

Simulation-based evaluation of susceptibility distortion correction methods in dMRI

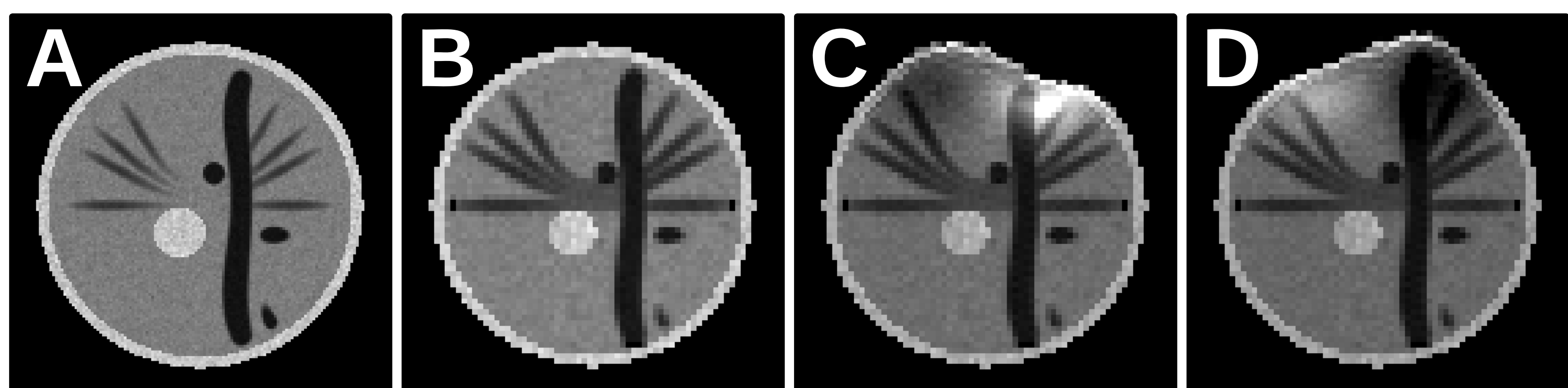
O. Esteban^{1,2} A. Daducci² E. Caruyer³ K. O'Brien⁴ MJ. Ledesma-Carbayo¹ M. Bach-Cuadra^{5,2} A. Santos¹



- Connectivity analyses rely on complex workflows to extract the network from dMRI datasets.
- One important pitfall that potentially bias the extracted connectome is susceptibility distortion, a typical artifact on dMRI [Irfanoglu et al. 2012].
- In this work, we evaluate three widely used methodologies for bias correction, originally proposed for fMRI data: fieldmap-based method (FMB, [Jezzard et al 1995]), reverse-encoding method (REB, [Cordes et al. 2000, Chiou et al. 2000]), and T2-weighted intensity-based registration (T2B, [Kybic et al. 2000]).
- Benchmarking includes geometrical accuracy scores, signal recovery scores, and a preliminary study of impact on the extracted tractography and connectivity matrices.

Digital dMRI phantom & theory-based warping

We generated a test set using low-resolution dMRI phantoms (online available¹) with corresponding T1-weighted and T2-weighted images at high-resolution. We simulate the artifact (geometrical distortion and signal dropout) using a synthetic fieldmap.



A) T2w; B) undistorted b_0 volume; C, D) distorted b_0 volumes with opposed phase encoding directions, maximum displacement of 3.80 mm.

Evaluation framework

We use *nipype*², a powerful tool for building processing pipelines in neuroimaging. The evaluation framework includes the phantom distortion module, the three correction methodologies, DTI&HARDI reconstruction methodologies, tractography, and a final module to analyse downstream impact on geometry, tractography and connectivity.

