Transmit Field Bias Correction of T1w/T2w Myelin Maps

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Abstract

T1-weighted divided by T2-weighted (T1w/T2w) myelin maps were initially developed for neuroanatomical analyses such as identifying cortical areas, but they are increasingly used in statistical comparisons across individuals and groups with other variables of interest. Existing T1w/T2w myelin maps contain residual radiofrequency transmit field (B1+) biases, which may be correlated with these variables of interest, leading to potentially spurious results. Here we propose multiple methods for correcting these transmit field biases using either explicit measures of the transmit field or alternatively a 'pseudo-transmit' approach that is highly correlated with the transmit field. We find that the resulting corrected T1w/T2w myelin maps are both better neuroanatomical measures (e.g., for use in cross-species comparisons), and more appropriate for statistical comparisons across individuals and groups (e.g., sex, age, or body-mass-index). We recommend that investigators who use the T1w/T2w approach for mapping cortical myelin use these B1+ transmit field corrected myelin maps going forward.

Introduction

The T1-weighted divided by T2-weighted (T1w/T2w) myelin mapping approach was originally developed over a decade ago as a high-resolution non-invasive measure of cortical architecture that correlates with postmortem measures of cortical myelin content for the purpose of mapping cortical areas (Glasser and Van Essen 2011). The approach has proven highly successful at this purpose, revealing many cortical areal boundaries that co-localize with those of other non-invasive measures of cortical properties, such as cortical thickness, resting state functional connectivity, task activation, and within-area topographic maps measured with functional connectivity (Glasser et al., 2016a). Initial efforts were focused at the group level, given artifacts from imperfect surface reconstructions in individuals from relatively low resolution 1mm isotropic T1w and T2w scans (Glasser and Van Essen 2011). Markedly improved individual participant T1w/T2w myelin maps (Glasser et al., 2013; 2014) were later achieved by improving spatial resolution to 0.7mm or 0.8mm isotropic, less than half of the

minimum 1.6mm cortical thickness (Glasser et al., 2016b), and improving FreeSurfer (Fischl 2012; Zaretskaya et al., 2018) surface reconstructions by using the full resolution of the T1w images for white and pial surface placement and using the T2w image for exclusion of dura and blood vessels during pial surface placement. These individual myelin maps, collected and processed by the Human Connectome Project (HCP), were of sufficient quality to enable individual participant identification of cortical areas using a machine learning algorithm when combined with the other multi-modal non-invasive measures mentioned above (Glasser et al., 2016a). Importantly, the methodological improvements in T1w/T2w myelin maps in the context of the HCP Pipelines have to date focused solely on ensuring that they are appropriate for the above neuroanatomical use cases.

In parallel, there has been increasing interest in using non-invasive myelin maps, including T1w/T2w myelin maps, to make comparisons across individuals and groups, for example in healthy adults (Teraguchi et al., 2014; Shafee et al., 2015; Yang et al., 2020), over the course of development (Bozek et al., 2018; Norbom et al., 2020; Kwon et al., 2020), aging (Grydeland et al., 2013; 2019; Vidal-Piñeiro et al, 2016), in brain diseases (Teraguchi et al., 2014; Granberg et al., 2017; Rowley et al 2018; Nakamura et al., 2017; Du et al., 2019; Wei et al., 2020; Qiu et al., 2021), and for exploration of other neurobiological questions (Grydeland et al., 2016; Ma and Zhang 2017; Burt et al., 2018; Fukutomi et al., 2018; Li et al., 2019; Toschi et al., 2019; Gao et al., 2020; Liu et al.m 2020; Paquola et al., 2020); however, additional considerations arise when using T1w/T2w myelin maps to address such questions. For example, although in the absence of head motion there is an exact correction of radiofrequency (RF) receive field (B1-) effects after taking the ratio of T1w and T2w images, this ratio produces only a partial correction of RF transmit (B1+) field effects (Glasser and Van Essen 2011; see Theory section below). The residual B1+ effects are particularly noticeable in the HCP Young Adult (HCP-YA) dataset because of the smaller 56cm body transmit coil that was necessary to accommodate the stronger HCP-SC72 gradients (100 mT/m) of the customized Siemens 3T Skyra 'ConnectomS' platform (Uğurbil et al., 2013). Additionally, these design compromises resulted in the participants' heads lying 5cm above the magnetic field isocenter and also off-center, relative to the body transmit coil, resulting in reduced B1+ uniformity through the head. These residual B1+ effects on the T1w/T2w myelin maps are most easily appreciated as cross-hemisphere asymmetries in the cortical T1w/T2w ratio (Figure 5). Because such asymmetries are not expected neurobiologically (see below) and were smaller in the original Siemens 3T Trio datasets (Conte69 from Glasser and Van Essen 2011), an ad hoc correction method was initially developed to minimize them. This approach entailed subtracting the heavily surface-smoothed difference between a left/right symmetric group average Siemens 3T Trio T1w/T2w myelin map and each individual's T1w/T2w myelin map, resulting in the "bias-corrected (BC)" MyelinMap (MyelinMap_BC) results produced by the HCP Pipelines and released by the HCP (Glasser et al., 2013; Figure 18).

Although this "MyelinMap_BC" approach works well for localizing cortical areas in individual participants (Glasser et al., 2016a), it was never intended for cross-participant statistical analyses of the sort mentioned above, as it removes both artifactual and real cross-participant T1w/T2w myelin map differences. Further, although the Siemens 3T Trio-based maps have

reduced residual B1+ effects relative to the customized HCP-YA scanner, these residual transmit effects are not zero, meaning that all prior T1w/T2w myelin maps contain at least some biases, as biases along the anterior posterior and superior inferior axes cannot be removed by symmetrization (such as a symmetric center versus periphery bias). Finally, the actual B1+ transmit field is modulated by the loading of the body coil by both the participant's head and body, meaning that geometric variables such as head size and body size, and compositional variables such as weight and Body Mass Index (BMI), will have spurious correlations with uncorrected T1w/T2w myelin maps across participants. This modulation occurs because the scanner transmit voltage must be set higher for larger heads and bodies, leading to different B1+ transmit fields in participants of different sizes (see table 1). Further, asymmetries in the circularly polarized B1+ transmit field are induced by the upper bodies of participants and increase when they are larger (Glover et al., 1985; Sled and Pike 1998; Ibrahim et al., 2001; cf Figures 9 for children with smaller bodies on average and 12 for adults with larger ones obtained using the same sequences).

Because the HCP consortium was aware of these limitations, an Actual Flip angle Imaging (AFI) scan (Yarnykh 2007), which measures the effect of the non-uniform B1+ transmit field on the flip angle obtained in every voxel, was acquired in most participants during the same imaging session as the T1w and T2w structural images. However, use of these AFI scans to improve the myelin maps has remained unexplored until now. Additionally, subsequent HCP projects including the Human Connectome Development project (HCD), Human Connectome Aging project (HCA), and Connectomes Related to Human Diseases projects (CRHDs) required shorter protocols than the HCP-YA project, necessitating removal of the AFI scans and creation of an alternative approach to B1+ transmit field mapping based on the images that were acquired. Finally, we apply the B1+ correction of T1w/T2w myelin maps to a non-human primate study, the Non-Human Primate Neuroimaging and Neuroanatomy Project (NHP_NNP), as T1w/T2w myelin maps are useful in cross-species comparisons (Glasser et al., 2014; Mars et al., 2018; Autio et al., 2020; Hayashi et al., 2021).

Here, we demonstrate a novel set of B1+ transmit field correction methods of T1w/T2w myelin maps using actual and "pseudo" measures of the B1+ transmit field. We fit corrections of the T1w/T2w myelin maps at the group and individual levels, relying on (1) the hemispheric asymmetry of B1+ transmit field effects (at least on Siemens scanners in our experience to date), and (2) the neuroanatomically grounded assumption that asymmetries in cortical myelin content should not be correlated with asymmetries in the B1+ transmit field. We show that this correction removes not only the hemispheric asymmetries, but also additional symmetric biases. Finally, we show that this correction reduces B1+ correlated hemispheric asymmetries and spurious results in statistical comparisons of basic biological parameters of interest such as age, sex, or BMI.

Methods

Theory

The T1w/T2w myelin mapping approach is based on the idea that these images represent a combination of contrast correlated with myelin content and a multiplicative bias field as illustrated in Equation #1. In a given voxel, x represents contrast for myelin and b represents the image intensity bias field (Glasser and Van Essen 2011).

$$T1w/T2w \approx (x * b) / ((1/x) * b) \approx x^2 \quad (1)$$

Ideally, the multiplicative bias field inherent in MRI images would cancel, leaving a bias free ratio image with increased contrast of interest. This approach has an advantage over algorithmic bias corrections in that it makes no assumptions about tissue intensity uniformity. There is a large literature exploring non-uniformity of the cortical grey matter—reviewed in Glasser et al., 2014—and the white matter also contains genuine non-uniformities. Importantly, these grey and white matter tissue non-uniformities vary independently, meaning that an algorithmic correction attempting to achieve white matter uniformity will induce a bias in grey matter and vice versa. Indeed, it was this observation that led to development of the T1w/T2w approach originally (Glasser and Van Essen 2011). Equation #1 can be expanded to consider the effects of the receive (rb) and transmit (tb) bias fields separately as in Equation #2.

$$T1w/T2w \approx (x * rb * tb_{T1w}) / ((1/x) * rb * tb_{T2w}) \quad (2)$$

This simplifies to Equation #3, as the receive bias field cancels out exactly in the absence of motion and the myelin contrast is enhanced.

$$T1w/T2w \approx x^2 * (tb_{T1w} / tb_{T2w}) \quad (3)$$

Unfortunately, tb_{T1w} / tb_{T2w} does not equal 1 in any location where the scanner does not achieve the prescribed flip angle. The T1w/T2w ratio approach does reduce the residual intensity bias seen, even if the receive bias has already been corrected on the scanner (e.g., with Siemens PreScan Normalize), as was seen in the original T1w/T2w myelin mapping dataset (Glasser and Van Essen 2011). The reason for the difference in the transmit field effects on the T1w and T2w images is that the spin echo-based T2w SPACE acquisition employs a substantially different RF pulse train (e.g. 90 degree excitation and multiple variable flip angle refocusing pulses up to 180 degrees; Mugler 2014) that requires more transmit power than does the gradient echo-based T1w MPRAGE image pulse sequence (e.g. ~8 degree excitation pulses). As such, the signal intensity of the T2w SPACE image is more affected by transmit field inhomogeneities than is the signal intensity of the T1w scan, which has the added benefit of using an adiabatic inversion pulse for the initial 180-degree preparation pulse that is designed to be resilient to transmit field inhomogeneities (Garwood and Ugurbil 1992). Notably, the effect of tb_{T1w} / tb_{T2w} on T1w/T2w varies spatially according to the transmit field and is multiplicative (Bonny et al., 1998; Collewet et al., 2002; Wang et al., 2004; Wang et al., 2005; Weiskopf et al 2011; Delgado et al., 2020) but does not substantially affect image contrast over the range of flip angles at 3T (Mugler 2014). Importantly, this effect is expected to vary across participants based on interindividual differences such as weight, Body Mass Index (BMI), or head size due to differential transmit coil loading and dielectric effects. These variables may

themselves correlate with variables of interest such as sex or age, or even be intrinsically of interest themselves (such as BMI). Thus, if we have a measure of the transmit field (TF), we can use it to attenuate these effects in the T1w/T2w data by determining the slope and intercept that relate the TF to the residual bias in the T1w/T2w ratio with a linear approximation, as in Equation #4 where tb_{T1w} / tb_{T2w} = 1/(TF*slope+intercept).

$$(T1w/T2w)_{corr} = (T1w/T2w)_{orig} /(TF*slope+intercept) \quad (4)$$

For simplicity, the TF is already scaled such that a value of 1 means that the reference flip angle was reached exactly (in the case of the AFI approach in HCP-YA data, the reference flip angle was set to 50 degrees, and to compute TF we divide the flip angle map by 50). If we assume that the T1w/T2w values should not be changed where the achieved flip angle equals the reference value, slope is the only parameter to optimize, as in Equation #5.

$$(T1w/T2w)_{corr} = (T1w/T2w)_{orig} /(TF*slope+(1-slope)) \quad (5)$$

We investigated two different ways for optimizing the slope value, by either using asymmetry (Equation #6) or correspondence with a previously bias-corrected template (Equation #7). Comparison to a template is more robust because it does not require significant asymmetry in the B1+ maps, making it a better option for individual-level correction. Instead, it relies on the strong central to peripheral gradient in the transmit field illustrated in Figure 5, which is more consistent in individual participants than the L-R asymmetry. However, to use this approach, we first must generate a corrected group template (or find the value equivalent to the reference flip angle in the pseudo-transmit approach described below) using the asymmetry method (Equation #6), as there is no existing unbiased template to evaluate against. We make the neurobiologically plausible assumptions that the actual cortical myelin distribution is generally symmetric across the left and right hemispheres and thus that hemispheric asymmetries present in both the group T1w/T2w ratio and the group B1+ transmit field are spurious. This approach only works if the B1+ transmit field is asymmetric, which it is in the 3T Siemens Trio, HCP 'ConnectomS' (Figure 4), and Prisma scanners. Indeed, even at 3T with a relatively uniform circularly polarized body coil as the transmitter, such an asymmetry exists due to the fundamental dependence of the B1+ field pattern on the electrical properties (i.e., permittivity and conductivity) and geometry (e.g., size) of the sample (Glover et al., 1985; Sled and Pike 1998; Ibrahim et al., 2001). Thus, we used the cost function in Equation #6 to find the optimum slope to produce a corrected group T1w/T2w myelin map, where L is the left hemisphere and R is for the right hemisphere.

$$Cost=sum(abs((L-R)/((L+R)/2))) \quad (6)$$

We chose to optimize this cost function using the golden search algorithm (Kiefer 1953; aka golden section search). This optimization produces a group average transmit field corrected T1w/T2w myelin map (see Figure 6). Using this corrected T1w/T2w group surface map as a

reference template (T), one can correct individual participants (I) using a more robust cost function in Equation #7.

$$Cost=sum(abs((I-T)/(T)) \quad (7)$$

Importantly, when applying Equation #7 we want to allow for the existence of real per-participant differences in the spatial mean of the T1w/T2w ratio. To prevent these per-participant differences from biasing the cost function in Equation #7, we do the following: (1) identify the surface vertices where the calculated actual flip angle is close to the target flip angle (+/- 5%, where the B1+ correction will not cause appreciable changes to the values at those vertices); (2) find the median T1w/T2w ratio within those vertices for both the group and individual T1w/T2w maps; and (3) divide the ratio of the medians out of the individual map before computing the cost function. This process avoids over- or under-correcting the spatial pattern because of per-participant differences in the spatial mean of the T1w/T2w ratio (see also the section on *Covariates for Downstream Statistical Analyses Involving Groups of Individuals* for more details on controlling for nuisance effects on the T1w/T2w ratio mean).

