

# PTB Tools Documentation

Version 0.0.1

PTB-Division 8.14

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### 1 Description

PTB Tools is software designed to run various  $B_1^+$  Mapping and Shimming algorithms in the context of Magnetic Resonance Imaging (MRI) at ultrahigh magnetic fields, used by division 8.14 of the Physikalisch-Technische Bundesanstalt (PTB).

The software's frontend is designed using HTML, CSS, and Vanilla JavaScript. The backend is programmed in Python, using the Flask library. The graphical interface of the software is expected to be run in the Google Chrome browser.

### 2 Installation Instructions

Follow the next steps to install everything needed to run this software.

1. Clone the Github repo:

```
git clone https://github.com/Spoksonat/PTB-Tools.git
```

2. Navigate to the project directory:

```
1 cd PTB-Tools-main
```

3. Install dependencies:

```
pip3 install -r requirements.txt
```

### 3 Execution

To access the graphical interface, you must follow the next steps:

Change the access permissions to the main bash script:

```
chmod u+x execute_app.sh
```

2. Execute the main bash file

```
1 ./execute_app.sh
```

Note: If the software stops running and you want to access the graphical interface again, just execute the second step.

### 4 General Structure and Use

After executing the command:

./execute\_app.sh

the main window of PTB Tools opens, as shown in Figure 1.

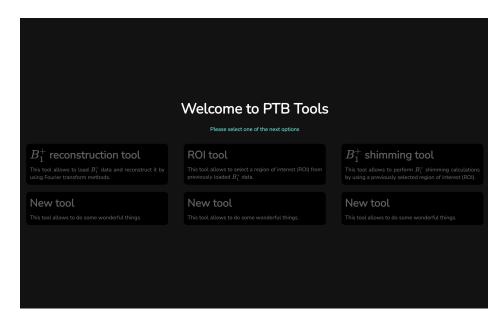


Figure 1: Main Menu of PTB-Tools.

The main menu features six tools, and you can access each of them by clicking the corresponding button. Let's take a closer look at each of the six tools.

### 4.1 $B_1^+$ Mapping Tool

Once you click the " $B_1^+$  reconstruction tool" button, the menu for the  $B_1^+$  Mapping Tool will appear, as shown in Figure 2.

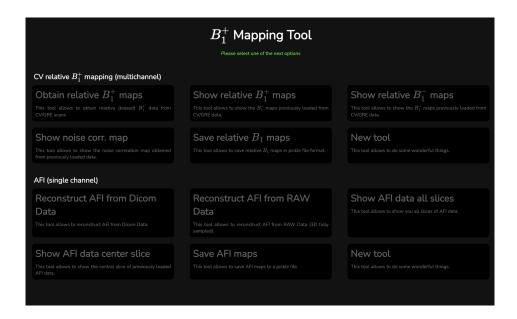


Figure 2:  $B_1^+$  Mapping Tool.

The  $B_1^+$  Mapping Tool consists of two main features:

- CV relative  $B_1^+$  mapping (multichannel): This tool allows you to load and reconstruct CV/GRE scans, as well as obtain and visualize the corresponding maps of  $B_1^+$ ,  $B_1^-$ , and Noise correlation.
- AFI (single channel): Note that this tool has not been implemented yet.

The instructions for using each of these features will be described in detail below.

#### 4.1.1 CV relative B<sub>1</sub> mapping (multichannel)

First, you must load the CV/GRE scan data by clicking the corresponding button. After clicking the button, a window will appear, as shown in Figure 3.

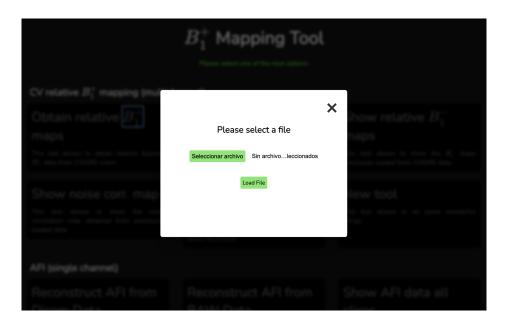


Figure 3: CV/GRE load window.

If you click the "Select File" button, the file explorer will open. You must select the file with the .dat extension that you want to open<sup>1</sup>. Once you select the desired file, you should press the "Load File" button to load the file.

