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

2020— Associate 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
 - 2023 Savonen, C. (2023). "Open-source tools for training resources-OTTR". Journal of Statistics and Data Science Education 31.1, pp. 57–65.
 - Zhao, C. (2023). "ModelArray: an R package for statistical analysis of fixel-wise data". Neuroimage 271, p. 120037.
 - 2022 Karas, M. (2022). "Comparison of accelerometry-based measures of physical activity: retrospective observational data analysis study". JMIR mHealth and uHealth 10.7, e38077.
 - Pham, D. D. (2022). "ciftiTools: a package for reading, writing, visualizing, and manipulating CIFTI files in R". Neurolmage 250, p. 118877.
 - Qi, G. (2022). "Genome-wide association studies of 27 accelerometry-derived physical activity measurements identified novel loci and genetic mechanisms". Genetic epidemiology 46.2, pp. 122-138.
 - Redd, A. D. (2022). "Curating the evidence about COVID-19 for frontline public health and clinical care: the novel coronavirus research compendium". Public Health Reports 137.2, pp. 197-202.
 - Sharrock, M. F. (2022). "Bayesian deep learning outperforms clinical trial estimators of intracerebral and intraventricular hemorrhage volume". Journal of Neuroimaging 32.5, pp. 968-976.
 - Smirnova, E. (2022). "Predictive performance of selected breath volatile organic carbon compounds in stage 1 lung cancer". Translational Lung Cancer Research 11.6, p. 1009.
 - 2021 Bennett, T. D. (July 2021). "Clinical Characterization and Prediction of Clinical Severity of SARS-CoV-2 Infection Among US Adults Using Data From the US National COVID Cohort Collaborative". JAMA Network Open 4.7, e2116901–e2116901. eprint: https://jamanetwork.com/journals/jamanetworkopen/articlepdf/2781923/ $bennett_2021_oi_210506_1625605077.56265.pdf.$
 - Bowring, M. G. (2021). "Outcome-stratified analysis of biomarker trajectories for patients infected with severe acute respiratory syndrome coronavirus 2". American journal of epidemiology 190.10, pp. 2094-2106.

Broll, S. (2021). "Interpreting blood glucose data with R package iglu". *PloS one* 16.4, e0248560.

Garibaldi, B. T. (2021). "Patient trajectories among persons hospitalized for COVID-19: a cohort study". *Annals of Internal Medicine* 174.1, pp. 33–41.

Leroux, A. (2021). "Quantifying the predictive performance of objectively measured physical activity on mortality in the UK Biobank". *The Journals of Gerontology: Series A* 76.8, pp. 1486–1494.

Scully, E. P. (2021). "Sex and gender differences in testing, hospital admission, clinical presentation, and drivers of severe outcomes from COVID-19". *Open forum infectious diseases*. Vol. 8. 9, ofab448.

Sharrock, M. F. (2021). "3D deep neural network segmentation of intracerebral hemorrhage: development and validation for clinical trials". *Neuroinformatics* 19.3, pp. 403–415.

Torbati, M. E. (2021). "A multi-scanner neuroimaging data harmonization using RAVEL and ComBat". *Neuroimage* 245, p. 118703.

Tustison, N. J. (2021). "The ANTsX ecosystem for quantitative biological and medical imaging". *Scientific reports* 11.1, pp. 1–13.

Wang, G. (2021). "Moderated t-tests for group-level fMRI analysis". *NeuroImage* 237, p. 118141.

Wongvibulsin, S. (2021). "Development of severe COVID-19 adaptive risk predictor (SCARP), a calculator to predict severe disease or death in hospitalized patients with COVID-19". *Annals of internal medicine* 174.6, pp. 777–785.

Wrobel, J. (2021). "Diurnal physical activity patterns across ages in a large UK based cohort: the UK Biobank study". *Sensors* 21.4, p. 1545.

Software: R Packages

All download counts are from RStudio CRAN logs and are accurate as of May 23, 2024.

rscopus: Scopus Database API Interface, 269857.

neurobase: Neuroconductor Base Package with Helper Functions for nifti Objects, 78671.

diffr: Display Differences Between Two Files using Codediff Library, 67396.

fslr: Wrapper Functions for FSL (FMRIB Software Library) from Functional MRI of the Brain (FMRIB), 60895.

matlabr: An Interface for MATLAB using System Calls, 48878.

brainR: Helper Functions to misc3d and rgl Packages for Brain Imaging, 48675.

gifti: Reads in Neuroimaging GIFTI Files with Geometry Information, 44738.

mscstts: R Client for the Microsoft Cognitive Services

Text-to-Speech REST API, 44289.

WhiteStripe: White Matter Normalization for Magnetic

Resonance Images, 40874.

text2speech: Text to Speech Conversion, 32086.

kirby21.base: Example Data from the Multi-Modal MRI

Reproducibility Resource, 32078.

gcite: Google Citation Parser, 31759.

freesurfer: Wrapper Functions for Freesurfer, 31560.

cifti: Toolbox for Connectivity Informatics Technology Initiative

(CIFTI) Files, 30545.

neurohcp: Human Connectome Project Interface, 29213.

spm12r: Wrapper Functions for SPM (Statistical Parametric Mapping) Version 12 from the Wellcome Trust Centre for Neuroimaging, 27548.

kirby21.t1: Example T1 Structural Data from the Multi-Modal MRI Reproducibility Resource, 26857.

kirby21.fmri: Example Functional Imaging Data from the Multi-Modal MRI Reproducibility Resource, 26386.

glassdoor: Interface to Glassdoor API, 25900.

papayar: View Medical Research Images using the Papaya

JavaScript Library, 24868.

stapler: Simultaneous Truth and Performance Level Estimation, 23880.

leanpubr: Leanpub API Interface, 19500.

nsrr: Interface to National Sleep Research Resource, 17065. fedreporter: Interface to Federal RePORTER API, 14928. neurovault: Neurovault Database API Access, 1677.