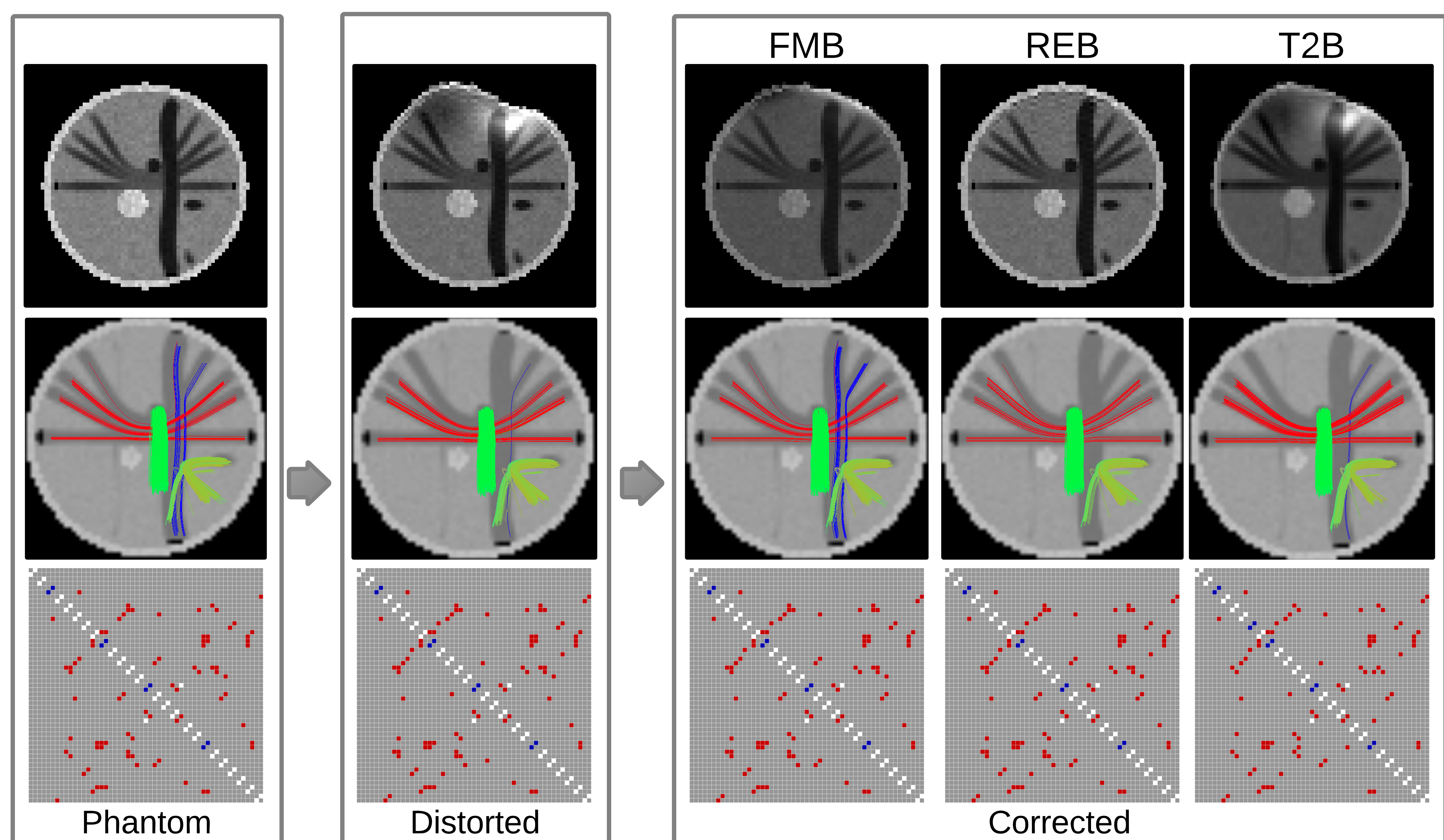
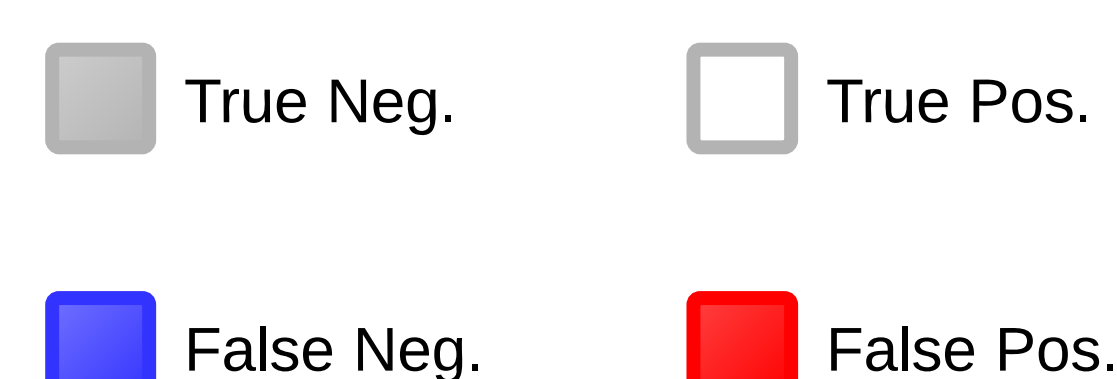
Visual results

Coronal section of b_0

Tractography (Only tracks connecting regions are shown)

Connectivity Matrices

Matrix elements w.r.t. ground-truth:



Quantitative results

Table : Accuracy results

| | Overlap (Jaccard Index, %) | | | | Signal Correlation (%) | |
|-----|----------------------------|-------|-------|-------|------------------------|-------------|
| | Av. | CSF | WM | GM | b_0 | DWIs |
| FMB | 93.00 | 88.57 | 96.74 | 94.02 | 80.05 | 96.26 ± .06 |
| REB | 96.64 | 94.31 | 98.26 | 96.75 | 91.00 | 97.65 ± .03 |
| T2B | 79.19 | 66.31 | 89.85 | 82.14 | 64.58 | 90.10 ± .13 |

Table : Tractography and connectivity results.

| | # tracks | length (mm) | FP | FN |
|-----------|----------|---------------|----|----|
| Original | 735 | 40.87 ± 13.55 | 40 | 4 |
| Distorted | 878 | 40.54 ± 13.73 | 42 | 4 |
| FMB | 743 | 40.04 ± 13.60 | 43 | 4 |
| REB | 830 | 39.87 ± 13.93 | 44 | 4 |
| T2B | 825 | 41.44 ± 12.85 | 40 | 5 |

Conclusions and references

- DTI dataset was rejected in evaluation: extracted connectivity matrices were biased, as the phantom is designed for high angular resolution methods.
- In terms of geometry, the results indicate that REB method ranks first.
- In terms of tractography, visual assessment and quantitative results suggest that FMB could be better.
- Connectivity matrices are evaluated, but a more appropriate phantom is required. A phantom with the connecting interface densely covered by the seeding regions may be the key to characterize the impact on connectivity.
- Connectome analyses demand the standardization of processing techniques and pipelining software tools to ensure replicability of experiments and reliability of results.

Links and references

1. emmanuelcaruyer.com/phantomas.php
2. nipy.sourceforge.net/nipype