

TRX: A community-oriented tractography file format

Francois Rheault^a, Valérie Hayot-Sasson^b, Robert E. Smith^{c1,c2}, Christopher Rorden^d, Jacques-Donald Tournier^e, Eleftherios Garyfallidis^f, Fang-Cheng Yeh^h, Christopher J. Markiewiczⁱ, Matthew Brett^j, Ben Jeurissen^k, Paul A. Taylor^l, D. Baran Aydogan^{m1,m2}, Derek A. Pisnerⁿ, Serge Koudoro^f, Soichi Hayashi^o, Daniel Haehn^p, Steve Pieper^q, Daniel Bullock^r, Emanuele Olivetti^s, Jean-Christophe Houde^u, Marc-Alexandre Côtéⁱ, Flavio Dell'Acqua^x, Alexander Leemans^y, Maxime Descoteaux^{u,v}, Bennett Landman^{a,w}, Franco Pestilli^g, and Ariel Rokem^z

Affiliations:

- ^aElectrical and Computer Engineering, Vanderbilt University, Nashville, USA;
^bDepartment of Computer Science and Software Engineering, Concordia University, Montreal, Canada
^{c1}Florey Institute of Neuroscience and Mental Health, Melbourne, Australia
^{c2}Florey Department of Neuroscience and Mental Health, The University of Melbourne, Melbourne, Australia
^dDepartment of Psychology, University of South Carolina, Columbia, USA
^eBiomedical Engineering Department, School of Biomedical Engineering and Imaging Sciences, King's College London, UK
^fDepartment of Intelligent Systems Engineering, Indiana University, Bloomington, USA
^gDepartment of Psychology, The University of Texas at Austin, Austin, USA.
^hDepartment of Neurological Surgery, University of Pittsburgh, Pittsburgh, USA
ⁱDepartment of Psychology, Stanford University, Stanford, CA, USA
^jBrain Imaging Center, University of California, Berkeley, Berkeley, CA, USA
^kDepartment of Physics, University of Antwerp, Antwerp, Belgium
^lScientific and Statistical Computing Core, National Institute of Mental Health, USA
^{m1}A.I. Virtanen Institute for Molecular Sciences, University of Eastern Finland, Kuopio, Finland
^{m2}Department of Neuroscience and Biomedical Engineering, Aalto University, Espoo, Finland
ⁿUniversity of Texas at Austin, Austin, USA
^oDepartment of Psychological and Brain Sciences, Indiana University, Bloomington, USA
^pMachine Psychology Lab, Department of Computer Science, University of Massachusetts, Boston, USA
^qDepartment of Neurosurgery, Brigham and Women's Hospital, Harvard Medical School, Boston, United States
^rCenter for Magnetic Resonance Research, University of Minnesota
^sNeuroinformatics Laboratory, Fondazione Bruno Kessler, Trento, Italy
^tMicrosoft Research Lab, Montréal, Canada
^uImeka Solutions, Sherbrooke, Canada
^vSherbrooke Connectivity Imaging Laboratory (SCIL), Département d'Informatique, Université de Sherbrooke, Canada;
^wVanderbilt University Institute of Imaging, Nashville, USA;
^xNatBrainLab, Dept. of Forensics and Neurodevelopmental Sciences, King's College London, London, UK
^yPROVIDI Lab, Image Sciences Institute, UMC Utrecht, Utrecht, The Netherlands
^zDepartment of Psychology and eScience Institute, University of Washington, Seattle, USA

Introduction

File formats that store the results of computational tractography were typically developed within specific software packages. This approach has facilitated a myriad of applications, but this development approach has generated insularity within software packages, and has limited standardization. Moreover, because tractography file formats were developed to solve immediate challenges, only a limited breadth of applications within a single software package was envisioned, sometimes also neglecting computational performance. Given the growing interest in tractography methods and applications, and the increasing size and complexity of datasets, a community-driven standardization of tractography have become a priority. To address these challenges, our community initiated a discussion to design a new file format and agreed to participate in its conception, development, and, if successful, its adoption. Here, we propose **TRX**, a tractography file format designed to facilitate dataset exchange, interoperability, and state-of-the-art analyses, acting as a community-driven replacement for **TCK** [1], **TRK** [2], **VTK**, **TT** [3], **TRAKO** [4], **qFIB** [5], **zFIB** [6], **NIML.TRACT** [10] and **DPY** [8].

Background

Following discussions on GitHub (<https://github.com/nipy/nibabel/issues/942>), during a videoconference between the developers of several software libraries for tractography, and at the DIPY online group meeting, we identified key features that were desired by the community (both developers and users):

1. Maintain functionality currently implemented by other file formats;
2. Computational efficiency (both in speed and file size);
3. Extensibility so the file format could evolve more easily.

