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Test-retest reliability of cerebral blood flow in healthy individuals using arterial spin labeling: Findings from the EMBARC study

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Abstract

Introduction—Previous investigations of test-retest reliability of cerebral blood flow (CBF) at rest measured with pseudo-continuous Arterial Spin Labeling (pCASL) demonstrated good reliability, but are limited by the use of similar scanner platforms. In the present study we examined test-retest reliability of CBF in regions implicated in emotion and the default mode network.

Material and methods—We measured absolute and relative CBF at rest in thirty-one healthy subjects in two scan sessions, one week apart, at four different sites and three different scan

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platforms. We derived CBF from pCASL images with an automated algorithm and calculated intra-class correlation coefficients (ICCs) across sessions for regions of interest. In addition, we investigated site effects.

Results—For both absolute and relative CBF measures, ICCs were good to excellent (i.e. > 0.6) in most brain regions, with highest values observed for the subgenual anterior cingulate cortex and ventral striatum. A leave-one-site-out cross validation analysis did not show a significant effect for site on whole brain CBF and there was no proportional bias across sites. However, a significant site effect was present in the repeated measures ANOVA.

Conclusions—The high test-retest reliability of CBF measured with pCASL in a range of brain regions implicated in emotion and salience processing, emotion regulation, and the default mode network, which have been previously linked to depression symptomatology supports its use in studies that aim to identify neuroimaging biomarkers of treatment response.

Keywords

Arterial spin labeling; Reliability; fMRI; Cerebral blood flow

1. Introduction

Arterial spin-labeling (ASL) is a noninvasive magnetic resonance imaging (MRI) method that uses magnetically labeled arterial blood water as an endogenous tracer for cerebral blood flow (CBF) perfusion imaging [1]. Unlike positron emission tomography (PET) and single-photon emission computed tomography (SPECT), which require radioligands, ASL uses an endogenous marker that allows repeated use without radiation exposure [1]. Importantly, ASL shows high concordance with these more invasive neuroimaging techniques [2–4]. For example, resting CBF measures in cortical regions acquired with ASL and H₂O PET were similar in healthy subjects [3]. The non-invasive nature of ASL has facilitated its use in animal studies [5,6] and human studies with both healthy [7,8] and clinical populations [9–12], as well as pharmacological studies [13–16].

Given the promise of ASL imaging to help identify biomarkers in psychiatric disorders [1,9], it is important to demonstrate its reliability during repeated testing, both within and between scanner platforms. Several studies suggest that ASL is a reliable and reproducible measure of regional CBF over time using the same scanner [13,14,17–32], as well as across different sites using the same MRI scanner platforms [20,24,25,29,30]. Only two studies from the same group explored ASL reliability across two different scanner platforms, finding good reliability both at rest and during finger tapping in the motor cortex [33], and good reliability in mean global gray matter CBF at rest [34]. To date, only one study investigated ASL reliability across scanners of three different platforms [31], finding good reproducibility for most regions of interest studied, except posterior cingulate cortex, and concluding that differences in sequence parameters have greater impact on ASL reproducibility compared to differences in hardware/software across MRI scanner platforms. However, the study did not examine reliability across the three platforms as one unique group, but rather as two parallel investigations. Moreover, most other ASL reliability studies investigated CBF stability within a few hours [20,25,30,31] or within several days [20,24,25,29]. Overall, these studies

reported high reliability for scans completed within the same day, and a decrease in reliability after several days. Potential biomarkers of treatment-associated changes are more likely to yield clinical translation when measured after a longer interval, such as one week, rather than after few hours or days due to the time required for even the earliest treatment effects to emerge.

In the current study, we aimed to assess, for the first time, the test-retest reliability of ASL in healthy subjects scanned one week apart at four sites across three different MRI scanner platforms. This healthy cohort is part of a larger study, Establishing Moderators and Biosignatures of Antidepressant Response in Clinical Care (EMBARC), investigating treatment in major depressive disorder (MDD) [35]. We focused our analysis on brain regions previously reported to be dysfunctional in MDD, including regions implicated in emotion and salience processing, implicit and effortful emotional regulation, and regions of the default mode network [36]. Regions in these networks also show abnormalities in intrinsic (resting state) functional connectivity in MDD [36] and are therefore potential neural targets for antidepressant medications.

