# MRVelocityMapping V1.3.1 Manual

# 1. Open Tools

Double Click 'MRVelocityMappingV1.3.1.exe'. When you fail to open the tools panel like as Fig.1 then install MS Visual C++ Runtime libs first. Run 'vcredist\_x86.exe' or 'vcredist\_x64.exe' which can be downloaded from Microsoft official websites.

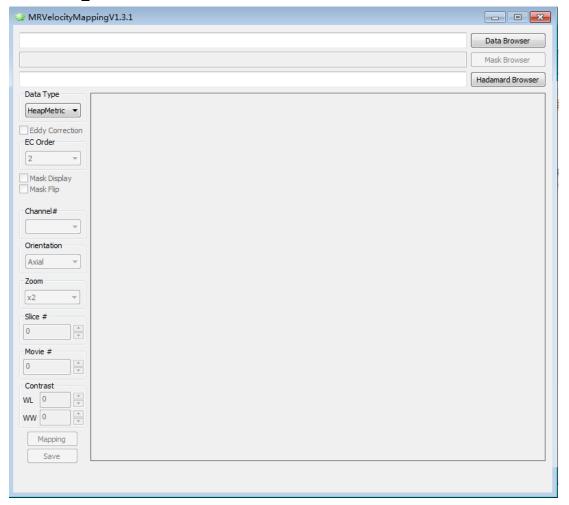


Figure 1

# 2. HEAP-METRIC Velocity Mapping

- 2.1 Reconstruction without mask file.
  - 2.1.1 Select the 'Data Type' box to be 'HeapMetric'.
  - 2.1.2 Click 'Data Browser' button. Then select a data folder which is acquired by using 'pcHeapMetric' sequence on Bruker ParaVision 6.0.1 system.
  - 2.1.3 Click 'Hadamard Browser' button. Then select a Hadamard Matrix file '\*.sqmat'.
  - 2.1.4 Click 'Mapping' Button. Then results will be reconstructed on 5 channels: magnitude, phase raw (no unwrapped), unwrapped phase, velocity map (with

- sign), absolute velocity map (no sign). Select 'Channel#' box to display different channel results shown as Fig.2.
- 2.1.5 Click 'Save' button to save results. Only the results on current selected channel would be saved. If you hope to save all channel results then you have to change the 'Channel#' and click save button repeatedly. All results are saved as 'Analyze 7.5' format (\*.hdr +\*.img).



Figure 2

- 2.2 Reconstruction with mask (With eddy current filed correction).
  - 2.2.1 Use images reconstructed from **Step 2.1** to manually trace a mask on global static tissues. Then saved your mask as **'Analyze 7.5'** format (\*.hdr +\*.img).
  - 2.2.2 Repeat Steps 2.1.1 2.1.3
  - 2.2.3 Click **'Eddy Correction'**, then click **'Mask Browser'** button. Chose the mask file saved from **Step 2.2.1**.
  - 2.2.4 Repeat **Step 2.1.4**. (Optional operations: now you can change **'EC Order'** to change eddy field fitting harmonics; you can click **'Mask Display'** button to show mask to confirm the match like as **Fig.3**; you can click **'Mask Flip'** to flip images for sometimes mask saved are reversed on phase encoding orientation). You can

