Neuroconductor: An R Platform for Medical Imaging Analysis



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What is Neuroconductor?

Neuroconductor (https://neuroconductor.org/) is a a centralized repository of R software dedicated to medical image analysis.

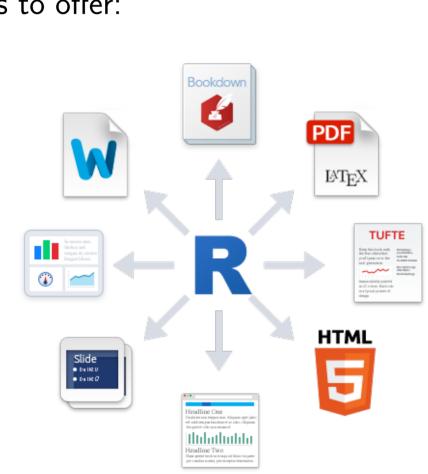
Goals of Neuroconductor

- Disseminate quickly software updates
- Educate a large, diverse community of scientists using detailed tutorials and short courses
- Ensure quality via automatic and manual quality controls
- Promote the reproducibility of image data analysis

Benefits of Imaging in R

Allow medical imaging to use all R has to offer:

- Statistics and Machine Learning
- Package versioning, testing, and distribution
- Reproducibile reports and analyses (knitr and rmarkdown)
- Shiny applications for the web



(Image from http://rmarkdown.rstudio.com/images/RMarkdownOutputFormats.ng)

Potential Downsides to Neuroconductor

- More control over the workflow = more work (e.g. for statisticians)
- Users need external software (versions/installation)
- No control over external software
- if maintainer changes something, not much recourse
- Need the content (buy-in from the imaging/R communities)

References

- [1] Brandon Whitcher, Volker J. Schmid, and Andrew Thornton. "Working with the DICOM and NIfTI Data Standards in R". In: Journal of Statistical Software 44.6 (2011), pp. 1–28.
- Xiangrui Li et al. "The first step for neuroimaging data analysis: DICOM to NIfTI conversion". In: Journal of
- Neuroscience Methods 264 (2016), pp. 47–56.