Based on previous findings indicating that intra-class correlation coefficients of ASL measures may differ between scans acquired a few hours apart from those acquired several days apart [18,19,21,23,25–29,32] and limited information on possible effects of data acquisition using different scans platforms, we aimed to determine the test-retest reliability of ASL measures in these networks across sites/platforms on scans acquired one week apart.

2. Material and methods

2.1. Establishing moderators and biosignatures of antidepressant response in clinical care (EMBARC) study

The EMBARC study (<http://embarc.utsouthwestern.edu>) [35] is a large multisite randomized controlled trial aiming to identify neuroimaging and other moderators and mediators of treatment response in depressed subjects with MDD, and also included recruitment of healthy control subjects to test reliability of the biomarkers, which is the focus of the current manuscript. All study subjects undergo neuroimaging assessment at baseline and after one week [35,37,38], which provides an ideal opportunity to acquire an ASL dataset across different sites using the same protocol.

2.2. Subjects

Forty healthy subjects were tested twice, one week apart, in one of four sites (10 different subjects per site). Subjects were between the ages of 18 and 65, fluent in English and scored lower than 8 on the Quick Inventory of Depressive Symptomatology-Self Report (QIDS-SR) -a self-report measure of depressive symptomatology [39]. Exclusion criteria included: lifetime history of Axis I mood disorder, anxiety disorder, psychotic disorder, or eating disorder; presence of epilepsy or other conditions requiring an anticonvulsant; presence of thyroid medication for hypothyroidism unless stable on thyroid medication for 3 months; history of a general medical condition that required hospitalization or deemed clinically significant; any clinically significant abnormal laboratory results and any clinically

significant neurological injury or deficit. The usual MRI exclusion criteria also applied, including being pregnant or breastfeeding, presence of ferromagnetic metal in one's body. The study was approved by the Institutional Review Board (IRB) at each site and all subjects signed informed consent.

2.3. Imaging methodologies

2.3.1. Data acquisition—Neuroimaging data were collected at four different sites using 3 Tesla scanners: Columbia University (CU – General Electric scanner), Massachusetts General Hospital (MGH - Siemens scanner), University of Michigan (UM – Philips scanner), and the University of Texas Southwestern (UTSW – Philips scanner). Pseudo-continuous arterial spin labeling (pCASL) protocols were optimized for performance and quality while maintaining similar parameters across sites (whole brain coverage with 29 axial slices, 35 pairs, 2D EPI acquisition, labeling duration = 1516 ms, post-labeling delay = 1500 ms, sequence duration = 5 min 12 s, voxel size: $3.4 \times 3.4 \times 5 \text{ mm}^3$, TR/TE = 4460/17 msec, FOV = $220 \times 220 \text{ mm}$, matrix = 64×64 , excitation flip angle 90°). In order to reduce variability across scanners the coil sensitivity was corrected by using a reference scan that was part of the scan protocol and the images were corrected for B1 heterogeneities. Labeling efficiency was set at 0.86 across subjects based on a group-average efficiency estimated in our previous study [40]. The labeling plane was positioned at 84 mm below the anterior-posterior line as proposed in the same study [40] and subsequently adapted by the ASL white paper [41]. Furthermore, the pCASL labeling module has been shown to be relatively robust to B0 inhomogeneities. T1-weighted Structural 3D axial images were acquired in the same session. The ASL sequence was acquired at rest with eyes open and fixated on a visual cue [42]. It was the last sequence in one-half hour scan session, which sets a high bar for reliability. However, staff received training to minimize subject head movement.