We demonstrate below that this approach indeed works if an AFI map is available, as is the case for the HCP-YA data (though unfortunately due to acquisition artifacts, the map is imperfect at the individual participant level as described below and shown in Figure 4). However, in some HCP-style studies no AFI map is available, so an alternative approach is needed. A candidate alternative emerged in the course of developing an improved receive field bias correction approach for fMRI data termed the "SEBASED" correction (Glasser et al., 2016a), which enables biological interpretation of fMRI parameter maps (e.g., so that estimated task fMRI betas have a consistent scale across space). In particular, differential transmit field effects were also present in the gradient echo EPI (echo-planar imaging) fMRI images and the spin echo EPI field map images (for the same reason that they are present in the T1w and T2w images as described above, i.e. the signal intensity depends on sin(FA*TF) for gradient echo images and sin^3(FA*TF) for spin echo images, FA=Flip angle; TF=Transmit Field; Bonny et al., 1998; Collewet et al., 2002; Wang et al., 2004; Wang et al., 2005; Weiskopf et al 2011; Delgado et al., 2020). For the purposes of fMRI receive bias correction, such transmit effects were nuisances that needed to be appropriately modeled (together with T2* induced susceptibility dropouts) to generate an accurate receive field estimate. However, this "pseudo" transmit field can also be used for T1w/T2w myelin map correction if it is collected during the same session as the structural data. Indeed, at the group average level, this pseudo-transmit field has a remarkably similar spatial pattern to the AFI-based transmit field (compare Figures 5 versus 9 and 12). Thus, the pseudo-transmit field can be used in the same way as a real transmit field if a reference value can be established, analogous to the reference flip angle of 50 degrees in the AFI transmit field (see the pseudo-transmit specific methods section below).

The above theoretical considerations entail several assumptions. 1) *No significant change in head position between the T1w and T2w images*. For the receive field to exactly cancel, it must be identical in both images. Because the head coil is stationary during a scan, head motion changes where the brain anatomy is inside the receive bias field. The use of on-scanner

corrections such as Siemens PreScan Normalize will mitigate this issue, as a fixed receive field is removed from the images by the scanner, and thus prior to downstream motion correction (which will align the anatomy but not the receive fields, given the differential relationship between brain anatomy and B1- receive field when head motion occurs). Thus, if on-scanner B1- corrections are not used, it is helpful to separately acquire data to measure the B1- receive field and apply this correction to the data in a way that accounts for the head movement within a static bias field (see 'BIAS' scans in *AFI Image Preprocessing)*. This assumption is addressed in the methods section below and also applies to the use of GRE and SE EPI images for pseudo-transmit field computation. 2) *No significant change in head position between the B1+ transmit field map and T1w and T2w images*. Changes in head position may intrinsically change B1+ inhomogeneity due to differential coil loading and reduce the accuracy of B1+ field correction. This assumption is difficult to address, but it is expected to be a small effect at 3T given most 3T scanners use a large body transmitter coil. This is also not an issue for non-human primate imaging because the head is almost always fixed during imaging. 3) *The B1+ transmit field can be related to the residual B1+ bias with a linear approximation*. The typical non-linear effect of the B1+ field on spin echo-based T2w images is conditioned by an optimized (vendor dependent) variable flip angle echo train in the SPACE sequence to yield a small (and approximately linear) dependence of signal intensity and contrast on flip angle variation (i.e., ~ +/-20%) typically observed in the human brain at 3T (Mugler 2014). This assumption is also empirically supported by the results in the manuscript. 4) *The pseudo-transmit field reference value correctly indicates where the scanner achieves the intended flip angle.* Individual differences in T2 and T2* may influence the pseudo-transmit field map beyond the real transmit field. T2* from large scale magnetic field inhomogeneity is dealt with using the thresholding described below, but it is clear that the reference value found by optimizing the group-level left/right cost function is slightly different in the HCD and HCA datasets due to such effects. Similarly, the scanner is assumed not to differentially scale the GRE and SE images automatically in different subjects (and such behavior has not been found on Siemens scanners).

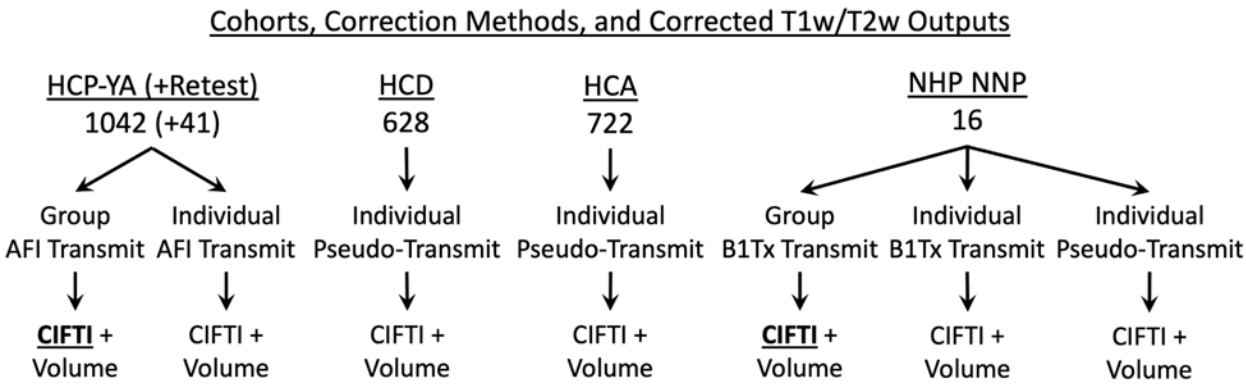

Figure 1 illustrates the datasets, specific analyses, and corrected outputs in this study. **Bold underlined** font indicates the reference T1w/T2w outputs (which are computed only for the surface). Surface data are stored in the CIFTI format (Glasser et al., 2013), which combines both hemispheres into a single file.

Participants

Data from the Human Connectome Young Adult (HCP-YA), Development (HCD), and Aging (HCA) Projects were acquired and used with approval of the Washington University Institutional Review Board (IRB) and those of partner institutions (Van Essen et al., 2013, Bookheimer et al., 2019; Harms et al., 2018; Somerville et al., 2018). The Non-Human Primate Neuroimaging and Neuroanatomy Project (NHP_NNP) data were acquired with approval from the RIKEN Kobe Japan institutional animal care and use committee (Autio et al., 2020; Hayashi et al., 2021). Figure 1 illustrates the cohorts, specific analyses, and corrected T1w/T2w outputs produced in this study. For HCP-YA, 1042 participants had successful MSMAll registration and had AFI scans available (ages 22-35, 470 men and 572 women, BMI 16.5-47.8, head size 2433cc to 4461cc, and body coil reference voltage 183V-356V). For HCD, 628 participants had successfully completed preprocessing and quality control at the time of the study and were publicly released as part of the "Lifespan HCP Release 2.0" (ages 8-22 (5-7 year olds were not included due to small numbers at this time), 292 boys and 336 girls, BMI 13.1-43.2, head size 2133cc to 4260cc, and voltage 206V to 317V). For HCA, 722 participants had successfully completed preprocessing and quality control at the time of the study and were publicly released as a part of the "Lifespan HCP Release 2.0" (ages 36-90+, 319 men and 403 women, BMI 15.7-43.7, head size 2522cc to 4719cc, and voltage 217V-337V). The correlations of these demographic variables are listed in Table 1. The NHP_NNP acquired data from 16 macaque monkeys (13 *M. mulatta* and 3 *M. fasicularis*, 16 male and 0 female, ages 3.5-7.0, weights 3.5-9.8kg, head size 342cc to 570cc, and voltages 266V-282V).

Table 1	Sex			Age			BMI			HeadSize			Voltage		
Study	HCP-YA	HCD	HCA	HCP-YA	HCD	HCA	HCP-YA	HCD	HCA	HCP-YA	HCD	HCA	HCP-YA	HCD	HCA
Sex	1.00	1.00	1.00	-0.22	0.04	0.02	0.06	-0.04	0.03	0.73	0.45	0.67	0.55	0.26	0.39
Age	-0.22	0.04	0.02	1.00	1.00	1.00	0.07	0.50	-0.14	-0.12	0.57	0.00	-0.06	0.48	-0.08
BMI	0.06	-0.04	0.03	0.07	0.50	-0.14	1.00	1.00	1.00	0.43	0.61	0.43	0.70	0.79	0.72
HeadSize	0.73	0.45	0.67	-0.12	0.57	0.00	0.43	0.61	0.43	1.00	1.00	1.00	0.78	0.75	0.69
Voltage	0.55	0.26	0.39	-0.06	0.48	-0.08	0.70	0.79	0.72	0.78	0.75	0.69	1.00	1.00	1.00

Table 1 contains the demographic Pearson correlations of the three human studies, HCP-YA (n=1042), HCD (n=628), and HCA (n=722).

Scans HCP-YA

The acquisition of the HCP-YA data has been described previously (Glasser et al., 2013; Uğurbil et al., 2013). Briefly, 0.7mm isotropic resolution T1w MPRAGE images (Mugler and Brookeman 1990) were acquired with TI=1000ms, flip angle=8 degrees, TE=2.14ms, and TR=2400ms. 0.7mm isotropic resolution T2w SPACE images (Mugler et al., 2000) were acquired with TE=565ms and TR=3200ms. One or two T1w and T2w images were used depending upon image quality. A 4.5 min AFI scan (Yarnykh 2007) was acquired at 2mm isotropic resolution, TR1=20ms, TR2=120ms, 50-degree nominal (i.e. reference) flip angle, and TE=1.9ms. Additionally, 2mm isotropic "Bias_32CH" and "Bias_32BC" images were acquired with a gradient echo sequence (TR=250ms, TE=1.01ms) with the only difference being whether the 32-channel head coil (32CH) or body coil (BC) was used for image receive (each ~ 0.5 min). These images were acquired to measure the B1- receive field, as it was desired in the HCP-YA to do all

image processing offline and Siemens PreScan Normalize (PSN) was not used for any acquisitions (we recommend the use of Siemens PreScan normalize for future studies for all acquired images including structural, functional, diffusion, and spin echo field maps). The images were obtained on a customized Siemens 3T 'ConnectomS' scanner with 100mT/m gradients using a Siemens standard 32 channel head coil. All acquisitions used the body coil for radiofrequency transmission. All images had the face and ears removed using a defacing algorithm (Milchenko and Marcus 2013) to prevent identification of participants. Forty-one Test-Retest participants were scanned and processed a second time through all modalities (interval: 18-328 days; mean: 135 days). In a separate scanning session, gradient echo EPI fMRI data were acquired at 2mm isotropic resolution along LR and RL phase encoding directions with matching phase reversed spin echo fMRI data (Smith et al., 2013), data which were used for MSMAll areal feature-based surface alignment (see below).

Scans HCA and HCD (HCP-Lifespan)

The acquisition of the HCD and HCA data has been described previously (Harms et al., 2018). Briefly, 0.8mm isotropic resolution T1w multi-echo MPRAGE images (van der Kouwe et al., 2008) were acquired with an internal EPI-based navigator (Tisdall et al., 2012; vNav) with TI=1000ms, flip angle=8 degrees, TE=1.8/3.6/5.4/7.2ms, and TR=2500ms. Only the average of the first two echo times were used because of artifacts that were subsequently discovered in the later echo times (a multi-echo MPRAGE was used in an effort to equalize receiver bandwidth between the T1w and T2w images; however, such an approach is not required as readout distortions can be corrected in T1w and T2w images (Glasser et al., 2013) and a short TE single echo MPRAGE as used in the HCP-YA has higher SNR and fewer artifacts). 0.8mm isotropic resolution T2w SPACE images (Mugler et al., 2000) were acquired with an internal EPI-based navigator (Tisdall et al., 2012; vNav) with TE=564ms, TR=3200ms. Spin echo (SE) EPI images with both anterior-to-posterior (AP) and posterior-to-anterior (PA) phase encoding polarities were acquired at 2mm isotropic resolution. Gradient echo (GRE) EPI-BOLD fMRI images were acquired with identical geometric specifications, again with both AP and PA polarity. These images were obtained on a standard Siemens 3T Prisma with 80mT/m gradients using a Siemens standard 32 channel head coil. For the T1w and T2w acquisitions, both PSN and unfiltered (non-normalized) reconstructions were generated. The unfiltered versions were used in initial HCP Pipelines preprocessing to match the approach used in HCP-YA. Defacing (Milchenko and Marcus 2013) was run on the 3D T1w and T2w images including separately on the PreScan normalized and unfiltered images.

Scans NHP_NNP

The acquisition of the NHP_NNP data has been described previously (Autio et al., 2020). Briefly, 0.5mm isotropic resolution T1w MPRAGE images were acquired with TI=900ms, flip angle=8 degrees, TE=2.2ms, and TR=2200ms. 0.5mm isotropic T2w SPACE images were acquired with TE=562ms and TR=3200ms. B1+ mapping ("B1Tx") was acquired using a vendor-provided slice-selective RF-prepared sequence (called 'B1map' by Siemens) with a turbo fast low-angle-shot (FLASH) read-out with 2.0mm isotropic resolution, flip angle=8 degrees, nominal target flip

angle=80 degrees, TE=2.5ms, and TR=20s (Chung et al., 2010). Because the preparation RF pulse excites slightly thicker slices than the FLASH acquisition (requiring a gap between the slices), even and odd slices (with a gap 2 mm) were acquired in separate runs for full brain coverage (each 40s in duration). A pair of reversed phase-encoding (left-to-right [LR] and right-to-left [RL]) spin echo EPI images were acquired at 1.25mm isotropic resolution together with reversed phase-encoding gradient echo EPI-BOLD fMRI images with identical geometric specifications. These images were obtained on a standard Siemens 3T Prisma with 80mT/m gradients using a customized macaque-sized 24 channel head coil (Autio et al., 2020) that enables HCP-style pulse sequences to be acquired in the macaque (commercially available from Rogue Research Inc., Montreal/Takashima Seisakusho Co., Tokyo). PSN was used on all images (as recommended for all human and non-human primate studies going forward). The macaques were anesthetized using dexmedetomidine and low-dose isoflurane in a protocol described previously (Autio et al., 2020).

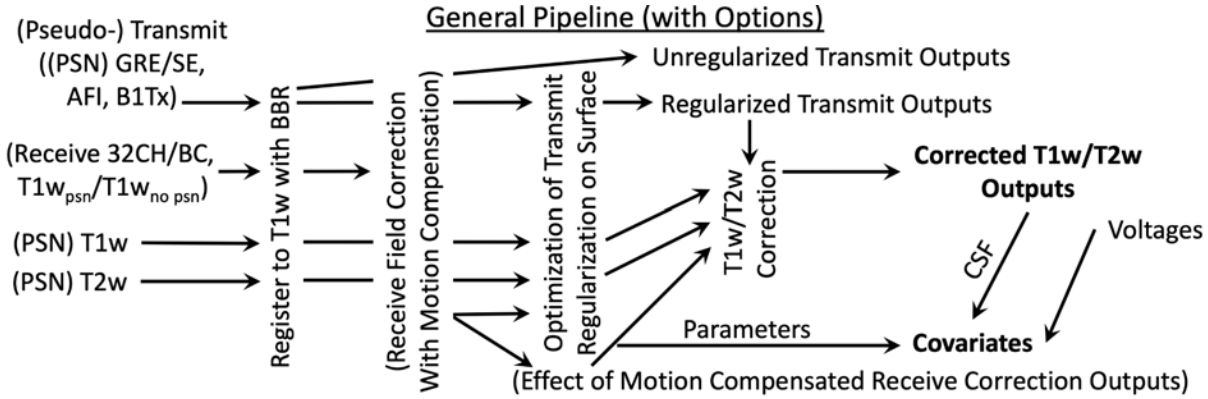

Figure 2 illustrates the overall T1w/T2w transmit field correction pipeline operations and outputs, with inputs and outputs oriented horizontally and processing steps oriented vertically. Options (depending on specifics of data acquisition) are indicated with () and arrows passing beneath a step. The simplest form of the pipeline would require that the T1w and T2w images (and GRE/SE images if the pseudo-transmit approach is used) have undergone Siemens PreScan Normalize (PSN) or the equivalent from another vendor, which permits skipping the Receive Field Correction With Motion Compensation step. **Bold** text indicates the primary outputs of interest for downstream analyses. Other outputs are helpful for investigating the effects of each correction and group analyses. In all cases, outputs are in CIFTI, NIFTI physical volume, and NIFTI MNI volume space.

