John Muschelli

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Education

2012–2016 PhD, Biostatistics,

Johns Hopkins Bloomberg School of Public Health, Baltimore, MD.

Computational Methods for Neuroimaging in R: Stroke Hemorrhage in X-ray Computed Tomography Scanning. Advisor: Ciprian Crainiceanu, PhD

Relevant Experience

2016— Assistant Scientist, Department of Biostatistics, Johns Hopkins Bloomberg School Present of Public Health, (Research-track Faculty).

Peer-Reviewed Publications from past 3 years

* denotes authors contributed equally

2020 Muschelli, J. (2020). "A publicly available, high resolution, unbiased CT brain template". International Conference on Information Processing and Management of Uncertainty in Knowledge-Based Systems, pp. 358-366.

Ding, T (2020). "An improved algorithm of white matter hyperintensity detection in elderly adults". NeuroImage: Clinical 25, p. 102151.

Garibaldi, B. T. (2020). "Patient trajectories among persons hospitalized for COVID-19: a cohort study". Annals of Internal Medicine.

Hannawi, Y. (2020). "Postcardiac arrest neurological prognostication with quantitative regional cerebral densitometry". Resuscitation.

Hansen, B. M. (2020). "Relationship of white matter lesions with intracerebral hemorrhage expansion and functional outcome: MISTIE II and CLEAR III". Neurocritical Care, pp. 1-9.

Kross, S. (2020). "ari: The Automated R Instructor". The R Journal 12.1, pp. 258–265. Leroux, A. (2020). "Quantifying the predictive performance of objectively measured physical activity on mortality in the UK Biobank". The Journals of Gerontology: Series Α.

Minhas, D. S. (2020). "Statistical methods for processing neuroimaging data from two different sites with a Down syndrome population application". International Conference on Information Processing and Management of Uncertainty in Knowledge-Based Systems, pp. 367-379.

Rothstein, J. D. (2020). "'The doctor said formula would help me': health sector influences on use of infant formula in peri-urban Lima, Peru". Social Science & Medicine 244.C.

Sharrock, M. (2020). "3D deep neural network segmentation of intracerebral hemorrhage: development and validation for clinical trials". medRxiv.

Valcarcel, A. M. (2020). "TAPAS: a thresholding approach for probability map automatic segmentation in multiple sclerosis". NeuroImage: Clinical.

2019 **Muschelli, J.** (2019b). "ROC and AUC with a binary predictor: a potentially misleading metric". Journal of Classification, pp. 1–13.

Muschelli, J. (2019a). "Recommendations for processing head CT data". *Frontiers in Neuroinformatics* 13, p. 61.

Hadavand, A. (2019). "Analysis of student behavior using the R package crsra". *Journal of Learning Analytics* 6.2, pp. 140–152.

Ryan, S. M. (2019). "Template creation for high-resolution computed tomography scans of the lung in R software". *Academic Radiology*.

2018 **Muschelli**, **J.** (2018). "freesurfer: connecting the Freesurfer software with R". *F1000Research* 7.

Muschelli, J. (2018). "Neuroconductor: an R platform for medical imaging analysis". *Biostatistics*, kxx068. eprint: /oup/backfile/content_public/journal/biostatistics/pap/10.1093_biostatistics_kxx068/1/kxx068.pdf.

Commowick, O. (2018). "Objective evaluation of multiple sclerosis lesion segmentation using a data management and processing infrastructure". *Scientific Reports*.

Smith, C. H. (2018). "Feasibility of coping effectiveness training for caregivers of children with autism spectrum disorder: a genetic counseling intervention". *Journal of genetic counseling* 27.1, pp. 252–262.

Valcarcel, A. (2018c). "TAPAS: threshold adjustment to probability map automatic segmentations". *MULTIPLE SCLEROSIS JOURNAL*. Vol. 24, pp. 629–630.

Valcarcel, A. M. (2018b). "MIMoSA: an automated method for intermodal segmentation analysis of multiple sclerosis brain lesions". *Journal of Neuroimaging*.

Valcarcel, A. M. (2018a). "MIMoSA: an approach to automatically segment T2 hyperintense and T1 hypointense lesions in multiple sclerosis". *International MICCAI Brainlesion Workshop*, pp. 47–56.

Software: R Packages

All download counts are from RStudio CRAN logs and are accurate as of October 13, 2020.

rscopus: Scopus Database API Interface, 67336.

fslr: Wrapper Functions for FSL (FMRIB Software Library) from Functional MRI of the Brain (FMRIB), 38196.

neurobase: Neuroconductor Base Package with Helper Functions for nifti Objects, 33972.

brainR: Helper Functions to misc3d and rgl Packages for Brain Imaging, 30298.

matlabr: An Interface for MATLAB using System Calls, 28363.

diffr: Display Differences Between Two Files using Codediff Library, 26895.

WhiteStripe: White Matter Normalization for Magnetic Resonance Images using WhiteStripe, 24097.

kirby21.base: Example Data from the Multi-Modal MRI Reproducibility Resource, 17085.

gifti: Reads in Neuroimaging GIFTI Files with Geometry Information, 17025.

freesurfer: Wrapper Functions for Freesurfer, 16661.

spm12r: Wrapper Functions for SPM (Statistical Parametric Mapping) Version 12 from the Wellcome Trust Centre for Neuroimaging, 16041.

gcite: Google Citation Parser, 15853.

mscstts: R Client for the Microsoft Cognitive Services

Text-to-Speech REST API, 15762.

papayar: View Medical Research Images using the Papaya

JavaScript Library, 14853.

cifti: Toolbox for Connectivity Informatics Technology Initiative

(CIFTI) Files, 14699.

kirby21.t1: Example T1 Structural Data from the Multi-Modal MRI

Reproducibility Resource, 13857.

kirby21.fmri: Example Functional Imaging Data from the Multi-Modal

MRI Reproducibility Resource, 13315.

neurohcp: Human Connectome Project Interface, 12702.

glassdoor: Interface to Glassdoor API, 12604.

stapler: Simultaneous Truth and Performance Level Estimation, 11337.

fedreporter: Interface to Federal RePORTER API, 11216.

text2speech: Text to Speech, 10000. leanpubr: Leanpub API Interface, 8600.

nsrr: Interface to National Sleep Research Resource, 7570.

neurovault: Neurovault Database API Access, 703.