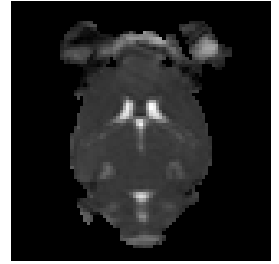


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DRBUDDI for MRI Processing

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Protocol status: In development

We are still developing and optimizing this protocol

Created: October 01, 2019

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Abstract

This protocol will provide a basic guide to utilizing the DRBUDDI tool in TORTIOSE.

Note: Steps may vary based upon image.





Introduction

- 1 DRBUDDI is a EPI distortion correction module in TORTOISE. It bases its corrections by using DTI data pairs (blip up and blip down). When processing MR images DRBUDDI is usually the second step in the pipeline.

DRBUDDI

- 2 Type the following in the terminal to view information

Command

```
DR_BUDDI_withoutGUI
```

- 2.1 You should have two corresponding data sets to edit, and you will need to distinguish them when you call DRBUDDI i.e. blip up and blip down
- 2.2 NOTE: There are two DRBUDDI versions available to use. There is a DRBUDDI with a GUI, and there is one without a GUI.
- 3 In terminal go to your directory with the DTI data.
- 4 In terminal type:



Command

```
DR_BUDDI _withoutGUI --up _data directory(blip  
up)/file_name_proc.list --down_data directory(blip  
down)/file_name_proc.list --structural  
directory_T2Data/file_name_structural.nii
```