General Preprocessing

All datasets were preprocessed using the HCP Pipelines (Glasser et al., 2013) or HCP-NHP Pipelines (Autio et al., 2020), which included the structural spatial preprocessing pipelines, the functional preprocessing pipelines, the multi-run spatial ICA+FIX (Salimi-Khorshidi et al 2014; Glasser et al., 2018) fMRI data denoising pipelines, and, in humans, multi-modal 'MSMAll' areal-feature-based surface registration (Robinson et al., 2014; 2018; the utility of MSMAll has yet to be established in the less variable macaque brain so cortical-folding-based 'MSMSulc' registration was used in macaques). The relevant results of these pipelines for the purposes of

this study are MSMAll (or MSMSulc in the case of macaques) aligned cortical T1w/T2w myelin maps and also T1w/T2w ratio volumes in MNI space (aligned using nonlinear FSL FNIRT-based registration of the T1w image). Figure 2 illustrates the general T1w/T2w transmit correction pipeline operations and outputs with multiple optional steps indicated. This pipeline is described in detail over the following sections. The group transmit (for generating reference T1w/T2w myelin maps) and individual transmit and pseudo-transmit nested optimization algorithms are illustrated in Figure 3 and described below.

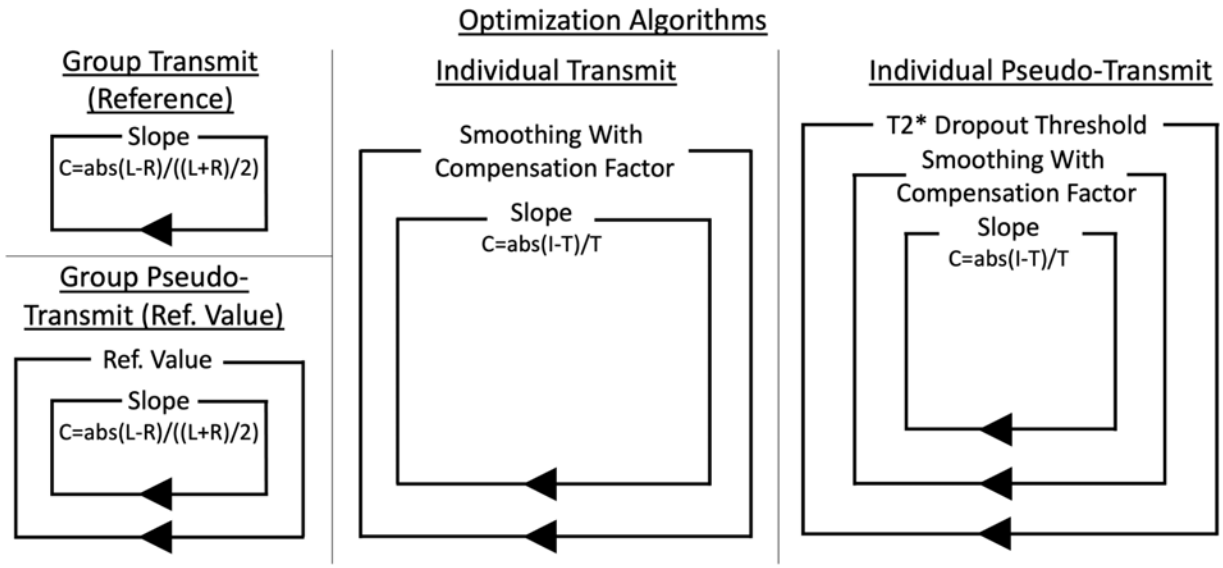

Figure 3 illustrates the four optimization approaches used in this study. Each box represents a loop, and anything the box surrounds is inside that loop body. The simplest is the group transmit optimization approach, which uses a single loop with the cost function in Equation #6 and generates species specific reference T1w/T2w myelin maps (the group average B1+ transmit fields are assumed not to require regularization due to cross-participant averaging for removal of artifacts). The reference map is then used in the individual transmit nested optimization algorithm together with the cost function in Equation #7. The outer loop finds the optimal amount of (spatially constrained) smoothing (for regularization) while compensating for the change in fraction of flip angles above and below the reference flip angle and the inner loop finds the optimal slope. For the pseudo transmit approach, the appropriate reference value (where the target flip angle is expected to have been achieved by the scanner) must be found for a given study, which is again done at the group level using Equation #6. Then the individual pseudo-transmit nested optimization algorithm is used together with the cost function in Equation #7, with the addition relative to the individual transmit algorithm of a T2* dropout threshold optimization loop. The reference map from the group transmit algorithm is used in both transmit and pseudo-transmit approaches. Appendix A contains a pseudocode representation of the individual pseudo-transmit algorithm.

AFI Image Preprocessing

The raw AFI volumes consist of two image volumes acquired at two different TRs. These need to be corrected for gradient nonlinearity distortions (as the other images have been) and rigidly aligned to the T1w scans in the participant's physical space. FreeSurfer boundary-based registration (BBR) (Greve and Fischl 2009) is used for this alignment after initialization with 6 DOF FSL FLIRT registration. At this point, the raw AFI volumes are aligned to the participant's physical space (and can be trivially resampled to MNI space or mapped onto the cortical surfaces). Unfortunately, the second AFI volume invariably contains ringing artifacts, which induce artifacts in the computed flip angle maps (Figure 4; Yarnykh 2007). These artifacts are most likely due to inadequate spoiling for the second volume (Nehrke 2009), but were of variable severity across the HCP-YA participants. Unfortunately, there is no easy way to eliminate these artifacts in individuals (averaging across participants eliminates them at the group level). Thus, we use the property that the B1+ transmit field map is mostly smooth in 3D to reduce the artifacts using spatial smoothing. Importantly, such relatively unconstrained 3D spatial smoothing is appropriate in this context *ONLY* because the B1+ transmit field is relatively spatially smooth in 3D in the volume at 3T. Because most brain neuroanatomy is not smooth in 3D, the traditional uses of unconstrained 3D smoothing in neuroimaging are almost exclusively inappropriate (Glasser et al., 2016b; Coalson et al., 2018).

Moreover, it turns out that totally unconstrained 3D spatial smoothing is also inappropriate for the B1+ transmit field. Figure 4 illustrates sharp discontinuities in the 3D structure of the B1+ transmit field at the fibrous falx and the tentorium in group average data without smoothing. This interesting property of the B1+ transmit field is presumably related to the electrical properties of different brain tissues (Vaidya et al., 2016), but has not, to our knowledge, been previously shown at this level of anatomical detail and has important implications for transmit field map regularization. When we initially used uniform smoothing in 3D, we found good correction of lateral cortex but residual bias medially, indicating that the fibrous dural tissue's effect on B1+ is present (though not easy to see) in the T1w/T2w images themselves. Better results were obtained by smoothing within three distinct regions (by a common amount for all three regions) generated from FreeSurfer's individual participant segmentations -- the left and right cerebral hemispheres separated by the falx, and the combination of the cerebellum and brainstem separated by the tentorium. We did allow smoothing across the corpus callosum and the midbrain, as they lack a fibrous discontinuity (anatomically and in the transmit field). Because valid flip angle data is present outside the grey and white matter (Figure 4), we also allowed smoothing to occur within the rest of the entire head to reduce edge effects. We smoothed in the participant's physical space because that was where the B1+ was generated. Because the artifacts vary in severity on a per participant basis (Figure 4), we used a golden search approach (similar to what was used above for finding the slope with Equation #7) to optimize the amount of smoothing in each participant (see Figure 3). Finally, because heavy smoothing changes the average location of the reference flip angle, we compensated the smoothed maps to ensure that a similar number of voxels are above and below the reference flip angle to avoid introducing biases in the T1w/T2w myelin map mean from the smoothing. At this point the final flip angle map in volume or surface space is ready for correction of the T1w/T2w myelin map using Equation #5.

The HCP-YA data have an additional consideration (relative to the above Theory section): they were not acquired with PreScan Normalize. Instead, two short gradient echo scans were acquired for estimating the receive bias field. These two scans ("BIAS_32CH" and "BIAS_BC") differed only in the receiver coil used (32-channel head coil vs body coil). Such data were not used when running the HCP Pipelines originally and thus the original T1w/T2w myelin maps have biases in them related to participant motion between the T1w and T2w scans. Thus, we used these scans to generate a smoothed (8mm FWHM) receive field restricted to the brain (due to poor estimation in osseous structures and differential defacing effects in the two images) and then dilated with extrapolation to estimate the field outside the brain to enable correction even in the setting of bulk head motion between the receive field scans and the T1w and T2w scans. To align this data to the T1w image, we used FreeSurfer BBR registration (Greve and Fischl 2009) using the pial surface instead of the default white surface given that grey matter/CSF contrast exceeded grey/white contrast in these images. This field was then applied so as to match the estimated participant movement between the T1w and T2w images (averaging the receive fields if two T1w or T2w images were used prior to dividing T1w by T2w) to produce a final field that represented the error from motion on the original T1w/T2w receive field correction. The error field was then used to correct the T1w/T2w ratio data prior to fitting the AFI transmit field map and final T1w/T2w correction.

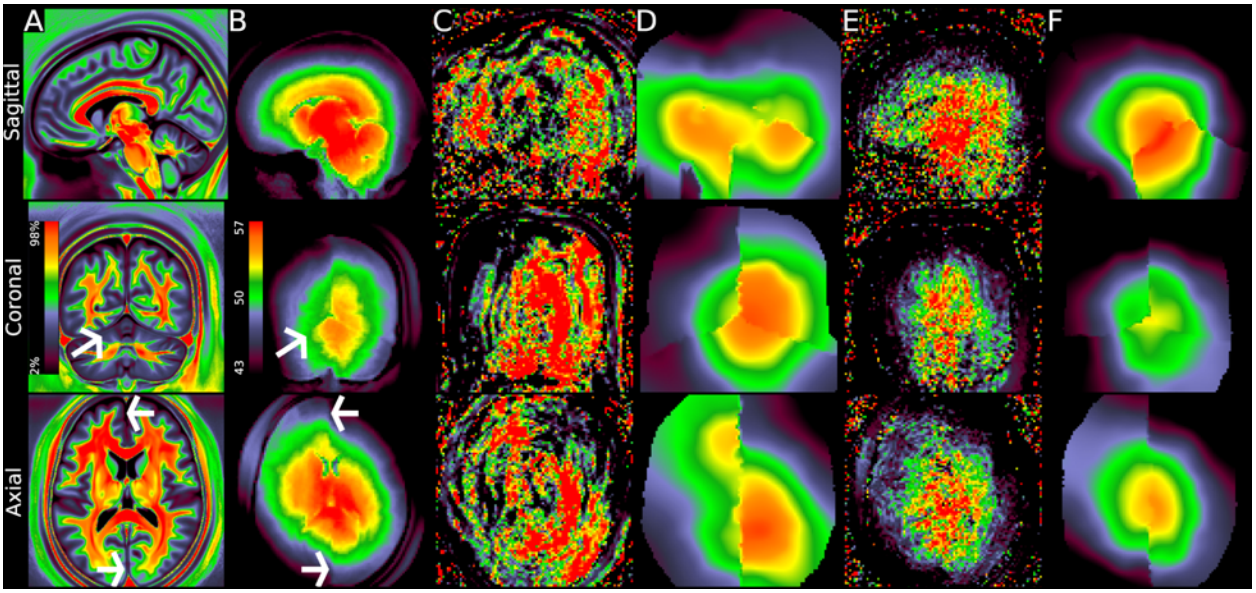

Figure 4 illustrates the group average original T1w/T2w volume and the unsmoothed group average AFI map together with unregularized and regularized AFI maps from two exemplar individuals. Panel A shows the group average original T1w/T2w average volume in sagittal (top), coronal (middle) and axial (bottom) slices. Panel B illustrates the group average AFI map, where the original AFI volumes (i.e., TR1 and TR2) are aligned to MNI space nonlinearly and then averaged across participants without smoothing; the AFI flip angle map is then computed from these results at the group level. The arrows mark the anterior and posterior falx and the left tentorium, which all show sharp discontinuities in the AFI map, indicating that the B1+ field is particularly affected by these fibrous dural reflections. Panels C and D illustrate a participant with the worst (worst correlation with the group average map) ringing artifacts before and after

regularization. Panel E and F illustrate a participant with the least (best correlation with the group average map) ringing artifacts before and after regularization. Other participants lie between these extremes. Additionally, Panel A shows a subtle correlation with the hemispherically asymmetric pattern in the B1+ map in Panel B within the deep white matter; however, contrast differences between tissues such as CSF/grey matter/white matter are much stronger than those within tissues and also the B1+ effect. This map also illustrates how partial volume effects between CSF/grey matter/white matter tissues overwhelm differences in myelin content within the cortical grey matter in group average T1w/T2w volume maps, making them inappropriate for cortical analyses.

Pseudo-transmit Field Generation and Preprocessing in Humans

As described above in the theory section, the idea behind using gradient echo and spin echo EPI images to compute a "transmit field"-like measure is that, just as the gradient echo T1w MPRAGE and spin echo T2w SPACE sequences experience differential amounts of transmit radiofrequency, gradient echo fMRI and spin echo field map images also experience different amounts of transmit field (i.e., the gradient echo image experiences an excitation flip angle of ~50 degrees but the spin echo image experiences and excitation flip angle of 90 degrees and then a 180 degree refocusing pulse). Unfortunately, they experience other effects that must also be considered, including magnetic inhomogeneity effects in the gradient echo image (i.e., signal dropout in regions of main magnetic field (b0) inhomogeneity from air-tissue interfaces, around blood vessels, and at tissue boundaries such as the brain and skull), potentially differential receive fields (if there is head motion between scans that did not all use PreScan Normalize), and slight differences in tissue contrast (due to T2* versus T2 weighting). Additionally, magnetic field inhomogeneity causes both types of images to suffer from signal pile up and rarefaction, and gradient echo images additionally have their intensities modulated by differential effective echo times across space (Deichmann et al., 2002; De Panfilis and Schwarzbauer 2005), requiring correction using a combination of Jacobian modulation and combining across opposed phase encoding directions. We used balanced amounts of data from opposed phase encoding directions to ensure that such contributions are appropriately compensated.

To produce the pseudo-transmit field, the average across phase encoding polarities of the single band reference gradient echo images ("SBRef;" after alignment and distortion correction) is divided by the corresponding average from the Jacobian modulated spin echo images. This division cancels the receive field and markedly reduces the tissue contrast in the image while preserving differential B1+ effects. For the pseudo-transmit approach, it is necessary to determine the "reference value" of the pseudo-transmit field that represents the location where the scanner achieves the prescribed flip angles (the locations equivalent to the 50-degree spots in the above transmit field maps). The reference value is found at the group level using the group average uncorrected myelin map and "raw" group average GRE/SE images, where first a reference value and then a slope are fit in a nested algorithm whose cost function minimizes the left to right asymmetry (Equation #6; see Figure 3). Because the raw group average GRE/SE images have significant T2* dropout regions, we did not produce a reference

map from this correction, and instead used the species-specific reference myelin map computed from the transmit field approach in the HCP-YA data or NHP data. Several remaining free parameters must be determined, which we accomplish in a nested fashion using the same golden search algorithm described above (i.e., the golden search for threshold (1) uses a cost function that runs an internal golden search on the smoothing (2), which in turn uses a cost function that internally runs a golden search on slope (3) as illustrated in Figure 3, and with pseudocode in Appendix A). *1) The threshold at which the GRE / SE-EPI pseudo-transmit field is contaminated with T2* dropout effects.* If the threshold is too small, holes in the field will appear in regions of susceptibility dropout, but if it is too large, the range of the field will be clipped for low flip angle regions. *2) The smoothing FWHM to reduce noise and tissue-specific effects* (using the same constrained 3D volumetric smoothing approach described above for the AFI scans with the same approach for compensating for the effects of smoothing on the mean of the pseudo-transmit field map). After thresholding but prior to smoothing, extrapolating dilation (using the local gradient to predict continued changes outside of the valid data region) is used to fill the head mask to avoid edge effects of smoothing at the edge of the valid data. This dilation is necessary because unlike the AFI approach above, valid data is not present outside of the brain because of the application of fat saturation to the gradient echo and spin echo EPI acquisitions. *3) The slope of the T1w/T2w pseudo-transmit field correction* (to use in Equation #5). The slope is fit using Equation #7 as the cost function. Because a group average reference T1w/T2w myelin map is available from the HCP-YA study, this template is not recomputed for the HCP-Lifespan data (but is for the macaques, see below).