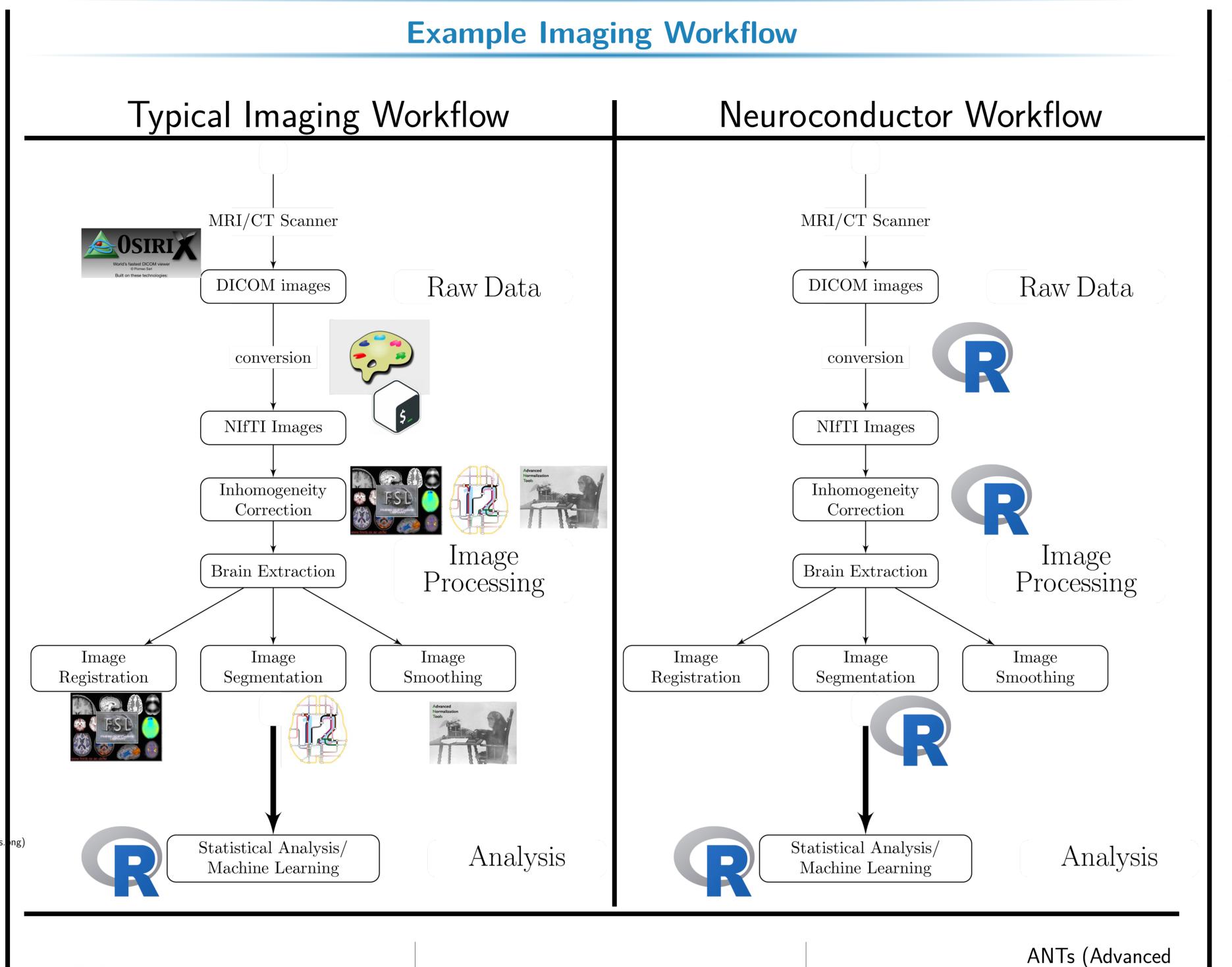
[3] Dirk Eddelbuettel et al. "Rcpp: Seamless R and C++ integration". In: Journal of Statistical Software 40.8 (2011),

- [4] Stephen M Smith et al. "Advances in functional and structural MR image analysis and implementation as FSL".
- In: Neuroimage 23 (2004), S208–S219. John Muschelli et al. "fslr: Connecting the FSL Software with R". In: The R Journal 7.1 (2015), pp. 163–175.
- Bruce Fischl. "FreeSurfer". In: Neuroimage 62.2 (2012), pp. 774–781.
- [7] William D Penny et al. Statistical parametric mapping: the analysis of functional brain images. Academic press,

[8] B. B. Avants et al. "A Reproducible Evaluation of ANTs Similarity Metric Performance in Brain Image Registra-

- tion". In: Neurolmage 54.3 (2011), 2033—2044. [9] Russell T Shinohara et al. "Statistical normalization techniques for magnetic resonance imaging". In: NeuroImage:
- [10] Bennett A Landman et al. "Multi-parametric neuroimaging reproducibility: a 3-T resource study". In: Neuroimage 54.4 (2011), pp. 2854–2866.
- [11] Kenichi Oishi et al. "Atlas-based whole brain white matter analysis using large deformation diffeomorphic metric mapping: application to normal elderly and Alzheimer's disease participants". In: Neuroimage 46.2 (2009),
- pp. 486–499. [12] Vladimir Fonov et al. "Unbiased average age-appropriate atlases for pediatric studies". In: NeuroImage 54.1
- [13] Vladimir S Fonov et al. "Unbiased nonlinear average age-appropriate brain templates from birth to adulthood".
- [14] Bennett Allan Landman et al. MICCAI 2012 Workshop on Multi-Atlas Labeling. CreateSpace Independent Publishing Platform, 2012.
- [15] David C Van Essen et al. "The WU-Minn human connectome project: an overview". In: Neuroimage 80 (2013),
- pp. 62–79.