2.3.2. Arterial spin labeling preprocessing—Perfusion data were preprocessed and analyzed with the Statistical Parametric Mapping software, Version-8 (SPM8, Wellcome Department of Cognitive Neurology, UK; www.fil.ion.ucl.ac.uk/spm) implemented in Matlab7 (Math Works, Natick, MA) and a customized perfusion script to calculate CBF (<https://cfn.upenn.edu/perfusion/software.htm>, accessed February 17, 2017) [43]. For each subject, functional images were first realigned to correct for head movement. Next, the mean image was co-registered with the anatomical image and segmented. Perfusion weighted image series were then generated by pairwise subtraction of the label and control functional images, followed by conversion to absolute CBF image series based on a single compartment pCASL perfusion model [9,43,44]. M_0 representing equilibrium magnetization was used as a global index in the perfusion model, and was determined from a center slice at the level of the thalamus. The M_0 calculation was done based on a binary mask that was individually estimated from a center slice at the level of the thalamus. The binary mask was created using a flood fill algorithm that selected an area containing brain tissue and excluded skull and cerebrospinal fluid. Each mask was inspected for quality control. A mean absolute CBF (aCBF) image was generated for each subject, normalized to the Montreal Neurological Institute (MNI) template using parameters derived from the segmentation, resampled to $2 \times 2 \times 2 \text{ mm}$ and smoothed with full-width at half maximum 8 mm kernel.

Relative mean CBF (rCBF) was calculated by dividing the aCBF image by the global individual mean aCBF at each voxel.

2.3.3. Cerebral blood flow quantitative quality assessment—Several factors can contribute to poor data quality of the CBF image measured with ASL (for review, see (41,45)). To assess data quality, we used two quantitative measures generated by an automated calculation system to decrease subjective assessment bias: 1) signal-to-noise-ratio (SNR) between global gray and white matter; and 2) the mean frame-wise displacement calculated for each subject, based on Power's method, which provides a summary statistic of translational and rotational head movement [46]. Subjects with SNR values < 2.3 and > 3.4 and mean framewise displacement > 0.319 , which represented outliers for these measures as determined by descriptive analysis, were excluded from further analysis.

2.3.4. Regions of interest—Measures of mean aCBF and rCBF were extracted from regions implicated in emotion and salience processing (amygdala, striatum, insula, dorsal anterior cingulate cortex (ACC)), implicit and effortful emotional regulation (medial prefrontal cortex (mPFC), rostral ACC, ventrolateral prefrontal cortex (vlPFC), and dorsolateral prefrontal cortex (dlPFC)), and the default mode network (mPFC, posterior cingulate cortex). Cingulate regions were defined by Beckman clusters [47] and the ventral striatum was defined based on a neuroimaging meta-analysis [48,49]. Remaining regions were defined by the Wake Forest Toolbox PickAtlas Talairach Daemon template [50,51].

2.4. Analytical method

2.4.1. Intra-class correlation coefficients - across all sites—Within scanner reliability was assessed using intra-class correlation coefficients (ICCs) [52–54] in IBM SPSSv21, a measure widely used in neuroimaging research investigating arterial spin labeling reliability [14,17–23,25–30]. ICCs reflect the ratio of the between subjects variance (variance due to the fact that different subjects participated in the study) and the total variance (which includes within subjects effects and error). Thus, ICC estimates the percent of variance accounted for by between-subjects effects and reflects the consistency of measurements from session 1 to session 2. ICCs were calculated using a two-way mixed model single measure (ICC(3,1)) as in previous ASL reliability studies [14,27]. ICCs were categorized as: Poor (< 0.4), Fair ($0.4 \leq \text{ICC} < 0.6$), Good ($0.6 \leq \text{ICC} < 0.75$), and Excellent (≥ 0.75) [53]. We calculated ICCs for both aCBF and rCBF images; however, it should be noted that previous work showed higher reliability for rCBF measures [21,55,56]. A comparison of aCBF and rCBF was considered to be significant if confidence intervals (95%) for ICCs did not overlap with each other [18].

2.4.2. Intra-class correlation coefficients - leave-one-site-out—We repeated the above analysis four times, each time leaving one site out to investigate the effect of site. Site was considered to have a significant effect if confidence intervals (95%) for ICCs for any of the sites removed did not overlap with ICCs for the other three sites [18]. We calculated ICCs for both, aCBF and rCBF.

2.4.3. Bland-Altman assessment across sites – gray matter—A Bland-Altman assessment for agreement was used to compare whole-brain gray matter aCBF at each time point and site. A range of agreement was defined as mean bias \pm 2 SD [57]. Proportional bias is present when the difference between measures is a function of the mean of the measures. A multiple linear regression analysis with the difference in CBF measure as the dependent variable and the mean CBF measure and site (transformed into dummy variables with site 3 as reference) as independent variables was used to test whether the model coefficient for mean CBF equaled zero. Rejection of that null hypothesis would be consistent with the presence of proportional bias ($p < 0.05$). Gray matter aCBF in young healthy subjects is considered normal between 40 and 100 mL/100 g/min [41].