Additionally, for the HCP-Lifespan data the T1w and T2w images were acquired with PreScan Normalize, but the unnormalized versions were used for preprocessing for historical reasons (which is not recommended going forward). Thus, the effect of any head motion between the T1w and T2w images (and GRE and SE images) must be compensated for, as described above. Explicit "BIAS" scans for estimating the receive field were not collected for HCP-Lifespan; however. Thus, for HCP-Lifespan, the receive field was computed by dividing the non-PreScan Normalized T1w image by the PreScan Normalized T1w image (the T2w image had additional on-scanner filters applied and could not be used in this way). Since these images entered the de-facing pipeline separately, de-facing differences were present due to the large discrepancies in image intensity and bias that affected the defacing pipeline registration algorithms (Milchenko and Marcus 2013). Thus, values outside of the brain for the receive field were unreliable and had to be inferred through extrapolating dilation as described above. Once computed, the effects of motion on this receive field can be modeled for the T1w and T2w images together with the GRE and SE images to ensure that the receive field properly cancels in all ratio images (T1w/T2w and GRE/SE).

Special Considerations for Macaque Transmit Field and pseudo-transmit Field Preprocessing

For the macaques, the data necessary for both real and pseudo-transmit field correction were available, so the two approaches could be directly compared. The transmit field consists of a "magnitude" image and a "phase" image. The magnitude image contains artifacts outside the brain that make registration initialization problematic. Because the macaques were

anesthetized with head fixed using adhesive medical tape, motion between the scans is expected to be minimal. Thus, registration was initialized using the existing FreeSurfer-based BBR registration between the scanner space fMRI and T1w images (under the assumption that the brain did not move much between the fMRI and B1Tx acquisitions) and then was fine-tuned with FreeSurfer BBR (Greve and Fischl 2009) to generate a precise registration between the B1Tx and T1w data. This approach avoids the need for failure-prone unmasked registration or arduous brain extraction of the B1Tx magnitude images for the initialization and likely could be used throughout the HCP and HCP-NHP Pipelines to reduce registration initialization failures. The reference flip angle from the Siemens B1+ transmit field phase map is 80 degrees (encoded in the image as flip angle * 10 or 800), so this value was used instead of 50 that was used in the HCP-YA AFI maps in humans. Similar to the other measures, the transmit field has some artifacts that are reduced by falx and tentorium constrained volumetric smoothing. Given the finer spatial resolution available and smaller brain sizes, distances for extrapolating dilation and outlier detection were reduced by half for the macaque relative to human scans. Because Siemens PreScan Normalize was used on all images, post-hoc receive field correction for motion was unnecessary and was skipped (Figure 2). As above for HCP-YA, a macaque T1w/T2w reference myelin map was generated using the unregularized group average transmit field and T1w/T2w myelin maps with the L-R cost function (Equation #6), as was the pseudo-transmit reference value, and then individual participants were corrected with both the transmit field and pseudo-transmit field approaches using the I-T cost function (Equation #7).

Covariates for Downstream Statistical Analyses Involving Groups of Individuals

The above methods aim to markedly reduce the effects of variable B1+ inhomogeneity, which correlate with interindividual differences such as BMI and head size and which may correlate with variables of interest in downstream statistical analyses (together with the effects of motion on B1- receive field correction across participants which introduce random artifacts in T1w/T2w myelin maps). That said, the T1w/T2w ratio is not an intrinsically quantitative measure such as, for example, a tissue T1 value. Because it has the potential to vary, we produce reference T1w/T2w values from the lateral ventricular CSF, which is not expected to vary with myelin content. Because there may be differences across ages in ventricle size and CSF flow artifacts, we chose the CSF T1w/T2w percentile value having the strongest correlation with the mean T1w/T2w value across participants (after eroding the lateral ventricles by 1 voxel to eliminate partial volume effects). Additional covariates we consider appropriate to include are measures of within-scan head motion if available (e.g., when using a 3D MPRAGE or SPACE with a navigator) and the correction parameters mentioned above (slope, mean B1+ measure, smoothing FWHM, pseudo-transmit threshold, and the value required to account for the effects of smoothing). Finally, we found that the reference voltage of the transmit coil was a useful additional covariate of no interest that helped to further remove dependence on the scanner generated B1+ field.

Toy Statistical Model for Demonstrating the Effects of Individual participant T1w/T2w Myelin Map Correction

To illustrate the beneficial effects of T1w/T2w myelin map correction on cross-participant analyses that relate participant demographics, behavior, and disease states to T1w/T2w myelin maps, we developed a "toy" statistical model. The variables of interest are sex, age, and Body Mass Index (BMI). We use multiple linear regression after normalizing the input variables to demonstrate simple linear relationships within the 360 cortical areas identified in the Human Connectome Project's multi-modal parcellation V1.0 (Glasser et al., 2016a). Importantly, in this work we focus exclusively on the effects of bias and nuisance from spurious B1+ variations on the T1w/T2w myelin maps used in these linear regressions and not on any neuroscientific questions of interest. Such neuroscientific questions will be properly addressed in subsequent publications using more advanced statistical models. For this reason, we do not assess the statistical significance of these effects and will comment on them only qualitatively insofar as it is useful to describe the effects of the transmit field correction.

T1w/T2w Myelin Maps for Neuroanatomical Use Cases After Transmit Field Correction

Although the covariates can be included in statistical models as described above, it is also helpful to produce T1w/T2w myelin maps with these parameters regressed out of the data (to correct the individual participant myelin map means and remove residual spatial biases) for neuroanatomical use cases. Thus, we also produced T1w/T2w myelin maps with these parameters regressed out. These maps may have some residual artifacts, for example, from (1) within-scan head motion (either in scans without the vNav motion correction or where the limit on the number of additional TRs reacquired due to motion is met), or (2) head coil malfunction in one scan but not the other (T1w or T2w). The "MyelinMap_BC" approach (described in the Introduction and in Glasser et al., 2013), using the new B1+ corrected reference group myelin map, may still be preferable to these regressed T1w/T2w myelin maps in these situations for neuroanatomical use cases, given that even covariate regression cannot eliminate all forms of low spatial frequency myelin map biases such as those described above.

Correction of T1w/T2w Ratio Volumes

Although the relationship between the T1w/T2w ratio and myelin is less well constrained outside of cortical grey matter (Glasser and Van Essen 2011), some investigators have been interested in exploring the T1w/T2w ratio in subcortical structures, including white matter. For all above data the B1- corrections were applied to the T1w/T2w volume appropriately, and the B1+ transmit field measure was regularized initially in volume space before being projected to the surface as described above. Thus, it can be trivially applied with the slope that was already computed on the surface, and so we also generate corrected T1w/T2w ratio volumes for all datasets. However, it is NOT appropriate to attempt "VBM-style" analyses of these T1w/T2w ratio volumes in lieu of proper surface-based cortical analyses for reasons described elsewhere (Glasser et al., 2016b; Coalson et al., 2018), and also because partial volume effects of CSF, grey matter, and white matter will completely dominate the differential effects of myelin content in traditional volume-based analyses (i.e., the differences in T1w/T2w ratio between tissues are much larger than those within tissues). Skeletonized methods such as Tract-Based-Spatial-Statistics (TBSS; Smith et al., 2006) offer the best option for voxelwise white matter analysis

(e.g. Operto et al., 2019), though tractography-based tract analyses would likely be even better (Chen et al., 2017; Thompson et al., 2020).

Results

HCP-YA: Group Average T1w/T2w Myelin Map Transmit Field Correction

We begin by illustrating the group average T1w/T2w myelin map (top two rows left) in Figure 5 together with the group average AFI map (top two rows right). The group myelin map is a simple surface-based average, whereas the TR=20 and TR=120 AFI datasets were separately averaged on the surface before computing the flip angle using the equation from (Yarnykh 2007; $\text{acos}\{(120/20 * \text{VolTR}_{120}/\text{VolTR}_{20} - 1)/(120/20 - \text{VolTR}_{120}/\text{VolTR}_{20})\}$). This approach is acceptable because the flip angle pattern is generally consistent across participants (same scanner and sequence), the real flip angles form a smooth map, and the nonlinear steps are nearly linear within the range of the data. As in Figure 4 the group average AFI map lacks obvious artifacts and can directly be used to correct the group average T1w/T2w myelin map. Importantly, the left-right asymmetry (computed using the asymmetry index in Equation #6) in the T1w/T2w myelin map (bottom row left) and AFI map (bottom row right) are highly correlated spatially (r=0.94) even though different in magnitude (necessitating the fitting procedure described in the methods).

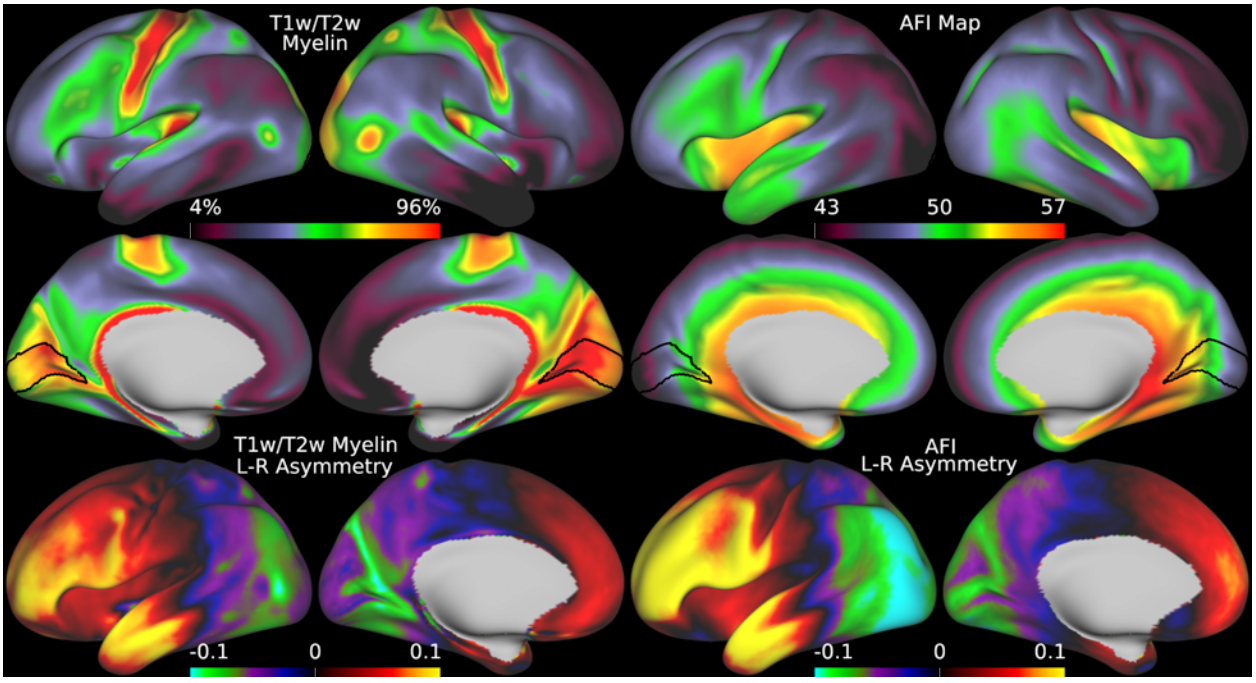

Figure 5 illustrates the group average T1w/T2w myelin maps (top two rows left), group average AFI maps showing the B1+ field as flip angles (top two rows right, in degrees), left-right asymmetry in T1w/T2w myelin maps (bottom row left), and left-right asymmetry in AFI maps (bottom row right). Area V1 is outlined in black.

Thus, minimizing left-right asymmetry in the group average T1w/T2w myelin map (Equation #6) is an appropriate cost function for optimizing the slope parameter in Equation #5. Figure 6 shows the original and corrected T1w/T2w myelin maps and their asymmetry maps. The correction has markedly reduced the spatial correlation of the left-right asymmetry between the AFI and T1w/T2w myelin maps (r=0.10). Additionally, beyond just correcting hemispherically asymmetric biases, the correction has also addressed *symmetric* biases in the myelin maps. For example, the homogeneity of visual area V1, which contains a strong central to peripheral flip angle gradient (see in Figure 5), is markedly improved by this correction as seen in Figure 6. Finally, there are some left-right hemispheric differences that are due to slightly different positions of the cortical areas relative to cortical folds after the folding-based registration used to align the FS_LR template across hemispheres (Van Essen et al., 2012), for example the positive/negative couplets near the MT+ complex, LIPv in the intraparietal sulcus, and along the precentral gyrus (see faint circles).

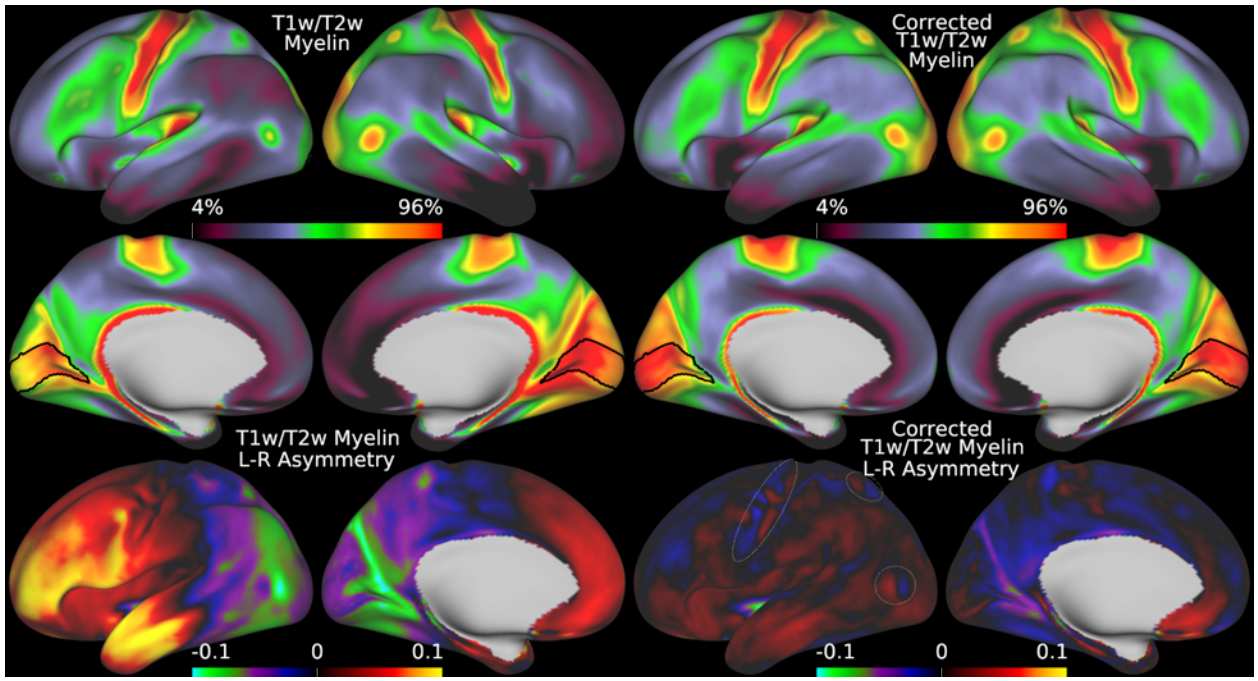

Figure 6 illustrates the group average original T1w/T2w myelin maps (top two rows left), group average corrected T1w/T2w myelin maps (top two rows right), left-right asymmetry in original T1w/T2w myelin maps (bottom row left), and left-right asymmetry in corrected T1w/T2w myelin maps (bottom row right). Area V1 is outlined in black. The faint circles represent the MT+ complex, LIPv, and M1.

