Supplementary materials for:Intensity warping for multisite MRI harmonization

J Wrobel^a, ML Martin^c, R Bakshi^{d,e}, PA Calabresi^f, M Elliot^g, D Roalf^h, RC Gur^{g,h,i}, RE Gur^{g,h,i}, RG Henry^j, G Nair^k, J Oh^{f,l}, N Papinutto^j, D Pelletier^j, DS Reich^{f,k}, WD Rooney^m, TD Satterthwaite^h, W Stern^j, K Prabhakaran^h,

NL Sicotteⁿ, RT Shinohara^c, J Goldsmith^b, on behalf of the NAIMS Cooperative^o

^aDepartment of Biostatistics and Informatics, Colorado School of Public Health, University of Colorado Anschutz Medical Campus

 b Department of Biostatistics, Mailman School of Public Health, Columbia University

 c Department of Biostatistics, Epidemiology, and Informatics, Perelman School of Medicine, University of Pennsylvania, Philadelphia, PA, USA

^dLaboratory for Neuroimaging Research, Partners Multiple Sclerosis Center, Department of Neurology, Brigham and Women's Hospital, Harvard Medical School, Boston, MA, USA ^eDepartment of Radiology, Brigham and Women's Hospital, Harvard Medical School, Boston, MA, USA

 f Department of Neurology, the Johns Hopkins University School of Medicine, Baltimore, MD, USA

^gDepartment of Radiology, Hospital of the University of Pennsylvania, Philadelphia, PA, 19104, USA

 h Brain Behavior Laboratory, Department of Psychiatry, Neuropsychiatry Section, Perelman School of Medicine, University of Pennsylvania, Philadelphia, PA, 19104, USA

 ${}^{i}L$ ifespan Brain Institute (LiBI) at the University of Pennsylvania and Children's Hospital of Philadelphia, Philadelphia, PA, 19104, USA

 j Department of Neurology, University of California - San Francisco, San Francisco, CA, USA

^kTranslational Neuroradiology Section, National Institute of Neurological Disorders and Stroke, National Institutes of Health, Bethesda, MD, USA

¹St. Michael's Hospital, University of Toronto, Toronto, Ontario, Canada

^mAdvanced Imaging Research Center, Oregon Health & Science University, Portland, OR, USA

ⁿDepartment of Neurology, Cedars-Sinai Medical Center, Los Angeles, CA, USA ^oThe North American Imaging in Multiple Sclerosis (NAIMS) Cooperative

1. Schematic of mica method

Figure [\(1\)](#page-1-0) shows a schematic of the mica process: images were bias corrected and skull-stripped, voxel-intensities were converted to CDFs, CDFs were

Preprint submitted to NeuroImage July 25, 2020

[∗]Corresponding author

Email address: julia.wrobel@cuanschutz.edu (J Wrobel)

aligned, and warping functions from CDF alignment were used to generate har-

⁵ monized images.

Figure 1: Harmonization pipeline. Raw images are N4 bias-corrected, skull-stripped, voxel intensities are converted to CDFs, CDFs are aligned by warping intensity values. The transformation of intensity values that produces this alignment is called a warping function, and the nonlinear transformation is applied to the raw images to produce harmonized images.

2. Expansion of manuscript Figure 1

Figure [2](#page-2-0) adds two panels to Figure 1 in the manuscript: a panel showing the histograms after White Stripe only and a panel showing the histograms after mica without White Stripe or other intensity normalization. This Figure shows ¹⁰ that both mica alone and White Stripe alone remove some scanner effects in the NAIMS data. With mica alone, the histograms are well aligned, but the final intensity domain depends on the scan chosen as the reference. With White Stripe alone, the intensities are in units of normal appearing white matter, but some areas of the histograms (lower intensity values, gray matter) are not well

¹⁵ aligned. *mica* plus White Stripe combines the superior histogram alignment of mica with the voxel intensity unit interpretability of White Stripe.

Figure 2: Histograms of NAIMS data for unnormalized and unharmonized images, White Stripe normalized images, mica harmonized images, and images that have been processed using White Stripe and mica.

3. PDFs of trio2prisma images across methods

Figure [\(3\)](#page-3-0) shows probability distribution functions (PDFs) of voxel intensities under different image harmonization and intensity normalization scenarios.

20 4. Sensitivity of leave N scans out

In order to understand how quickly the intensity warping function used to account for scanner effects stabilizes as the number of scans, N, used to construct this warping function increases. To examine this in the context of the trio2prisma study, we performed the following experiment.

- 25 1. Select 1 of the 10 subjects from the *trio2prisma* dataset. For number of scans $n \in 1:9$:
	- (a) Choose N subjects randomly from the remaining sample. For $N = 9$ this is the mica-loso process previously described.

Figure 3: Histograms of intensities before and after harmonization by tissue type in the trio2prisma study. Rows indicate tissue type, with whole brain, white matter, and gray matter shown in rows 1, 2, and 3, respectively. Columns correspond to different harmonization methods.

- (b) For each of the N subjects, calculate the warping function, $h_n^{-1}(x)$;
- 30 $n \in 1...N$, that maps the Trio scan to the Prisma scan for that subject.
	- (c) Calculate the leave $10 N$ scans out (lNso) intensity warping function, $h_{lNso}^{-1}(x)$, by averaging the *mica* warping function for the N selected subjects.
	- (d) For each N, calculate the root integrated square error (RISE) between the intensity warping function $h_n^{-1}(x)$ and the lNso warping function $h_{lNso}^{-1}(x)$ via

$$
\sqrt{\int \left[\left(h_n^{-1}(x) - h_{INso}^{-1}(x) \right) dx \right]^2}.
$$

³⁵ 2. Repeat steps (i) to (iii) for all 10 subjects.

Figure [4](#page-4-0) shows boxplots of the RISE for the 10 subjects across values of n. As n increases, both RISE and RISE variance tends to decrease, indicating that lNso warping estimation converges towards the true intensity warping function defining the scanner effect as the number of scans increases. We think this ⁴⁰ will be useful in the planning of future multisite studies, although we caution against over-interpreting these results to choose a priori the number of traveling

subjects necessary to account for scanner effects.

Figure 4: Boxplots of root integrated square error when different numbers of scans are used to calculate the warping function.