Feedback from different groups showed that developers desired features such as being based on open-source, well-maintained libraries, and robustness, while users prioritized new functionalities, simplicity, and low memory usage.

Results

A TRX file format candidate specification was designed to favor simplicity and human readability. Its data are organized following an intuitive naming convention that conveys dimensionality and data type. The specification allows storage of both data per vertex and per streamline (similar to TRK), streamlines-group, with both sparse and dense group indices representation. RASMM (world/scanner coordinates) for vertex coordinates is proposed (similar to TCK). The file format supports random-access with minimal upfront RAM requirements, and other low-level operations with virtually no RAM requirements, by using a data and offsets approach (similar to VTK9) and raw binary arrays that allow memory mapping. The file format supports choice over the data type for each raw data array (see Figure 1) and, optional, ZIP compression. Just like other file formats, efficient streamline compression [6] via linearization method is possible without modifying the current specification.

An initial benchmark for operations like partial file loading (Figure 2), concatenation (not shown), affine transformation (not shown), and file saving (Figure 2) was performed using the Python implementation and compared to Nibabel. A C++ prototype was also developed and proposed for community feedback.

Conclusion

The goal of TRX is to become the first, community-driven, standard amongst tractography file formats. As with other file formats like NiFTI [9], we believe that TRX will serve the community well and the growing computational needs of our field. We encourage community members to consider early contributions to our proposal so as to ensure the new standard will cover the needs of the wider audience of software developers, toolboxes, and scientists. Our long-term plan is to integrate TRX within the Brain Imaging Data Structure (BIDS) ecosystem [8]. Community contributions and related discussions are encouraged using: https://github.com/frheault/tractography_file_format/issues.

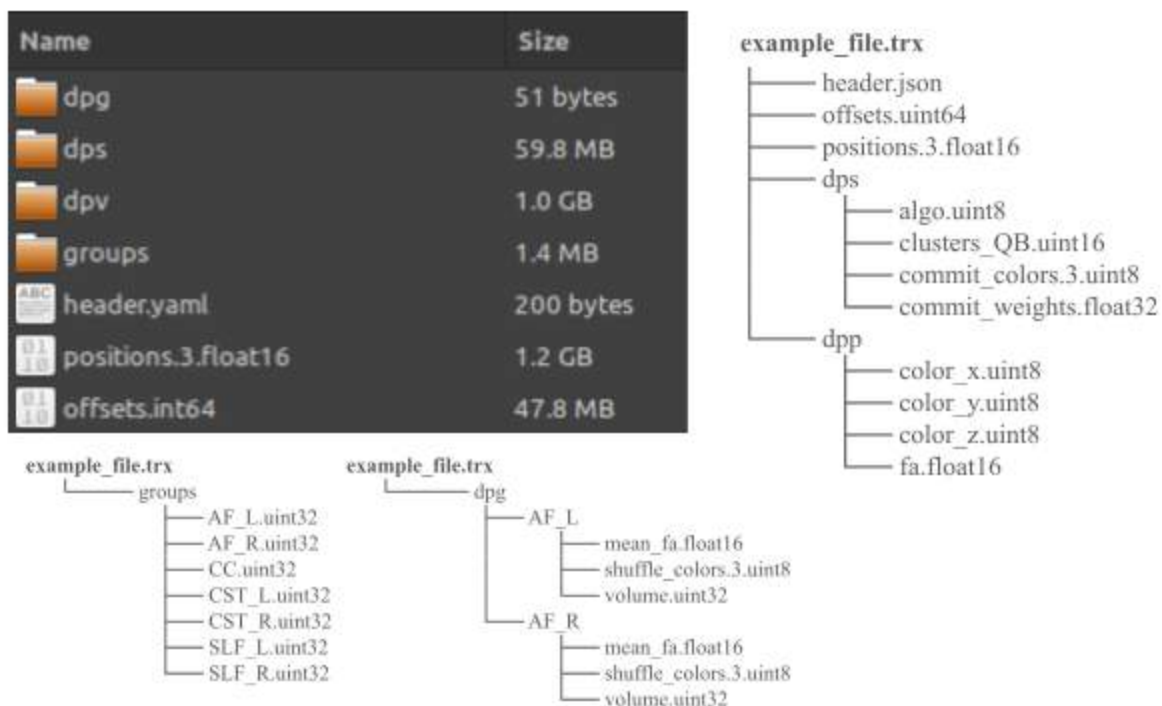


Figure 1: File structure of the TRX file format. A filename convention conveys the meaning, dimensionality and data type for each data array. The header, positions (coordinates of streamlines) and offsets (starting position of streamlines in the array) are mandatory files. Optional data include the data per streamline (dps), data per vertex (dpv), groups and data per group (dpg). Specifications can be found at https://github.com/frheault/tractography_file_format/blob/master/trx_file_memmap/specifications.md.