HCP-YA: Group Average Effects of Transmit Field Correction of Individual T1w/T2w Myelin Maps

Figure 7 shows a comparison of the group corrected myelin map (with the Left-Right cost function; Equation #6) and the mean individual corrected myelin map (with the Individual-Template cost function; Equation #7). These maps are nearly identical (spatial correlation of

r=0.99), illustrating that the individual participant correction with regularized AFI maps is in the aggregate well behaved.

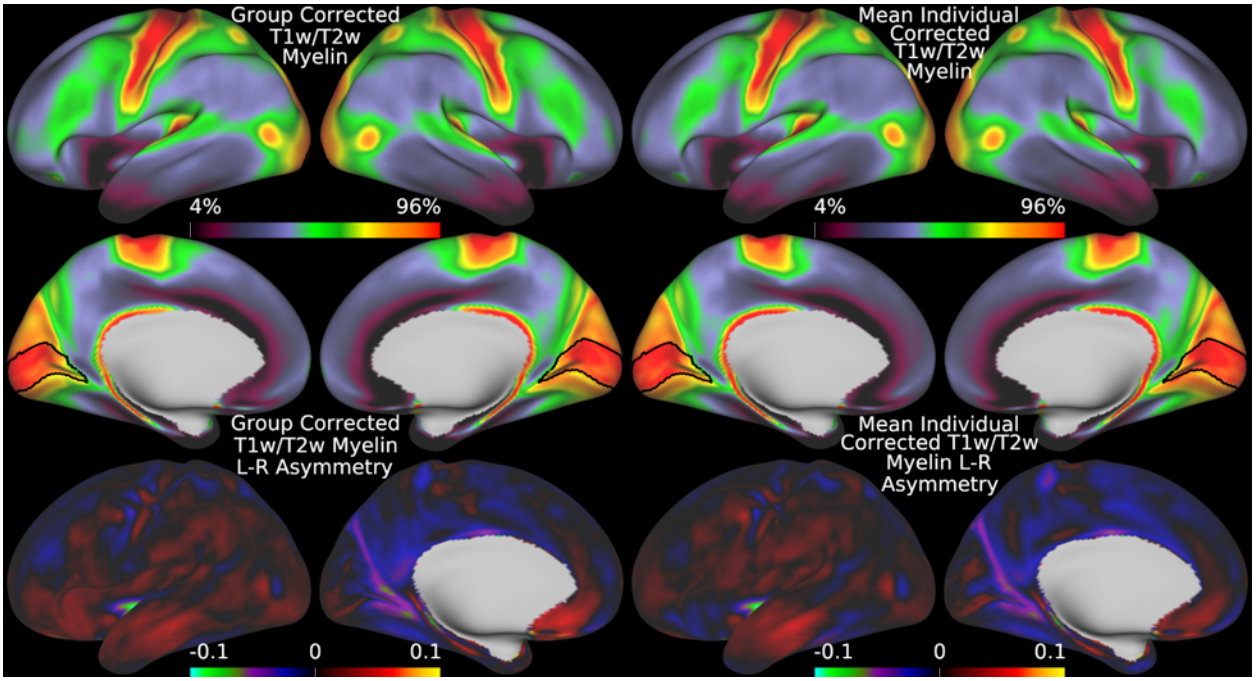

Figure 7 illustrates a comparison between the correction applied to the group average and the mean of the individually corrected myelin maps, showing that they are very similar.

HCP-YA: Statistical Effects of Individual T1w/T2w Myelin Map Transmit Field Correction

Although obtaining more accurate group average T1w/T2w myelin maps is useful neuroanatomically, the core goal of this paper is enabling cross-participant analyses that relate T1w/T2w myelin maps to other parameters of interest such as participant demographics, behavior, or disease state. Figure 8 illustrates a multiple linear regression of sex, age, and BMI with original T1w/T2w myelin maps, the AFI maps, and the B1+ corrected T1w/T2w myelin maps (with the participant-wise covariates of no interest included in the corrected T1w/T2w myelin map regression). This figure demonstrates the sizable biases present in the original maps, the fact that the same biases are present in the AFI maps, and elimination of these biases in the corrected maps. For example, the original uncorrected maps suggest higher T1w/T2w values in men than in women across most cortical areas, particularly in the right parietal cortex (left 3 columns). These spurious findings are reflected in the AFI maps, and thus they are absent in the corrected T1w/T2w myelin maps, where overall the differences are quite modest, and instead women have slightly higher T1w/T2w values than men across most of the brain, except for prefrontal cortex and left anterior inferior parietal cortex. This spurious confounding/biasing effect is most likely due to the differences in body and head size, shape, and composition between men and women leading to different flip angles achieved (see Table 1), which increases the T1w/T2w ratio in men (chiefly by reducing the denominator). Age (middle 3 columns) shows a different story. In young adults, head size and BMI are not

particularly correlated with age (see Table 1). As a result, there is not a strong relationship between age and transmit field (or voltage) across participants (center column in Figure 8). Uncorrected T1w/T2w values show relatively anatomically non-specific increases with age, and correction of the transmit field effects enhances the anatomical specificity (e.g., the motor cortex) of this relationship by reducing nuisance effects of head size and weight/BMI on the transmit field. Finally, BMI (right 3 columns) has a direct effect on the transmit field due to coil loading. In this figure, BMI shows a strong relationship with both the original T1w/T2w ratio and the AFI transmit field, with a left vs right asymmetry very similar to that illustrated in Figure 5. This effect is dramatically reduced in the corrected T1w/T2w myelin maps.

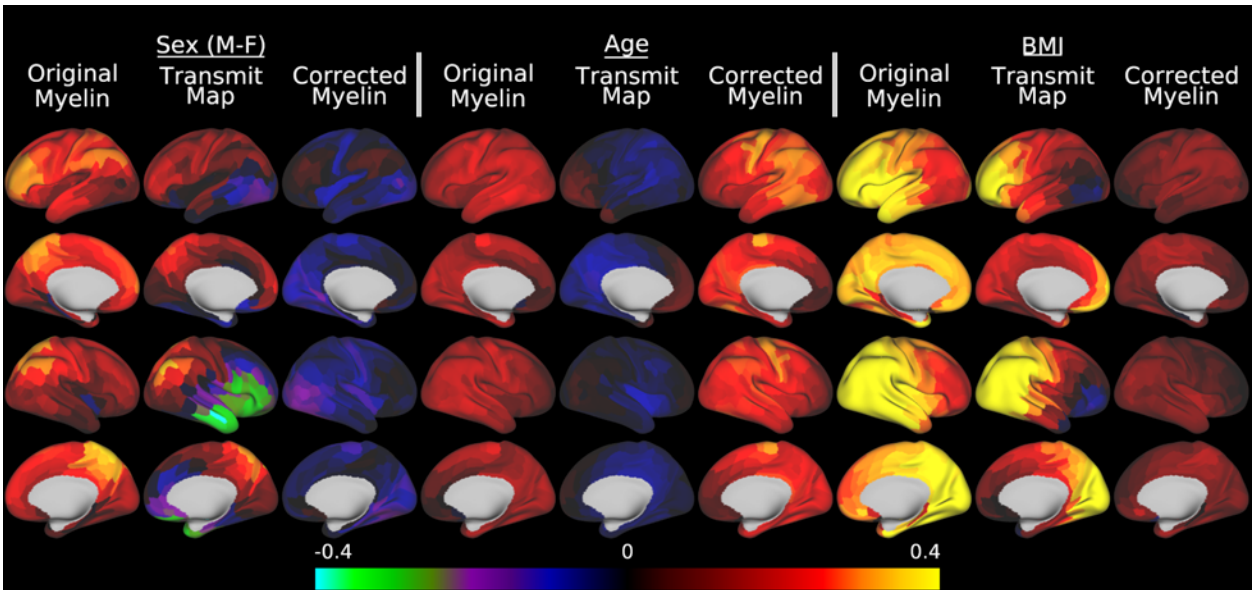

Figure 8 illustrates the effect of transmit field correction of individual T1w/T2w myelin maps on a toy statistical model that includes sex, age, and BMI in a multiple linear regression. Within each grouping by variable of interest, normalized beta values enable comparisons across original T1w/T2w myelin maps, the AFI transmit maps, and the B1+ transmit field corrected T1w/T2w myelin maps. For the corrected maps, the covariates of no interest (see Methods) are included in the model but not illustrated.

HCD: Group Average Effects of Pseudo-Transmit Field Correction of Individual T1w/T2w Myelin Maps

Figure 9 illustrates that the mean pseudo-transmit field has a similar pattern to the empirically measured (via AFI scans) transmit field in the HCP-YA data, particularly the center versus peripheral bias, and that the HCD data also have left-right asymmetry, though not as pronounced as the HCP-YA data. Figure 10 demonstrates that individual participant pseudo-transmit field correction using the reference T1w/T2w myelin map generated from the HCP-YA data also successfully removes these asymmetries in HCD data, as in the real transmit field correction in HCP-YA data, and the symmetric biases, producing a more homogeneous area V1.

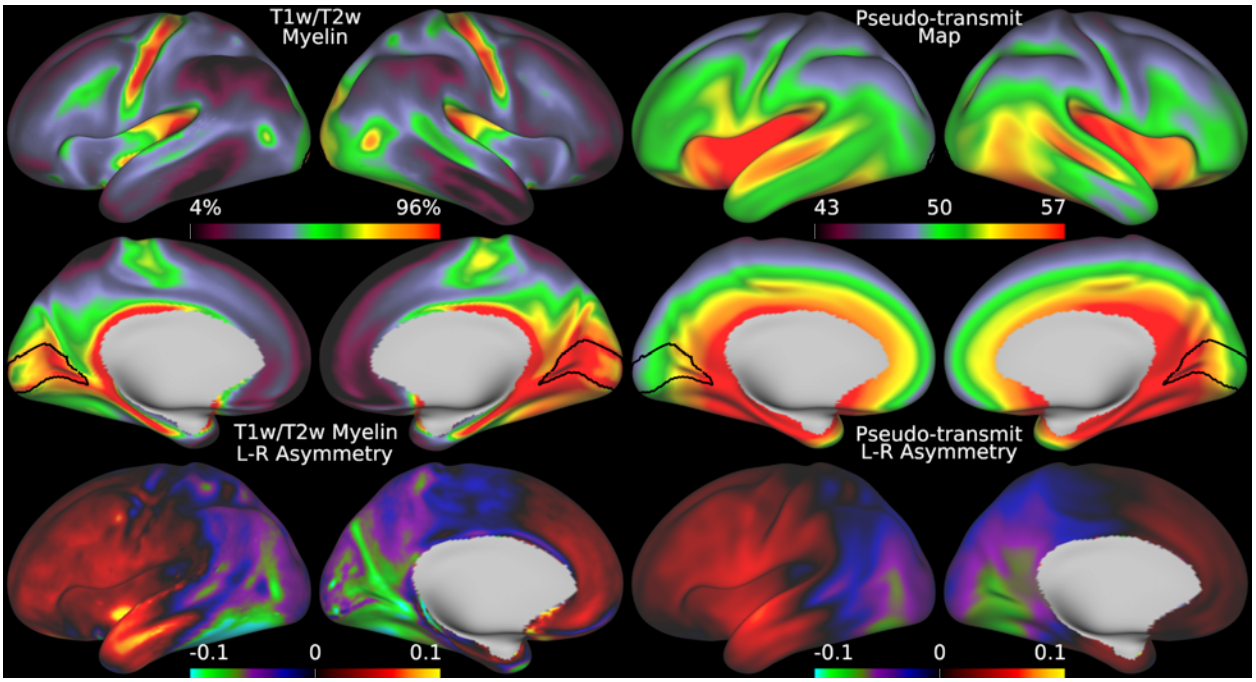

Figure 9 shows the original group average T1w/T2w myelin map from the HCD dataset in the top left two rows together with the regularized group average pseudo-transmit field map in the top right two rows. The bottom row shows the corresponding asymmetry maps.

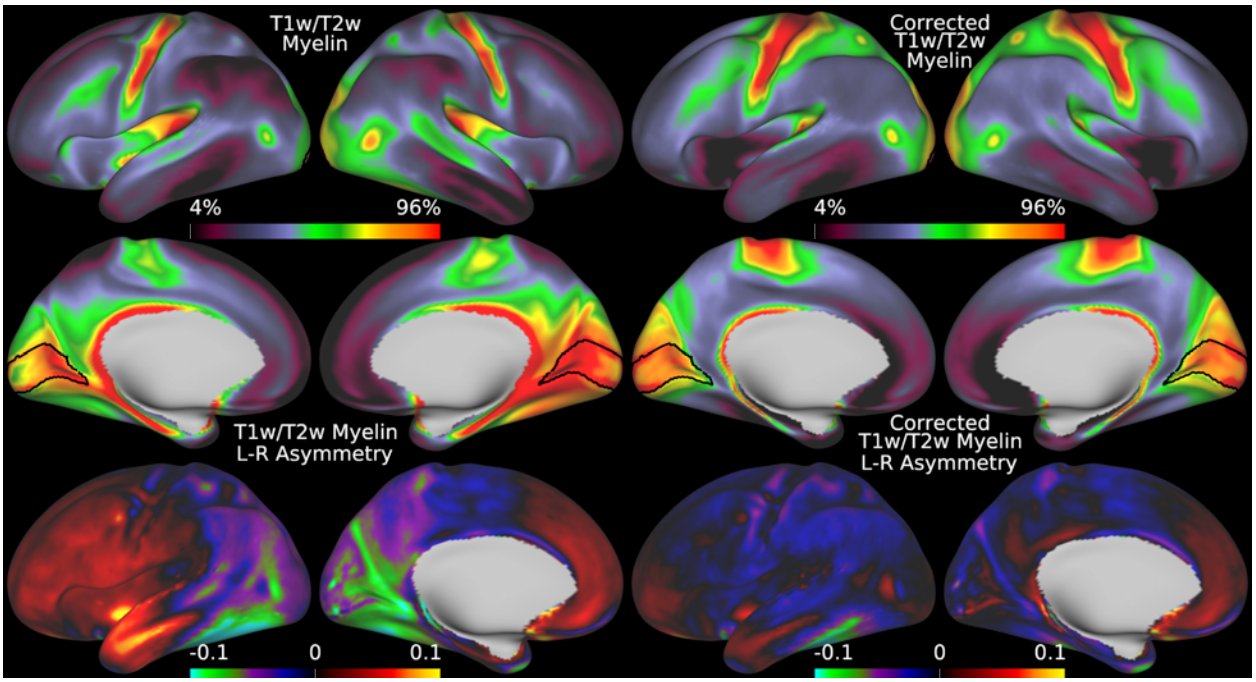

Figure 10 shows the original group average T1w/T2w myelin map from the HCD dataset in the top left two rows together with the mean corrected individual participant T1w/T2w myelin maps. The bottom row shows the corresponding asymmetry maps. The correction works well for most of the brain, but somewhat less well for areas of high susceptibility where the pseudo-transmit field must be imputed from surrounding valid data.