2.4.4. Assessing the effect of scanner site—To examine the effect of scanner site on whole-brain gray matter aCBF across sessions, we conducted a repeated-measures analysis of variance (ANOVA) with session as a within-subject variable and scanner site as a between-subjects variable. P values below 0.05 were considered significant.

3. Results

3.1. Demographic information and quantitative quality assessment

Six subjects were excluded because of low data quality (two subjects had gray/white matter SNR value > 3.4 and two subjects had values lower or equal to -2.3 ; and three subjects had mean framewise displacement > 0.32). One of these six subjects failed both data quality measurements. Four subjects did not complete both sessions. One subject did not complete both sessions and had low data quality. This yielded a final sample of $N = 31$ (Supplemental Fig. 1). There were no differences in age, gender, and years of education across sites (all $ps > 0.1$; Table 1). However, we found a trend towards significance for marital status ($p = 0.054$) and ethnicity ($p = 0.06$); and a significant difference for race ($p = 0.049$) across sites.

3.2. Absolute and relative cerebral blood flow reliability across all sites

The majority of brain regions had good to excellent aCBF and rCBF reliability, and six regions had fair reliability. The subgenual anterior cingulate and ventral striatum had the highest aCBF and rCBF reliability (excellent - both ICCs > 0.8) and the right amygdala had the lowest aCBF and rCBF reliability (fair - ICC = 0.4 Table 2).

The majority of brain regions had comparable aCBF and rCBF reliability. Four regions had numerically different ICCs between aCBF and rCBF, however, all had a difference lower than 0.2, and overlapping confidence intervals.

3.3. Absolute cerebral blood flow reliability - leave-one-site-out validation

The leave-one-site out analysis did not show a significant effect of site on the reliability of aCBF (Supplemental Table 1). For all regions, the confidence intervals of each of the four pair comparisons overlapped.

3.4. Relative cerebral blood flow reliability - leave-one-site-out validation

The leave-one-site out analysis did not show a significant effect of site on the reliability of rCBF (Supplemental Table 2). For all regions, the confidence intervals of each of the four pair comparisons overlapped.

However, it should be noted that the right amygdala reliability for absolute and relative CBF decreased to 0.02 when site 1 was left out of the analysis, although confidence intervals continued to overlap.

3.5. Bland-Altman assessment across sites – gray matter

Mean whole-brain gray matter aCBF across sites was 45.9 mL/100 g/min (SD = 10.8).

Bland–Altman plots together with 95% confidence interval (CI) of CBF measurements from gray matter in the 31 subjects subdivided by each of the four sites are shown in Fig. 1. Gray matter CBF difference plotted against mean gray matter CBF were randomly distributed and showed no dependence on mean gray matter CBF, as can be seen in the Bland–Altman plots shown in Fig. 1. Independent variables (site and mean aCBF measure) assessed with multiple linear regression did not predict change in the dependent variable, gray matter aCBF difference. Therefore, no proportional bias was found across sites ($p > 0.2$ for gray matter aCBF).

3.6. Assessing the effect of scanner site on mean aCBF levels - repeated-measures ANOVA

We observed no significant effect of session ($F(1,27) = 0.1$, $p = 0.8$) or session by site interaction ($F(3,27) = 0.3$, $p = 0.9$) for whole-brain gray matter aCBF. However, there was a significant scanner site effect ($F(3,27) = 6.3$, $p = 0.002$) with site 3 having lower whole brain gray matter aCBF relative to site 1 ($p = 0.034$) and 4 ($p = 0.003$); and a trend for lower aCBF relative to site 2 ($p = 0.08$) (Fig. 2).

4. Discussion

Resting aCBF and rCBF measured with ASL were comparable and demonstrated good to excellent reliability between sessions for the majority of brain regions examined and were stable across sites. In addition, there was no proportional bias within each site for whole-brain gray matter aCBF and the precision of the measures was not different across sites. However, one site had significantly lower aCBF relative to the other sites. These findings support the potential utility of pCASL measures in longitudinal studies, but suggest caution in multi-site studies.