#### Click 'Mapping' button over and over.

2.2.5 Save Results. See in Step 2.1.5.

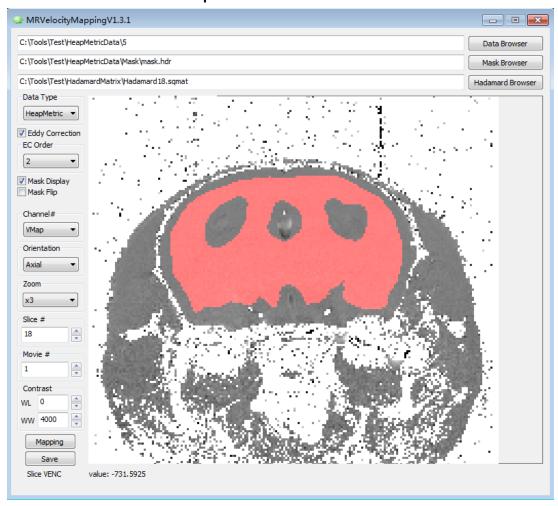


Figure 3

### 3. Trigger Cine Velocity Mapping

- 3.1 Reconstruction without mask file.
  - 3.1.1 Select the 'Data Type' box to be 'TriggerCine'.
  - 3.1.2 Click 'Data Browser' button. Then select a data folder which is acquired by using 'FLOWMAP' sequence with ECG trigger on Bruker ParaVision 6.0.1.
  - 3.1.3 Click 'Mapping' button. See in Step 2.1.4
  - 3.1.4 Save Results. See in **Step 2.1.5**. Results would be save as 4D format. The 4<sup>th</sup> dimension is the trigger delay time ordered.
- 3.2 Reconstruction with a mask file (With eddy current filed correction).
  - 3.2.1 Use images reconstructed from **Step 3.1** which is similar to **Step 2.2.1**.
  - 3.2.2 Repeat Steps 3.1.1 and 3.1.2
  - 3.2.3 Click 'Eddy Correction', then click 'Mask Browser' button. Chose the mask

file saved from **Step 3.2.1**. For single slice data **'EC order'** is better to be 1 or 0 (Show in Fig.4).

3.2.4 Repeat Steps 3.1.3 and 3.1.4.

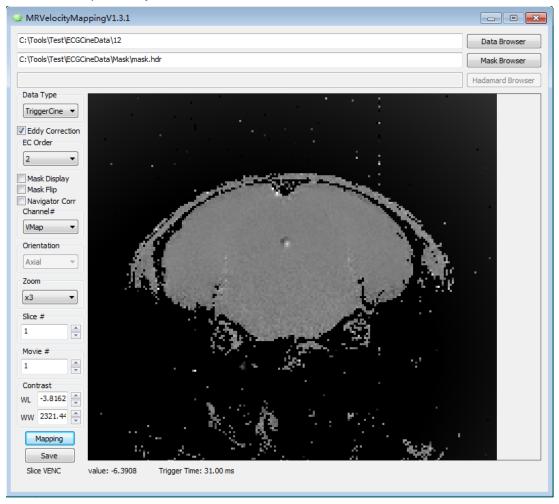


Figure 4

## 4. Directional Velocity Encoding Results Combination.

4.1 Repeat **Step 2.2** or **Step 3.2** to obtain three directional velocity maps (Read, Phase and Slice). Put the three results in a folder together. (See in Fig.5).

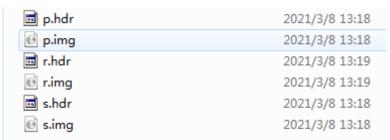
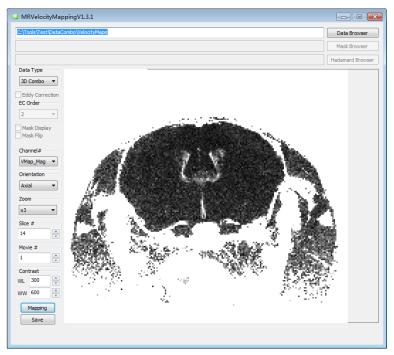


Figure 5

- 4.2 Select the 'Data Type' box to be '3D Combo'.
- 4.3 Click 'Data Browser' button. Then select a data folder containing three

directional velocity maps that is from Step 4.1.

- 4.4 Click 'Mapping' button. Then results will be reconstructed on 5 channels: Velocity Maps-Read, Velocity Maps-Phase, Velocity Maps-Slice, Velocity Maps-Magnitude (SOS combination), Velocity Maps-RGB (vector display). Show in Fig.6.
- 4.5 Click 'Save' button to save results which is similar to Step 2.1.5. What is the different is when the current 'Channel #' is 'VMap-RGB' then data will be saved as \*.bmp files.



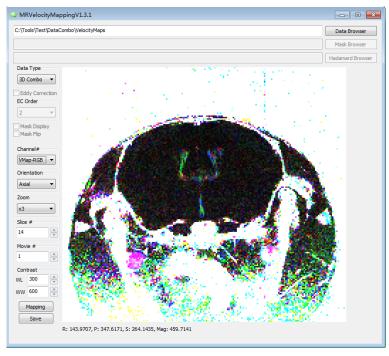


Figure 6