HCD: Statistical Effects of Individual T1w/T2w Myelin Map Pseudo-Transmit Field Correction

Similar to HCP-YA, the toy statistical analysis of sex, age, and BMI contains biases related to head and body size (Figure 11). For example, there is a similar pattern of boys > girls across most of the brain in the uncorrected data, whereas in the corrected data, a weak trend of girls > boys is seen across most of the brain. Although we expect a strong effect of age in developing participants, this is confounded with increasing head and body size (Table 1). There remains a strong effect of age after correction. Finally, BMI shows the presence of a strong positive but hemispherically asymmetric bias in the uncorrected maps, as was also the case for HCP-YA. Importantly, BMI in growing adolescents likely represents a different physiological construct than BMI in adults. Regardless of interpretation, the important point here is that the bias is eliminated in the corrected maps.

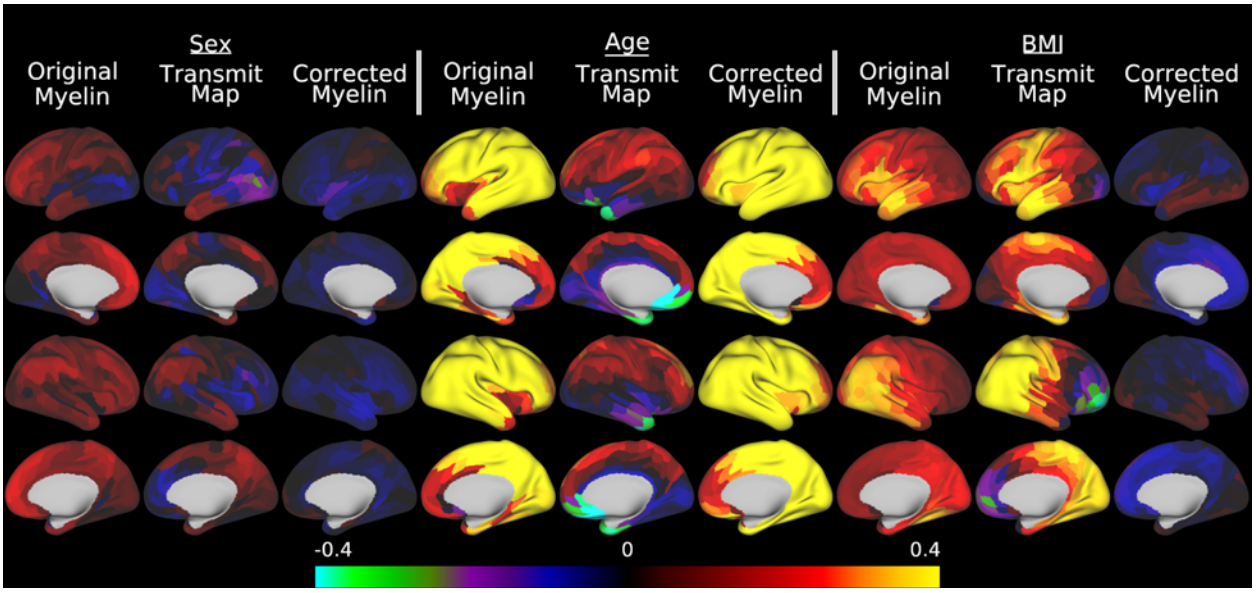

Figure 11 illustrates the effect of pseudo-transmit field correction of individual T1w/T2w myelin maps on a toy statistical model including sex, age, and BMI in a multiple linear regression. Within each grouping by variable of interest, normalized beta values are shown for original T1w/T2w myelin maps, the pseudo-transmit maps, and the pseudo-transmit corrected T1w/T2w myelin maps to enable comparison across different types of data. For the corrected maps, the covariates of no interest (see Methods) are included in the model but not illustrated.

HCA: Group Average Effects of Pseudo-Transmit Field Correction of Individual T1w/T2w Myelin Maps

Figure 12 illustrates that the mean pseudo-transmit field has a similar pattern to the estimated transmit field in the HCP-YA data and the pseudo-transmit field in the HCD data, and that the HCA data also have left-right asymmetry, with the magnitude of the asymmetry lying between the HCP-YA and HCD data. Figure 13 demonstrates that individual participant pseudo-transmit field correction using the reference T1w/T2w myelin map generated from the HCP-YA data also

successfully removes these asymmetries in the HCA data, as in the HCD data and the real transmit field correction in HCP-YA data, and produces a more homogeneous area V1.

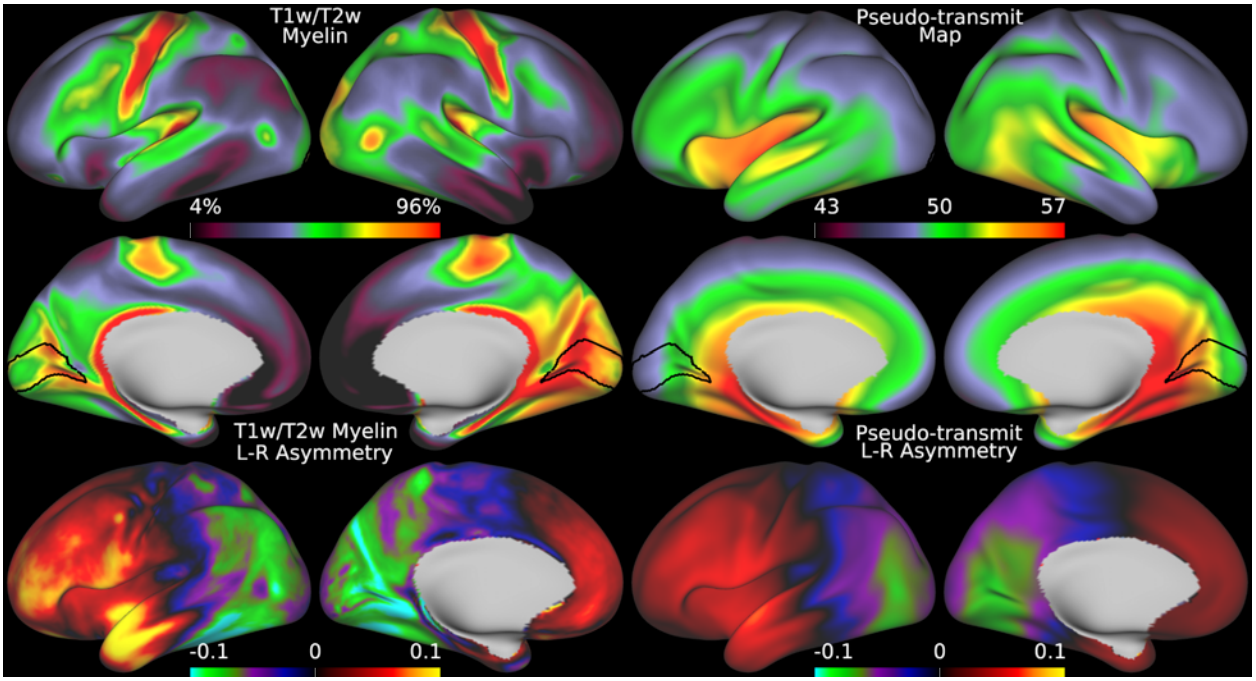

Figure 12 shows the original group average T1w/T2w myelin map from the HCA dataset in the top left two rows together with the regularized group average pseudo-transmit field map in the top right two rows. The bottom row shows the corresponding asymmetry maps.

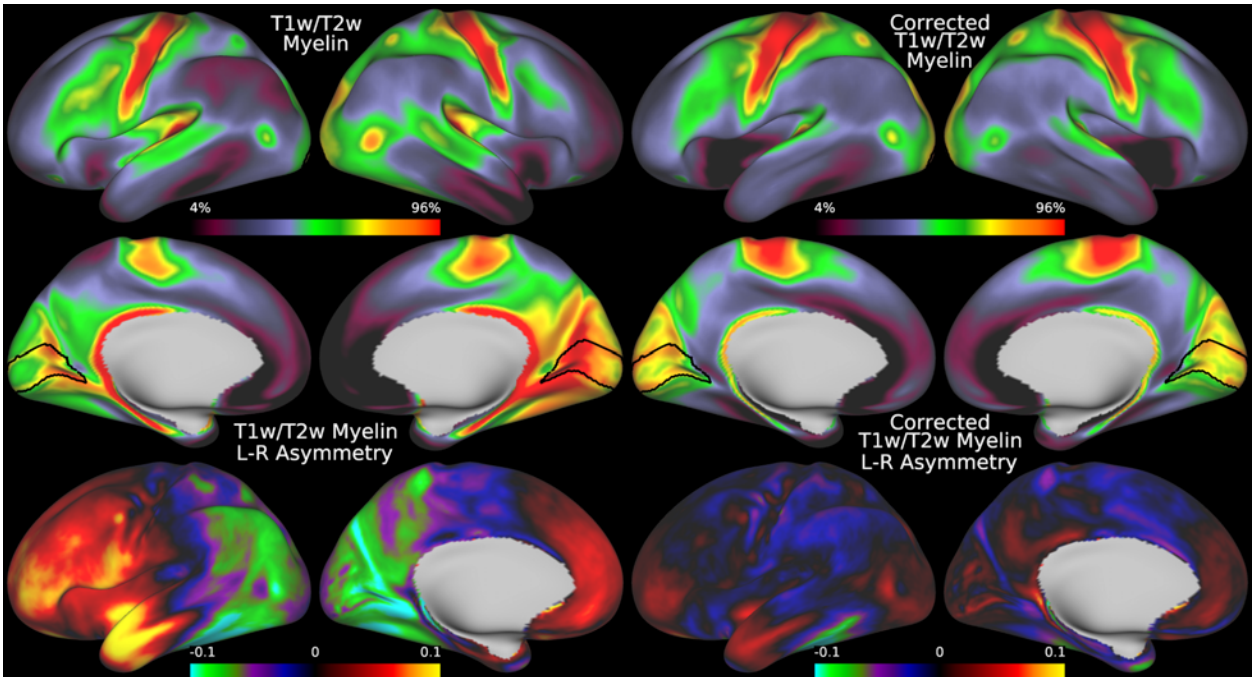

Figure 13 shows the original group average T1w/T2w myelin map from the HCA dataset in the top left two rows together with the mean pseudo-transmit corrected individual participant T1w/T2w myelin maps. The bottom row shows the corresponding asymmetry maps.

HCA: Statistical Effects of Individual T1w/T2w Myelin Map Pseudo-Transmit Field Correction

Similar to HCP-YA and HCD, the toy statistical analysis of sex, age, and BMI contains biases related to head and body size (Figure 14). In HCA participants, the correction still improves the symmetry of the maps for sex, age, and BMI. There is correlation between age and the transmit field map, but again the correction increases the relationship with age, likely by removing spurious nuisance correlations. Finally, BMI shows a hemispherically asymmetric pattern in the uncorrected data that is more similar to HCP-YA than HCD and shows a similar symmetrization in the corrected data, consistent with the strong correlation between BMI and B1+ voltage (Table 1).

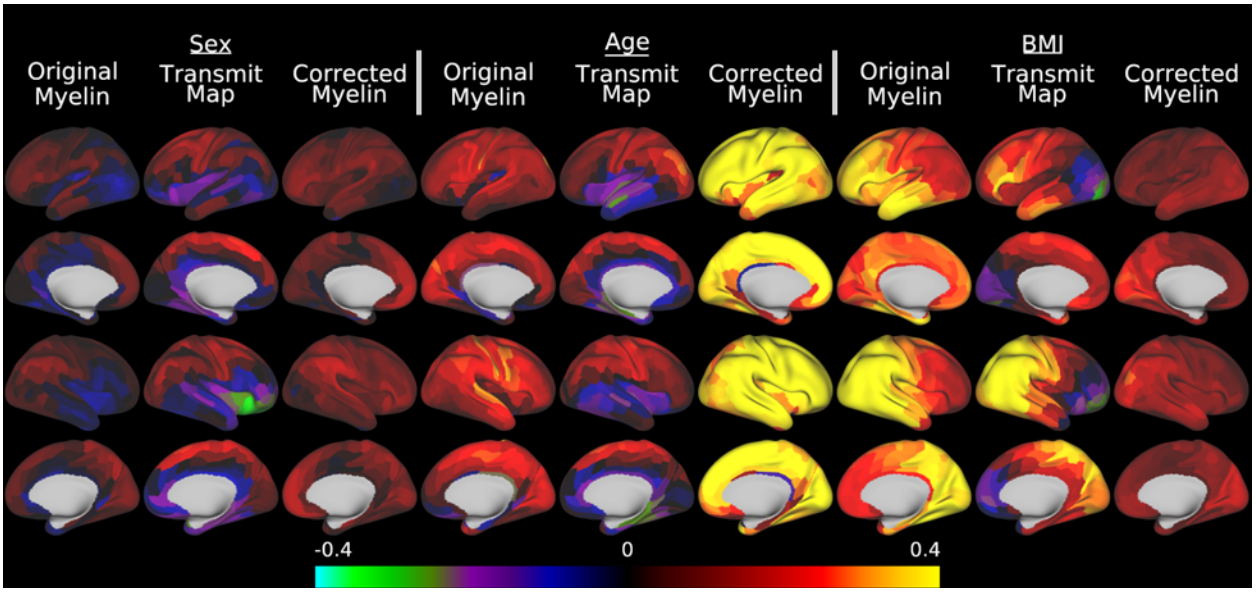

Figure 14 illustrates the effect of pseudo-transmit field correction of individual T1w/T2w myelin maps on a toy statistical model including sex, age, and BMI in a multiple linear regression. Within each grouping by variable of interest, normalized beta values are shown for original T1w/T2w myelin maps, the pseudo-transmit maps, and the corrected T1w/T2w myelin maps to enable comparison across different types of data. For the corrected maps, the covariates of no interest (see Methods) are included in the model but not illustrated.

NHP_NNP: Group Average T1w/T2w Myelin Map Transmit Field Correction in Macaques

Relative to humans, macaques have much smaller heads and bodies, and as a result the inhomogeneity of flip angles is much lower (Figure 15; flip angle range shown is +3 to -3 degrees in the macaque (units displayed are flip angle * 10 as generated by the scanner with a reference flip angle of 80 degrees instead of 50 degrees as in humans) versus +7 to -7 degrees in the human in Figure 5). Correspondingly, there is much less hemispheric asymmetry in both

the group average T1w/T2w myelin map and the B1+ transmit field map. Nevertheless, this asymmetry still shows a similar pattern and enables estimation of a corrected macaque myelin map (Figure 16). This map eliminates the left-right asymmetry and a symmetric center > peripheral bias in the macaque myelin map, just as in humans.

