# Report on Participation in the IronTract Challenge 2019

Nagesh Adluru<sup>1</sup>, Barbara B. Bendlin<sup>1</sup>, Vivek Prabhakaran<sup>1</sup>, and Andrew L. Alexander<sup>1</sup>

<sup>1</sup>University of Wisconsin-Madison

Essential details of processing and analysis, and some observations from participating in the IronTract challenge 2019 are presented in this report. Our approach ranked #2 on the training dataset for both the HCP (multi-shell) and overall (DSI) tracks. The ranking on the validation dataset placed our approach at #2 for the overall track, and #3 for the HCP track.

DESIGNER, MSMT\_CSD, iFOD2, heuristic length and include filtering Correspondence: adluxu@wisc.edu

## **Essential details**

The entire exercise was performed using tools from MRtrix3 (1), FSL (2), ANTS (3), and GNU parallel (4). The data was pre-processed through the DESIGNER (5) preprocessing which includes denoising (6, 7), deringing (8), B1 bias correction, and Rician correction. Constrained spherical deconvolution (9) was then performed using the multi-shell mutitissue response function (MSMT\_CSD) (10). The HCP resampled data was used for both the HCP and overall competitions. The seed region was binarized without any additional filtering.

Tractography was performed using the tokgen tool in MRtrix3. The main options were iFOD2, different angle thresholds from 10° through 90°, default stepsize of 0.35, and min and max length thresholds. The basic code, with a few additional parameters, for the tractography is shown below.

Six planar include filters, two for each of the x,y,z coordinates, were used to avoid looping tracts. Then, tract density images (TDI) (11) were created using the tckmap command. A set of thresholds were applied to each of the TDI image to include portions of the ROC curve for FPR $\in$  [0,0.3], which turned out to be a key factor for getting a competitive AUC. The TDIs from different angle thresholds were averaged for each of the threshold setting, which formed the final set of maps that were uploaded for scoring.

### **Observations**

The overall experience of participating in this challenge was akin to learning to drive a race car, with all its gears and gadgets to gain an edge in second or third decimal pace in the AUC scores. Most of the settings off the lot, seemed to perform quite competitively. The core tractography algorithms such as iFOD2 are constrained only locally, but to achieve a good score, imposing global constraints, such as using include filters seemed to have helped. It seemed like, if one could pick the settings of any tractography algorithm and execute it to generate an accurate and complete white matter mask, it would match the histological tracing perfectly. It is amazing that the TPR reaches 1 by around FPR of 0.3, implying that the AUC is on average above 0.93, actually. Perhaps because of this, simple default settings were sufficient.

### **ACKNOWLEDGEMENTS**

The following grants are acknowledged for providing support for this participation which enabled us to develop a good empirical understanding of the tractography tools available in MRtrix3, which in turn would help the main aims of these grants. Core grant to the Waisman Center from the National Institute of Child Health and Human Development (IDDRC U54 HD090256), NIH grants R01 NS092870, BRAIN Initiative R01-EB022883, UW CPCP Al117924, R01AG037639, R01AG027161, RF1 AG059312, P50AG033514, R01-NS105646, R01 AG037639, UF1 AG051216.

# **Bibliography**

- J-Donald Tournier, Robert Smith, David Raffelt, Rami Tabbara, Thijs Dhollander, Maximilian Pietsch, Daan Christiaens, Ben Jeurissen, Chun-Hung Yeh, and Alan Connelly. Mrtrix3: A fast, flexible and open software framework for medical image processing and visualisation. NeuroImage, page 116137, 2019.
- Mark Jenkinson, Christian F Beckmann, Timothy EJ Behrens, Mark W Woolrich, and Stephen M Smith. FSL. Neuroimage, 62(2):782–790, 2012.
- Brian B Avants, Nick Tustison, and Gang Song. Advanced normalization tools (ants). *Insight* j, 2:1–35, 2009.
- A. Ole Tange. GNU Parallel 2018. Ole Tange, March 2018. ISBN 9781387509881. doi: 10.5281/zenodo.1146014.
- Benjamin Ades-Aron, Jelle Veraart, Peter Kochunov, Stephen McGuire, Paul Sherman, Elias Kellner, Dmitry S Novikov, and Els Fieremans. Evaluation of the accuracy and precision of the diffusion parameter estimation with gibbs and noise removal pipeline. NeuroImage, 183:532–543, 2018.
- Jelle Veraart, Els Fieremans, and Dmitry S Novikov. Diffusion mri noise mapping using random matrix theory. Magnetic resonance in medicine, 76(5):1582–1593, 2016.
- Jelle Veraart, Dmitry S Novikov, Daan Christiaens, Benjamin Ades-Aron, Jan Sijbers, and Els Fieremans. Denoising of diffusion mri using random matrix theory. *NeuroImage*, 142: 394–406, 2016.
- Elias Kellner, Bibek Dhital, Valerij G Kiselev, and Marco Reisert. Gibbs-ringing artifact removal based on local subvoxel-shifts. Magnetic resonance in medicine, 76(5):1574–1581, 2016.
- J-Donald Tournier, Fernando Calamante, and Alan Connelly. Robust determination of the fibre orientation distribution in diffusion mri: non-negativity constrained super-resolved spherical deconvolution. *Neuroimage*, 35(4):1459–1472, 2007.
- Ben Jeurissen, Jacques-Donald Tournier, Thijs Dhollander, Alan Connelly, and Jan Sijbers Multi-tissue constrained spherical deconvolution for improved analysis of multi-shell diffusion mri data. NeuroImage, 103:411–426, 2014.
- 11. Fernando Calamante, Jacques-Donald Tournier, Graeme D Jackson, and Alan Connelly

Track-density imaging (tdi): super-resolution white matter imaging using whole-brain track-density mapping. *Neuroimage*, 53(4):1233–1243, 2010.



 $2 \mid \text{bio} \frac{\textbf{R}}{\chi}$ iv et al.  $\mid$  ITC