Once the file has been loaded and the respective  $B_1^+$ ,  $B_1^-$ , and Noise correlation maps have been obtained, these maps can be viewed by clicking the "Show relative  $B_1^+$  maps", "Show relative  $B_1^-$  maps", and "Show noise corr.map" buttons respectively. Figure 4 shows the window where the  $B_1^+$  relative maps are displayed. To return to the previous menu, you can press the "Go back" button. The same instructions apply to the windows where the  $B_1^-$  relative and Noise correlation maps are displayed. The maps were generated using the Python library called Bokeh.

<sup>&</sup>lt;sup>1</sup>The file must be located at PTB-Tools-main/main/files/RAW

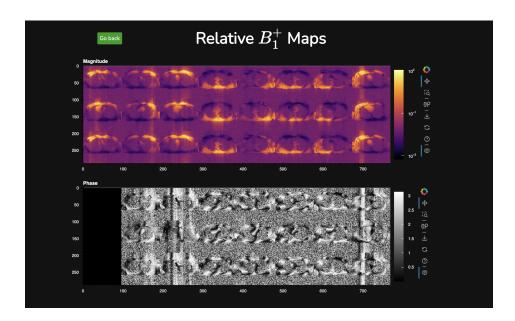


Figure 4: Example - Relative  $B_1^+$  maps.

Note: If the "Show relative  $B_1^+$  maps", "Show relative  $B_1^-$  maps", or "Show noise corr.map" buttons are pressed before loading the CV/GRE data, the alert shown in Figure 5 will appear.

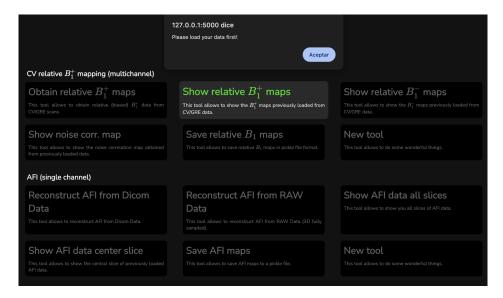


Figure 5: CV/GRE loading alert.

Note: If any "New tool" button, related to additional tools that might be implemented in the future, is pressed, the alert shown in Figure 6 will appear.

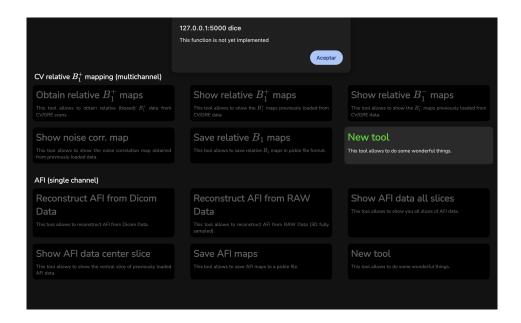


Figure 6: New tool alert.

If you want to save your  $B_1^+$ ,  $B_1^-$ , and Noise correlation maps, you must press the "Save relative  $B_1$  maps" button. When you press this button, the window shown in Figure 7 will appear. In the gray rectangular space, you can write the name under which you want to save your  $B_1^+$ ,  $B_1^-$ , and Noise correlation maps. Finally, you should press the "Save File" button. The file will be saved as a Python-pickle file with a *.pkl* extension in the PTB-Tools-main/main/files/B1p\_maps directory.

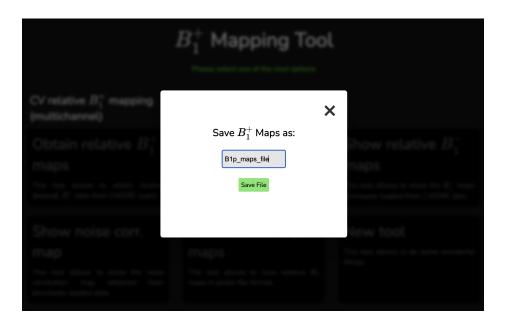


Figure 7: Save  $B_1$ -data tool.

#### 4.2 ROI Tool

To be developed

# **4.3** $B_1^+$ Shimming Tool

To be developed

### **5** Contact Information

For any questions or suggestions, please contact:

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