Clinical 6 (2014), pp. 9–19.





malf.templates

hcp

bash - shell scripting is usually required for command-line tools or pipelining

MRIcroGL - imaging

analysis suite, with

dcm2nii - DICOM to

NIfTI software



≜0SIRI**X**

FSL (FMRIB Software Library) suite of neuroimaging analysis tools

OsiriX - standalone

DICOM viewer



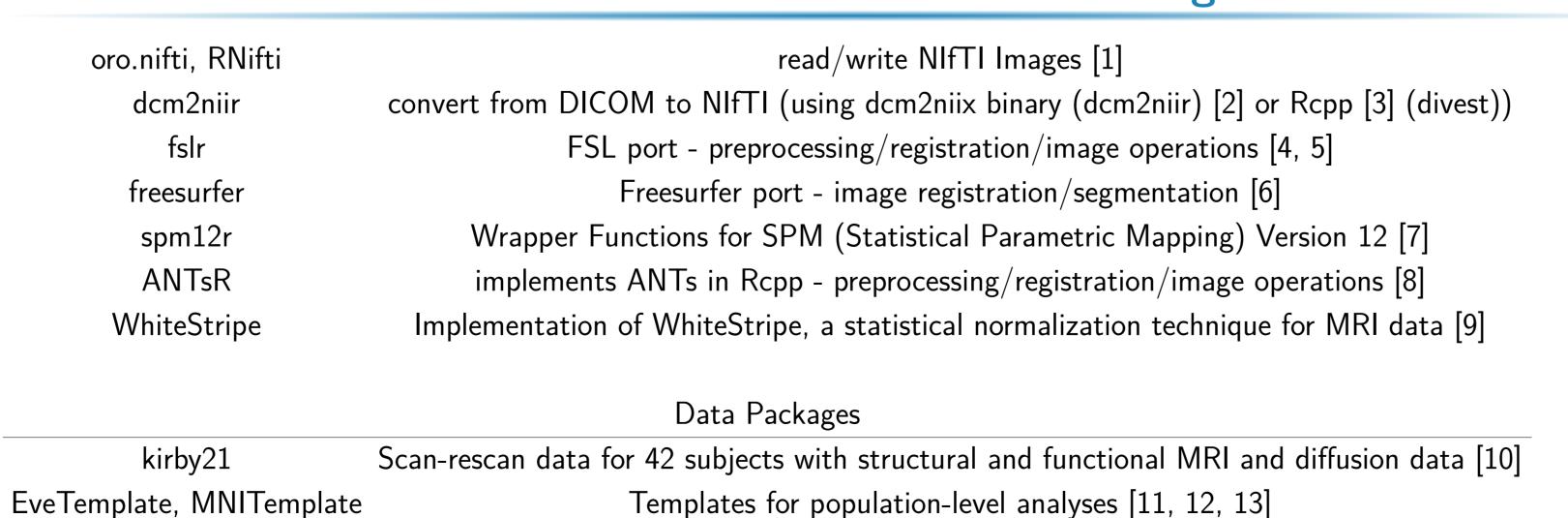
- state-of-the-art tools for neuroimaging analysis SPM 12 - statistical parametric mapping, requires MATLAB (Mathworks, Natick,

Normalization Tools)



