

# Simulation-based evaluation of susceptibility distortion correction methods in dMRI

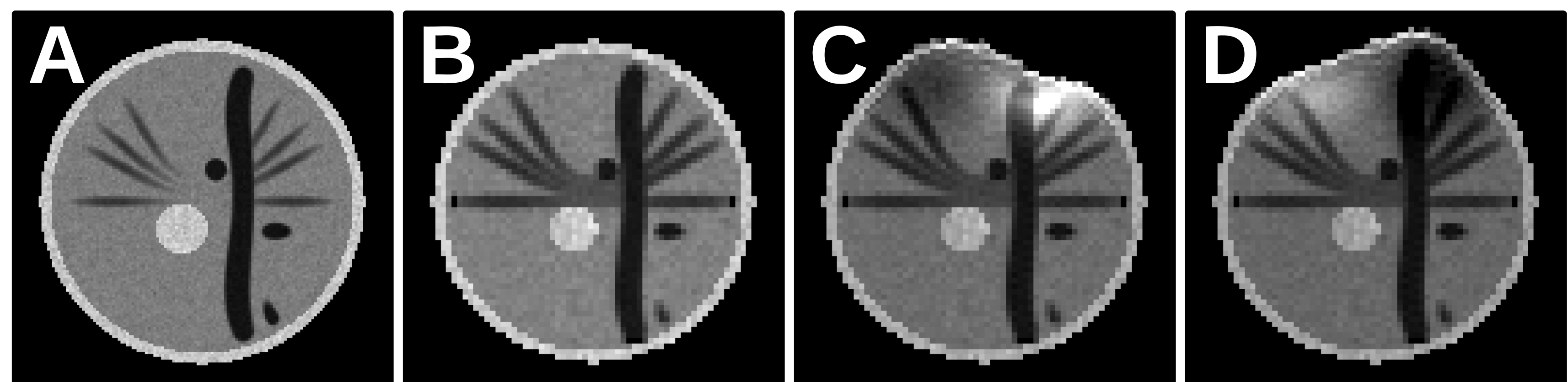
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- Connectivity analyses rely on complex workflows to extract the network from dMRI datasets.
- One important pitfall that potentially biases the extracted connectome is susceptibility distortion, a typical artifact on dMRI [Irfanoglu et al. 2012].
- 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 available<sup>1</sup>) with corresponding T1-weighted and T2-weighted images at high-resolution. We simulate the artifact (geometrical distortion and signal dropout) using a synthetic fieldmap.

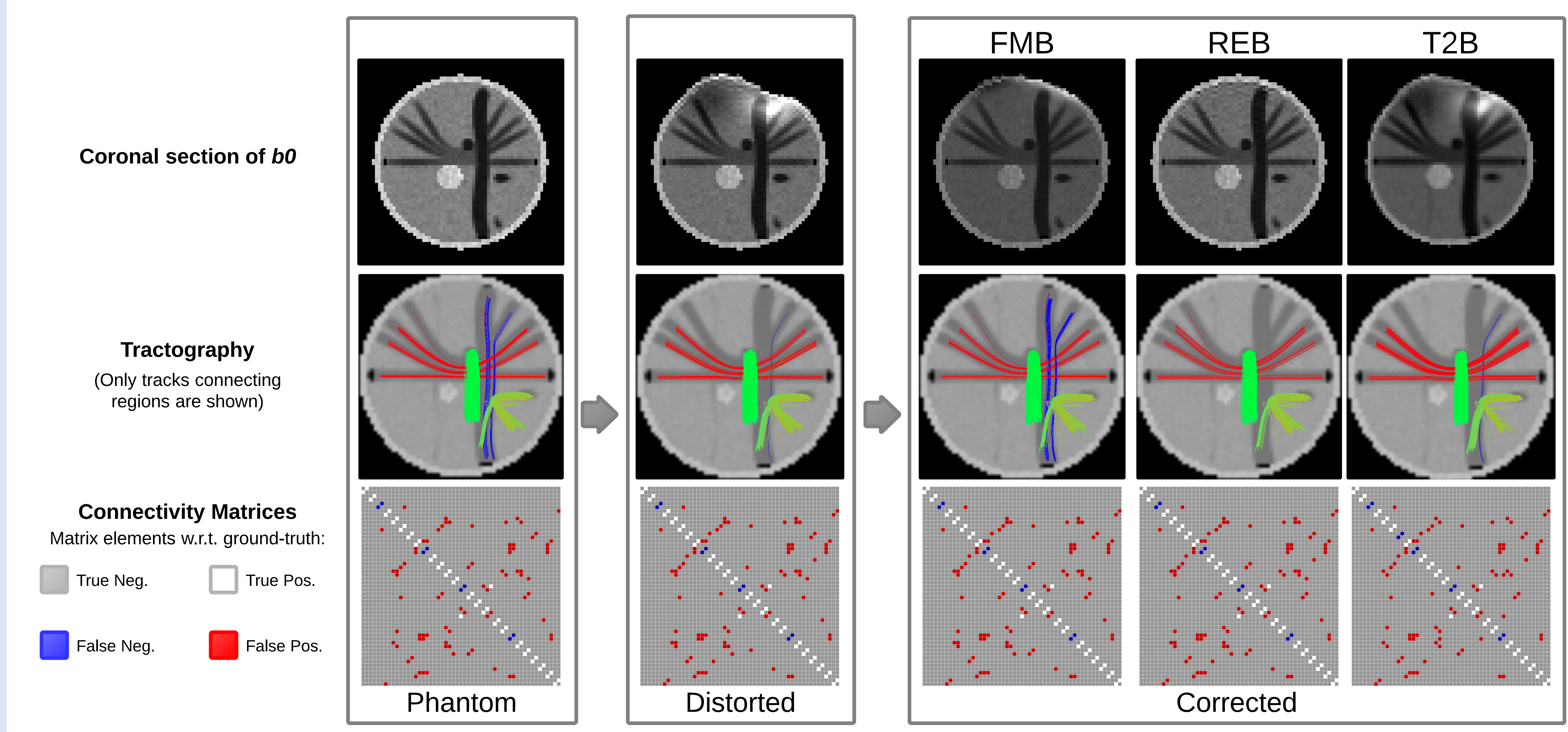


A) T2w; B) undistorted  $b_0$  volume; C, D) distorted  $b_0$  volumes with opposed phase encoding directions, maximum displacement of 3.80 mm.

## Evaluation framework

We use *nipype*<sup>2</sup>, 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



## 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](http://emmanuelcaruyer.com/phantomas.php)
2. [nipy.sourceforge.net/nipype](http://nipy.sourceforge.net/nipype)