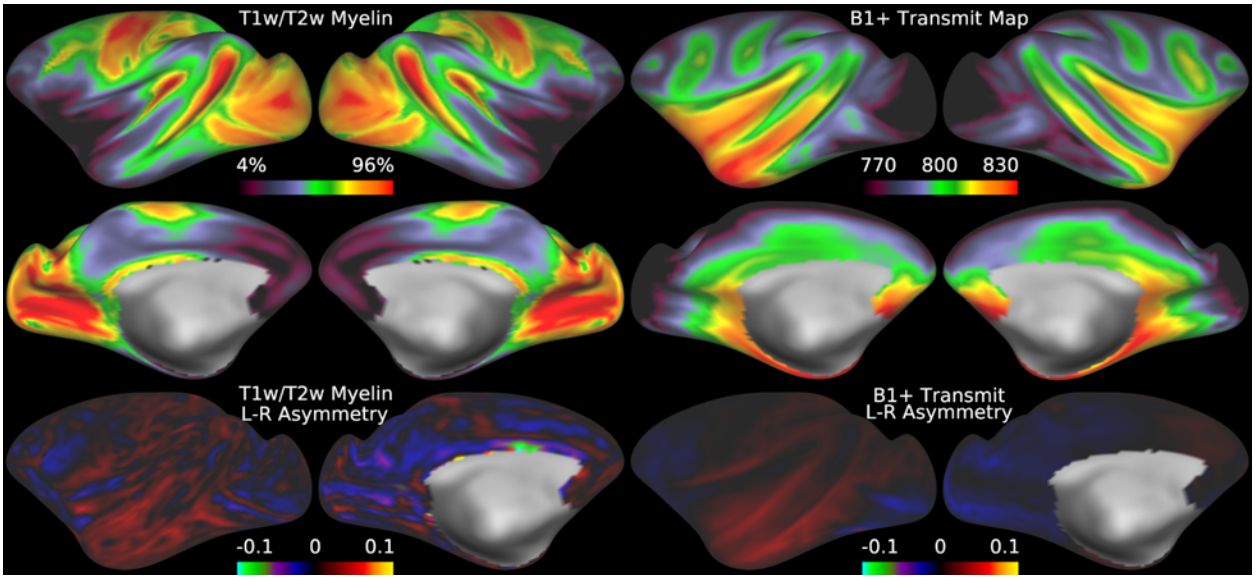

Figure 15 illustrates the group average T1w/T2w myelin maps (top two rows left), group average B1+ transmit field maps (top two rows right), left-right asymmetry in T1w/T2w myelin maps (bottom row left), and left-right asymmetry in group average B1+ transmit field maps (bottom row right). The units of the B1+ transmit field map are flip angle * 10, as generated by the scanner with a reference flip angle of 80 degrees instead of 50 degrees as in humans.

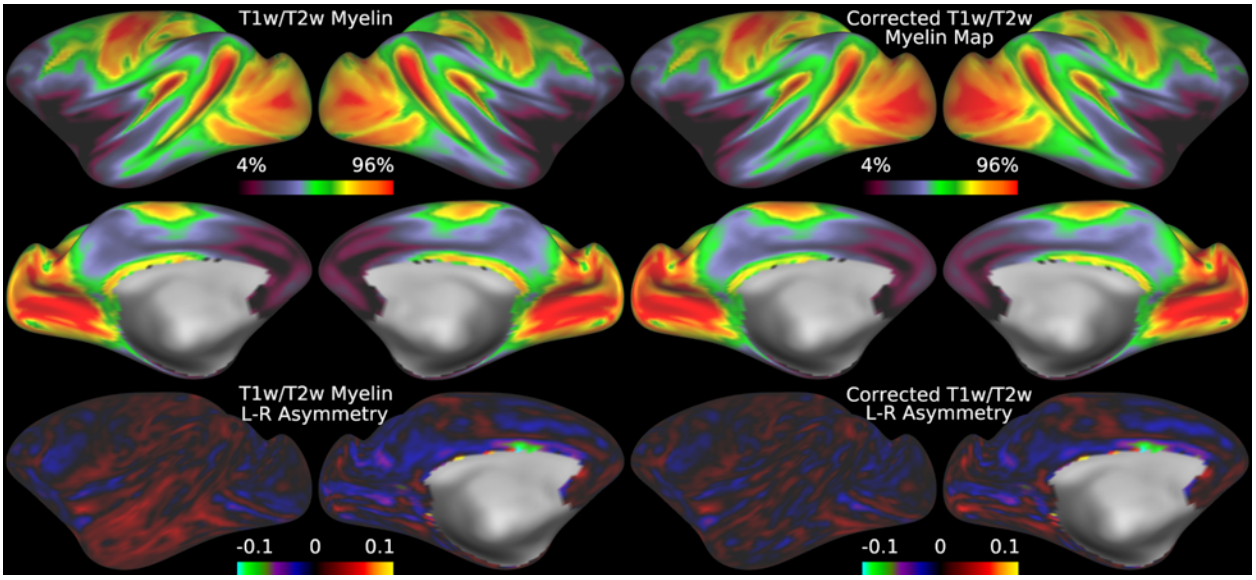

Figure 16 illustrates the group average original T1w/T2w myelin maps (top two rows left), group average B1+ transmit field corrected T1w/T2w myelin maps (top two rows right), left-

right asymmetry in original T1w/T2w myelin maps (bottom row left), and left-right asymmetry in corrected T1w/T2w myelin maps (bottom row right).

NHP_NNP: Group Average Effects of Transmit and Pseudo-Transmit Field Correction of Individual T1w/T2w Myelin Maps

Macaques are the only dataset in which both transmit field and pseudo-transmit field data are available for T1w/T2w myelin map correction. The spatial correlation between the mean of the individually corrected data and the group reference correction is r=0.999 for the transmit field approach and r=0.995 for the pseudo-transmit field approach. Visually, the mean individual corrected maps are practically indistinguishable (Figure 17) and the mean across participants regularized B1Tx and pseudo-transmit fields correlate at r=0.87.

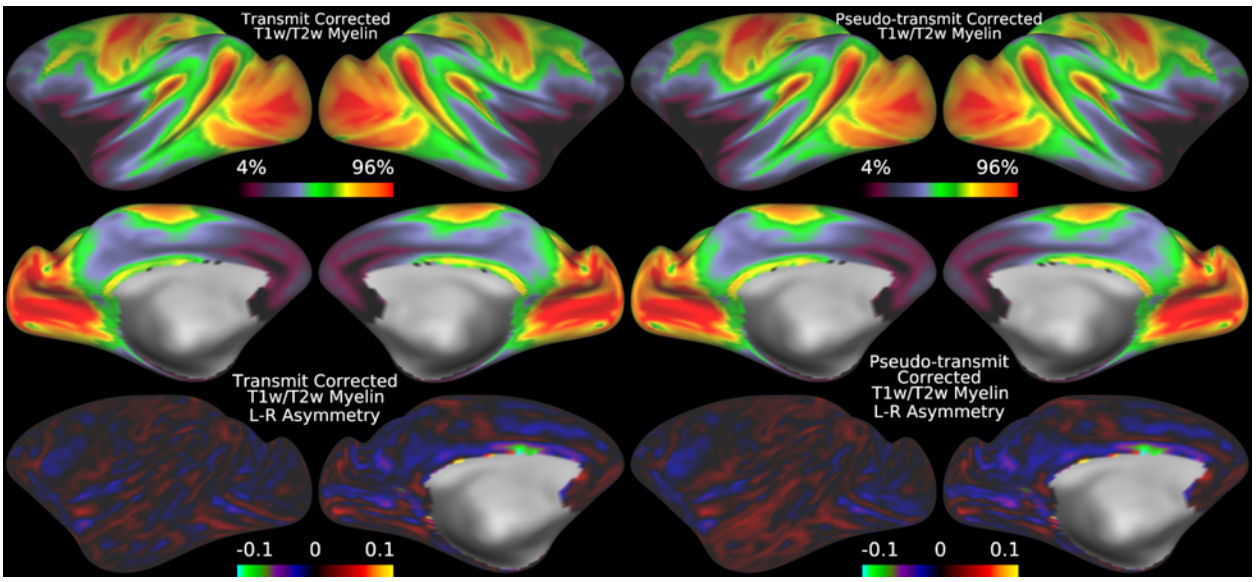

Figure 17 illustrates mean individual transmit field (top left two rows) and pseudo-transmit field (top right two rows) corrected T1w/T2w myelin maps. The corresponding asymmetry maps are shown in the bottom left and right row respectively.

Effects of Transmit and Pseudo-Transmit Field Correction of T1w/T2w Myelin Maps on NHP_NNP Cross-animal Variability and HCP-YA Within-participant Reproducibility

The 16-animal macaque dataset is the only one in which we can directly compare the transmit and pseudo-transmit field correction methods for T1w/T2w Myelin Maps. Although the HCP-YA acquired the gradient echo and spin echo EPI data that are prerequisites for the pseudo-transmit correction, it was always in a separate imaging session from the T1w and T2w scans. Thus, it would not be possible to tell whether any differences between transmit and pseudo-transmit field corrections were due to differing effectiveness or simply poorer matching of B1+ fields across different sessions due to differences in head position and/or B1+ homogeneity. Figure 18 illustrates this direct comparison. Eta^2 is used to quantify similarity (Cohen et al., 2008). Both transmit and pseudo-transmit field corrections improve agreement with the group

average data, but transmit field correction performs a bit better. Regressing out the covariates (or including them in the statistical model) further improves both transmit and pseudo-transmit data and makes their performance more similar. Thus, we recommend using the covariates (or regressing them out of the maps) depending upon whether one is analyzing a larger statistical model or simply using the maps for downstream neuroanatomical analyses. Both transmit and pseudo-transmit approaches are reasonable, though explicit transmit field mapping is a little better when available scanning time permits its use.

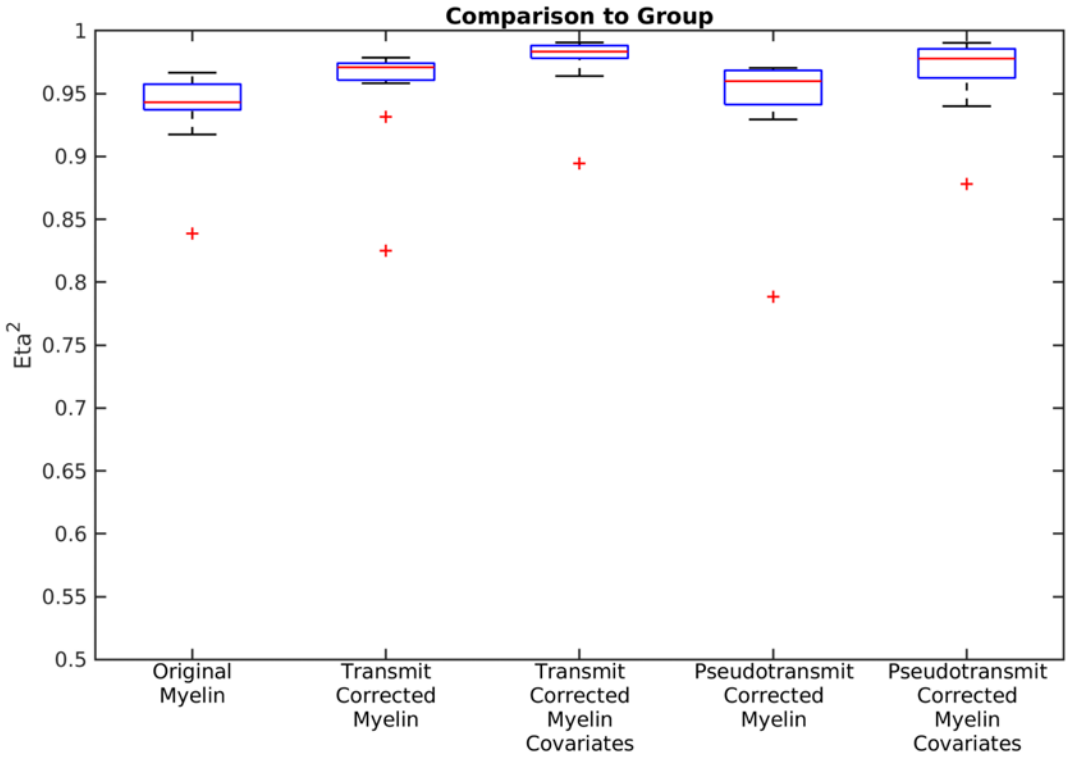

Figure 18 shows boxplots of the eta^2 between the group average corrected reference myelin map and the individual myelin maps for the n=16 macaque datasets. This is shown for the original T1w/T2w myelin values, the transmit field corrected T1w/T2w myelin values without and with covariate regression, and the pseudo-transmit field corrected T1w/T2w myelin values without and with covariate regression. The red line is the median, the edges of the box are the 25th and 75th percentiles (the interquartile range, IQR), the whiskers extend from the end of the IQR to the furthest observation within 1.5 * IQR, and the outliers (+'s) are the data points beyond those limits. The range of eta^2 is 0-1, with identical inputs yielding 1; inputs that are strictly the negative of each other yielding 0; and values in between represent varying degrees of 'similarity'.

The HCP-YA data includes 41 participants in which all modalities used in this study were acquired and processed twice in a test-retest design. Similar to Figure 18, Figure 19 top row illustrates the improvement in eta^2 with transmit field correction and further improvement with

covariate regression (or inclusion in the statistical model). A small amount of surface-based smoothing (4mm FWHM) further improves the similarity of the individual data to the group data, as it eliminates random noise and reduces small surface reconstruction differences. Median eta^2 is about 0.9 after correction and surface-based smoothing. The bottom row illustrates the test-retest eta^2 with and without smoothing for each T1w/T2w myelin map and also for unregularized and regularized AFI maps. Reproducibility across visits is minimally increased by the transmit field correction, peaking at median eta^2 of around 0.9 in smoothed data with correction and covariate regression. The raw AFI maps have markedly reduced reproducibility with median eta^2 of around 0.76. Regularization improves this to around 0.9; however. Notably, the transmit coil reference voltages have a test-retest eta^2 of 0.95, indicating that the poor test-retest reproducibility of the AFI maps is likely not due to marked differences in scanner B1+ transmit efficiency, but rather some combination of random artifacts and differences in head position inside the body coil. This finding also explains why test-retest reproducibility is similar between corrected and uncorrected data (i.e., the B1+ transmit field effects are reproducible themselves within participants). Overall, it is clear that if AFI is to be used routinely going forward, the artifacts shown here would need to be reduced to produce optimal corrections. If AFI cannot be improved, the B1+ mapping scans used in the macaques would be a better alternative if transmit field correction is desired, given that the B1Tx required less regularization (mean FWHM=13) than the AFI (mean FWHM=33mm).