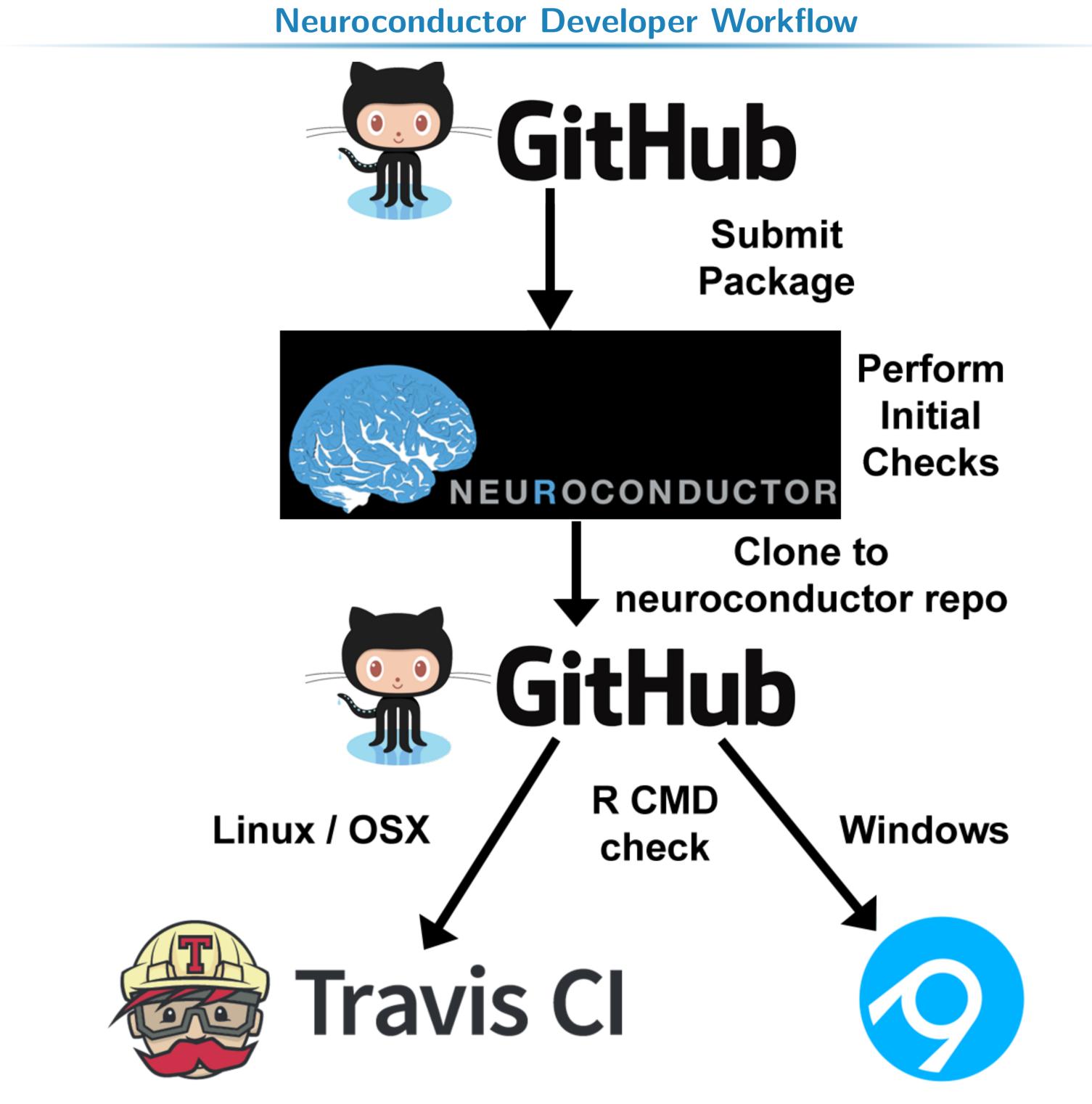
Massachusetts, USA) - analysis tools for PET/SPECT/fMRI

Overview of Neuroconductor Packages



Templates [14] for Multi-Atlas Label Fusion (MALF) and Skull Stripping

Download data from the Human Connectome Project [15]





GitHub - a online hosting service of git repositories. All Neuroconductor packages are hosted on GitHub.



Travis Cl

Before uploading to GitHub, checks are performed, a confirmatory email is sent (reduce spam), and Travis/Appveyor configuration files are added

Travis CI (continuous integration) - an online service of Linux/Mac OSX virtual machines that build and check pack-



AppVeyor - a similar CI service that builds and checks packages on Windows

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