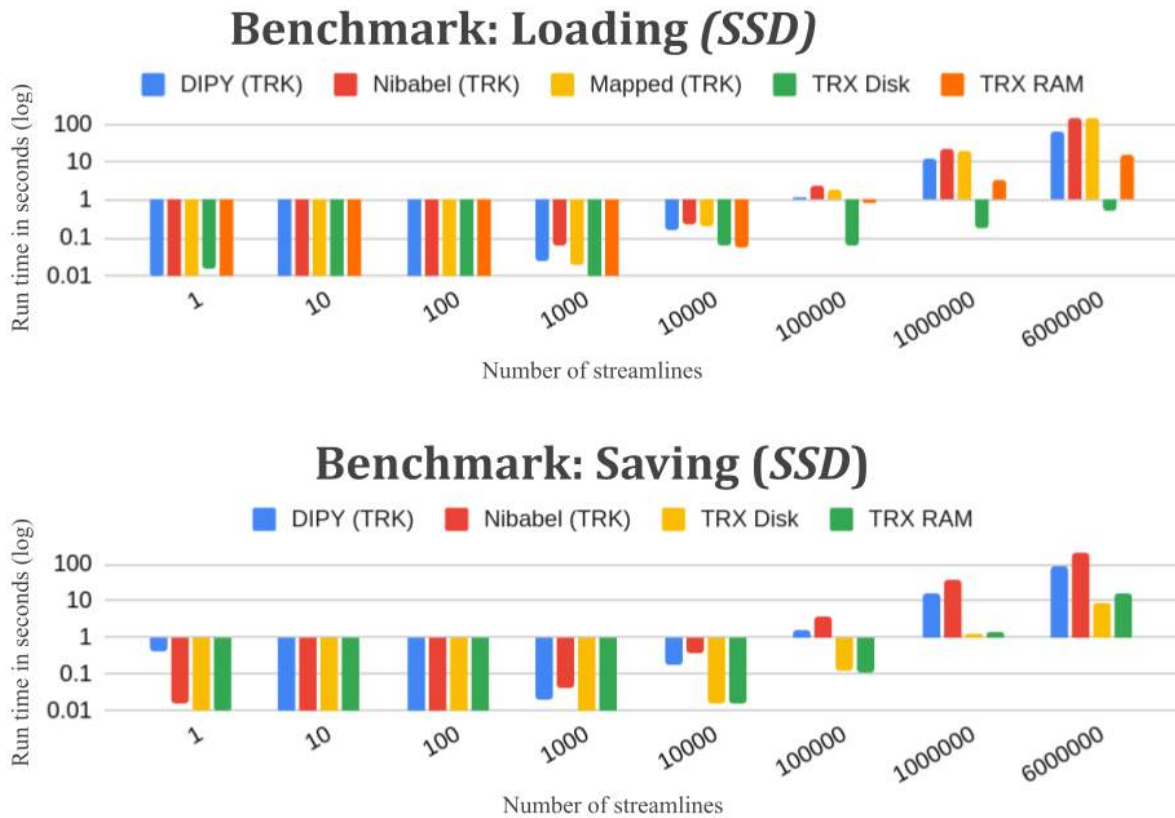


Figure 2: Load times (in seconds) for loading and saving a small (1 streamline) to a large file (6M streamlines). To facilitate visualization and comparison, the graph is shown on a logarithmic scale. *DIPY* is a Python implementation from DIPY [7] that does not support data per streamline / data per vertex (deprecated, TRK). *Nibabel* is the Nibabel API (version 3.0, TRK). *Mapped* is an implementation that allows random access (available on https://github.com/emanuele/load_trk, TRK). The loading operation of *TRX Disk* is in fact, simply the memory mapping initialization, there is no “loading”. For the saving operation, *TRX Disk* is simply a copy-paste operation, which showcases the baseline disk speed. *TRX RAM* is the memory mapping initialization and transfer of all data from disk to RAM. Python implementations are likely slower than the implementations of TrackVis or MI-Brain, but this nevertheless showcases the advantage of memory mapping even for slower languages. The Python prototype used for this benchmark can be found at https://github.com/frheault/tractography_file_format/tree/master/trx_file_memmap.

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Logo (optional)



