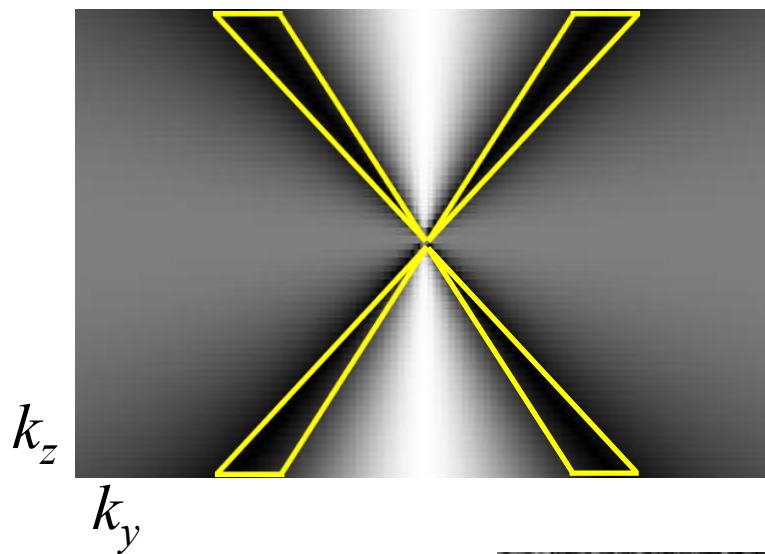
Corresponding Tissue Field Map:



In vivo QSM result with magnitude prior in k-space:



In vivo QSM result with magnitude prior in k-space:



Potential drawbacks of FOCUSS-QSM

- Computation time:
 - ❖ Dipole fitting for background removal \approx 2 hours
 - ❖ FOCUSS-QSM \approx 1 hours
 - ❖ Total processing time \approx **3 hours** for data of size [256x256x64]

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- Solution:

- ❖ Both algorithms solve Least Squares problems, Graphics Processing Card (GPU) implementation will greatly enhance the performance

Conclusion

- Starting with a multi-coil 3D GRE acquisition, we outlined coil combination and background phase elimination methods that yielded the tissue field map.
- We introduced a Quantitative Susceptibility Mapping algorithm that makes use of the magnitude image to facilitate the kernel inversion.