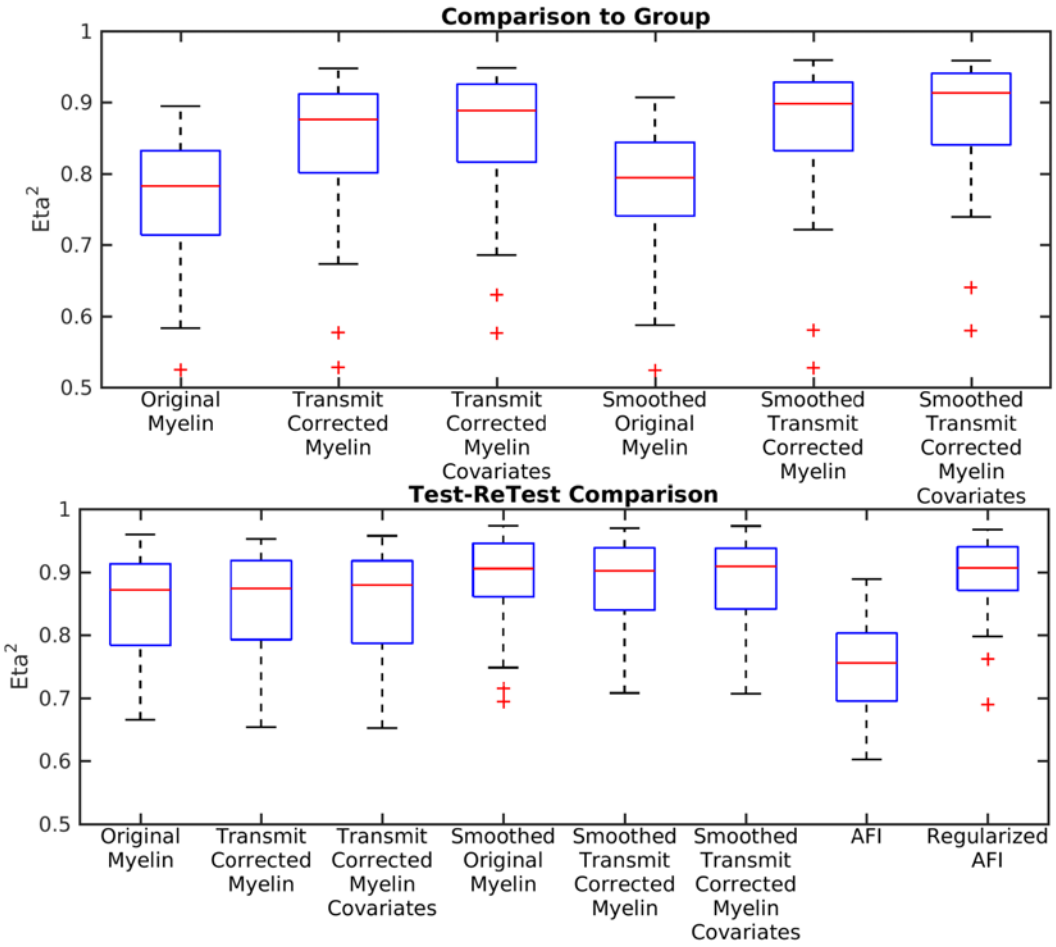

Figure 19 shows box plots across HCP-YA test-retest participants (n=41). Along the top row the eta^2 between each scan and the reference group average corrected myelin map is illustrated for both the 'Test' and 'ReTest' visits of the Test-ReTest participants. This is shown for the original T1w/T2w myelin values, the transmit field corrected T1w/T2w myelin values without and with covariate regression, and the transmit field corrected T1w/T2w myelin values without and with covariate regression with smoothing. Along the bottom row, the eta^2 between the Test and ReTest data is shown for each measure including unregularized and regularized AFI. Smoothing is 4mm FWHM on the surface, which improves test-retest reproducibility some due to reduction in variability from random noise and small surface reconstruction differences.

Discussion

We have demonstrated an accurate approach to B1+ transmit field correction of T1w/T2w myelin maps using either empirically measured transmit field maps or a derived analog that we term 'pseudo-transmit' field maps. We used the finding of highly correlated hemispheric asymmetries between the T1w/T2w cortical myelin maps and B1+ transmit field maps to develop an appropriate correction of both cortical T1w/T2w myelin maps and volume T1w/T2w

ratio maps at both the individual and group levels in humans and non-human primates. The resulting corrected T1w/T2w myelin maps are largely free of neurobiologically implausible asymmetries, additional symmetric biases, and spurious correlations with head and body size. Such B1+ corrected T1w/T2w myelin maps are now appropriate for more detailed studies of inter-participant differences with regard to behavior, development, aging, and disease, as well as cross-species differences, some of which will be presented elsewhere.

Interpretation of individual differences in cortical T1w/T2w myelin maps and their relationship to cortical thickness deserves comment. The following interpretational guidelines are likely to apply in situations that are not confounded by blurring from motion (which will tend to cause cortex to appear thinner (Reuter et al., 2015) due to errors in surface placement and increased partial voluming of grey matter with CSF and white matter contamination), edema (which will decrease the T1w/T2w ratio), or metal deposition (e.g., iron; which will increase the T1w/T2w ratio): 1) Apparent increases in cortical T1w/T2w myelin maps if cortical thickness is constant or increasing would likely be due to increases in myelinated axons (e.g., as could occur during development; Bozek et al., 2018). 2) If cortical T1w/T2w measured myelination is increasing at the same time cortical thickness is decreasing, the apparent increase in myelin would likely be due to a decrease in the non-myelinated, plasticity supporting cellular constituents of cerebral cortex (i.e., dendrites, spines, synapses, and glia; Glasser et al., 2014), resulting in a relative increase in myelin density in the remaining cortex (e.g., as may occur during aging), and not necessarily an absolute increase in myelination. 3) Decreases in T1w/T2w with unchanged cortical thickness would likely be due to demyelination (e.g., as could occur in a demyelinating disease). 4) Decreases in T1w/T2w with decreased cortical thickness would likely be due to direct cortical damage (e.g., as could occur in encephalomalacia). 5) Decreases in T1w/T2w with increasing thickness are not expected to occur in the absence of pathology. Thus, it is helpful to consider what is happening with cortical thickness when interpreting differences in cortical T1w/T2w myelin maps. That said, if image resolution is too low or the surface reconstruction is not optimized, apparent cortical thickness itself can be influenced by cortical myelin content (e.g., Figure 7 auditory core in Glasser and Van Essen 2011) or non-cortical structures such as dura or blood vessels, which can then cause T1w/T2w myelin map artifacts.

Interpretation of volume T1w/T2w maps is more challenging than cortical (surface) T1w/T2w myelin maps, as the relationship between the T1w/T2w MRI contrast and myelin content in the deep white matter has not been shown to be as tight as that seen in cortical grey matter (Glasser and Van Essen 2011). Figure 20 illustrates the group average B1+ corrected T1w/T2w volume map from the HCP-YA data. Several well-known fiber tracts are visible including the cortico-spinal tract and the optic radiations. Interestingly, while the optic radiations have high T1w/T2w signal, the cortico-spinal tract has relatively low T1w/T2w signal. Also, the motor and visual callosal fibers have relatively lower T1w/T2w signal (a similar pattern is seen with quantitative R1; Harkins et al., 2016). Conversely, the association fibers of the callosum have relatively higher T1w/T2w signal, as does the prefrontal white matter, which is the opposite pattern seen in cortical grey matter (with association areas generally having lower myelin content than primary areas). White matter T1w/T2w signal likely depends on more than the number of myelin wrappings per axon. Additional considerations such as average axonal

packing density and average axonal diameter will affect the myelin volume fraction of any given voxel (Stikov et al, 2015). Overall, within the white matter, the T1w/T2w ratio is likely related to the lipid-to-water content ratio. That said, as has been seen clinically by neuroradiologists for decades, frank white matter demyelination will result in decreased T1w signal and increased T2w signal leading to a decreased T1w/T2w ratio, and a normally developing young child's age can be estimated to within a few months based on white matter signal changes found in T1w and T2w images by a skilled pediatric neuroradiologist. Also, as stated previously, cortical myelin content is essentially uninterpretable even in unsmoothed group average T1w/T2w volumes due to cross-participant misalignments of the cerebral cortex, resulting in large partial voluming effects of CSF and white matter that swamp the intracortical grey matter heterogeneities. For this reason, Voxel Based Morphometry (VBM) style analyses are never appropriate in T1w/T2w studies of the cerebral cortex.

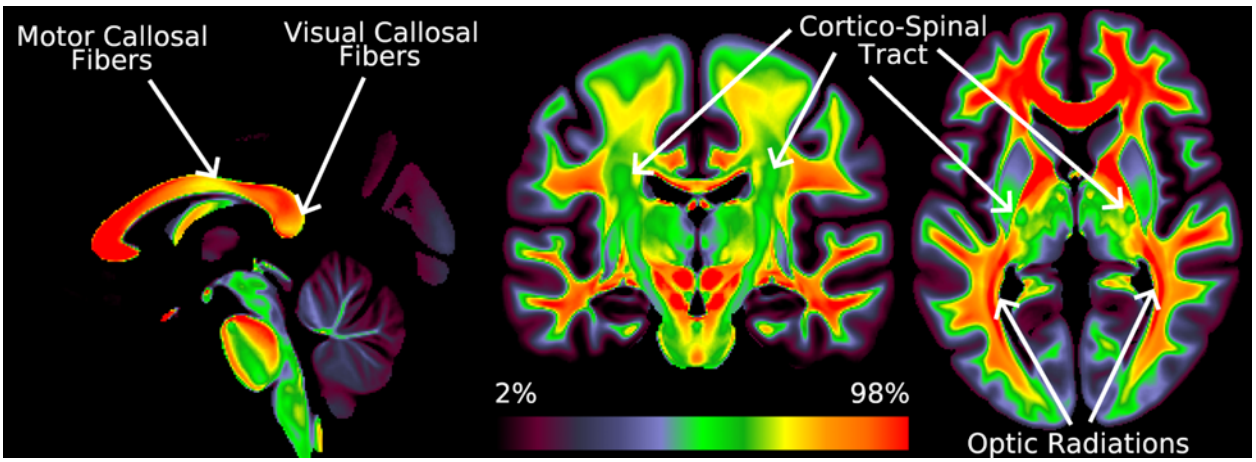

Figure 20 illustrates the group average T1w/T2w map after B1+ correction from the HCP-YA data.

There has been debate in the literature as to the optimal approach for non-invasive myelin mapping (Edwards et al., 2018; Campbell et al., 2018; Marques and Norris (2018); Cohen-Adad 2018; Fischl and Sereno 2018; Turner 2019; Shams et al., 2019). In particular, T1w/T2w myelin mapping has been criticized for potentially not being as specific for myelin as other measures such as quantitative T1, quantitative T2*, quantitative MT, or T2-based myelin water fraction (Edwards et al., 2018; Arshad et al., 2017; Hagiwara et al., (2018); Uddin et al., 2019). Indeed, T1w and T2w images vary due to a variety of effects within a given voxel, including lipid content, water content, and metal content. If these effects are strong, the T1w/T2w myelin measure will deviate from the actual myelin content in that location of the brain (e.g., vasogenic edema or iron deposition; Yarnykh 2016). The T1w/T2w measure has also been criticized relative to other measures such as quantitative T1 owing to difficulties in comparing across protocols and scanners (Edwards et al., 2018), though normalization methods have been developed (Ganzetti et al., 2014; Lee et al., 2015). Also importantly, those seeking to directly validate the T1w/T2w measure with histology in the same brains (Righart et al., 2017) have the important challenge that T1w contrast in particular changes early in the post-mortem time period (Fracasso et al., 2016; Shatil et al., 2018; Seifert et al., 2019), making post-mortem

T1w/T2w maps a poor surrogate for in vivo T1w/T2w maps (Patel et al., 2020). Despite these criticisms and the fact that quantitative approaches have made some improvements in image resolution and scanning time, T1w/T2w myelin mapping remains the most accessible approach to non-invasive myelin mapping as it relies on standard, widely available clinical 3D MPRAGE and SPACE sequences. Additionally, these high SNR sequences enable high spatial resolution (0.8mm isotropic or better) for precise mapping of the thin cerebral cortex at 3T in clinically reasonable acquisition times (c.f. Trampel et al., 2019; Dong et al., 2021). Finally, the T1w and T2w acquisitions are needed anyway for making high-quality surface reconstructions that are also used by other modalities such as fMRI or diffusion imaging in the HCP-Style approach to brain imaging analysis (Glasser et al., 2016b). Thus, we feel that it is reasonable to generate and analyze T1w/T2w myelin maps as accurately as possible. Investigators who wish to use more quantitative measures might additionally elect to acquire the additional measures for comparison that they consider important such as quantitative T1, quantitative T2*, quantitative magnetization transfer, or T2-based myelin water fraction (e.g., Cohen-Adad et al., 2012; Dick et al., (2012); Sereno et al., 2013; Cohen-Adad (2014); Sánchez-Panchuelo et al., 2014; Mangeat et al., 2015; Carey et al., 2018). Some high-field efforts enable acquiring more than one of these at the same time (Metere et al., 2017; Caan et al., 2019; Sun et al., 2020).

The need for B1+ correction of MRI-based myelin maps has been recognized by proponents of alternative myelin mapping technologies as well (e.g., Lutti et al., 2014; Yarnykh 2016; Marques et al., 2017; Hagberg et al., 2017; Haast et al., 2018). Interestingly, B1+ effects are also visible in certain diffusion measures such as NODDI (Zhang et al., 2012), likely through their effect on the efficiency of spin echo refocusing and thereby SNR (Fukutomi et al., 2018). Of the three approaches used for T1w/T2w myelin map B1+ correction in this study, the B1Tx approach used in the macaque seems to have the fewest artifacts and best overall performance, with a short acquisition time of additional scans (~ 2 min) and relies on a vendor supplied sequence. The only trick is that even and odd slices must be acquired in separate acquisitions to fill in the slice gap and then combined. The pseudo-transmit field approach is slightly less accurate but has the advantage that it allows use of images that are already being acquired in a typical HCP-style study. The ringing artifacts in the HCP-YA AFI acquisition could be avoided with adequate spoiling (Nehrke 2009), which would require careful study/scanner-specific piloting and AFI sequence adjustment. Other approaches for B1+ mapping exist including SA2RAGE (Eggenschwiler et al., 2012) and EPI-based approaches (Jiru and Klose 2006). Additionally, use of the T1w/T2w ratio at higher field strength (e.g. 7T and above) would require active B1+ shimming with parallel transmit to reduce the range of B1+ inhomogeneity to the more linear range seen at 3T to prevent loss of contrast and/or to satisfy power limitations. An alternative high-field approach involves the ratio of two gradient echo images with more similar B1+ effects (i.e. T1w/T2*w; De Martino et al., 2015; T1w/PDw; Van de Moortele et al., (2009)), though T2* has stronger b0 orientation dependence than T2 (Oh et al., 2013).

The approaches presented here enable correction of residual B1+ transmit field effects in T1w/T2w myelin maps, improving the accuracy of both group and individual maps. These new maps have advantages over the prior “MyelinMap_BC” correction method (Glasser et al., 2013) in that they (1) correct for the *symmetric* residual biases that were present in these maps (e.g.,

central to peripheral bias in intensity) and (2) correct individual biases related to head and body size that may correlate with variables of interest such as age (in development) or BMI without removing genuine individual differences in T1w/T2w ratio that may be of interest. The corrected ratio maps remain broadly applicable across species (just as the uncorrected or "MyelinMap_BC" corrected maps were), and may in fact enable improved identification of homologous brain regions, as a basis for more precise interspecies cortical surface registration. Prior literature using uncorrected myelin maps or “MyelinMap_BC” corrected myelin maps may need re-evaluation using the new approach depending upon whether results were affected by residual bias effects. Overall, we feel that these correction methods markedly improve the utility of T1w/T2w myelin maps for a wider range of studies when they are appropriate.

Acknowledgements

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Appendix A

Pseudocode for the pseudo-transmit (PT) field regularization optimization:

```
function optim_threshold {
  initialize T2* dropout threshold search range
  while T2* dropout threshold search range is larger than tolerance {
    pick new T2* dropout threshold and apply to PT data
    call optim_smoothing
    update search range
  }
  return best result
}

function optim_smoothing {
  initialize smoothing search range
  while smoothing search range is larger than tolerance {
    pick new smoothing amount and apply to PT data
    call optim_slope
    update search range
  }
  return best result
}

function optim_slope {
  initialize slope search range
  while slope search range is larger than tolerance {
    pick new slope value and use it as the correction for myelin (Eq. 5)
    compare corrected myelin to group template (Eq. 7)
    update search range
```

```
  }
  return best result
}
```

The `while` loop, `pick new value`, and `update search range` operations are all handled via a golden-section search (Kiefer 1953 PAMS).

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