One advantage of CBF measured with ASL is that it allows for calculation of an absolute quantification of CBF. Previous studies highlighted the greater reliability of relative over absolute CBF values when using ASL [21,56]. Relative measures can factor out global intra-subject CBF variation which is a significant source of noise. In the current study, aCBF and rCBF were highly comparable, and were measured with pseudo-continuous ASL, as per the current recommendation for this type of sequence [41]. Furthermore, absolute quantification can be used as a marker for pharmacological activity [13–16], which is useful for both

longitudinal and cross-sectional studies investigating psychopathology such as MDD, which is the focus of the EMBARC study.

Our findings are consistent with previous studies reporting good to excellent test-retest reliability of CBF measures one week apart in the same scanner [13,23,27,32]. However, ASL reliability studies have typically shown decreasing reliability as time between scan sessions increases [14,18–20,23,27,30]. For example, Chen et al. reported coefficients of variability < 11% (i.e. good reliability) within one hour apart and < 15% one week apart in different cerebral artery territories [23]. Similarly, Gevers et al. found excellent ICCs (> 0.85) at whole brain level and in different cerebral artery territories during the same session acquisition and a reduction in ICCs values after one to three weeks (> 0.7) [18]. Our findings indicate that ASL measures have high test-retest reliability, as shown by image acquisition one week apart. We show that this is also the case during the four leave-one-site-out analyses. These findings suggest that variations in ASL measures over time are probably more related to errors caused by subject repositioning, data processing, and physiological fluctuations, rather than scanner instability.

Furthermore, we found that the precision of the whole gray matter aCBF measure between baseline and week 1 was not predicted by a site effect analyzed with multiple linear regression, and that the leave-one-site-out analyses revealed no effect of site on test-retest reliability of aCBF and rCBF measures, in accordance with previous ASL studies examining reliability across different sites, but using the same MRI platforms [20,24,25,29]. Petersen et al. published the largest ASL reliability study to date, with approximately one hundred ninety one subjects in 28 sites, each using a 3 Tesla Philips MRI scanner. Subjects were scanned twice, two weeks apart [20], with reasonable reproducibility across sites, but with a wide range of values (gray matter ICC ranged between 0.07 and 0.81), which the authors explained by a noticeable correlation between mean CBF within the sites and their data acquisition success rate ($r = 0.66$). Furthermore, Gevers et al. reported whole-brain pCASL perfusion varying < 20% over repeated measurements 1–3 weeks apart within the same six subjects across three 3 Tesla Philips MRI scanners [24]. Intra-scanner reliability has been shown to be typically higher than inter-scanner reliability, but the latter was still shown to be high (visual cortex ICC > 0.7 and several regions of interest ICCs between 0.4 and 0.97) across two 3 Tesla GE scanners in two studies by the same group with twelve and eight subjects examined 10–15 days apart [25,29]. Two studies from the same group explored ASL intra and inter scanner reliability across GE and Philip platforms in twenty-two subjects two weeks apart from each session. The first study found low coefficient of variation in global gray matter CBF at rest within platforms (GE: 10.8%; Philips: 11.3%) and between platforms (13.6%) [34]. The second study found good reliability at rest (GE: IC-C = 0.69, Philips: ICC = 0.77) and during finger tapping task (GE: ICC = 0.66, Philips: ICC = 0.75) in the motor cortex. Moreover, the inter scanner reliability was also good at rest (ICC = 0.69) and during motor activation (ICC = 0.61) [33]. Only one small study explored more three 3 Tesla MRI platforms – GE, Philip and Siemens [31]. The authors compared similar pCASL acquisition parameters within the same day in two parallel studies, one with eleven subjects, and a second with fourteen subjects (eleven subjects participated in both studies). The authors found high inter-scanner reliability, albeit with coefficients of variation ranging from 11.2% to 46% depending on the gray matter region, leading the authors to

conclude that it is possible to acquire comparable cerebral CBF maps on scanners of different platforms.

Our current findings add to the literature by demonstrating high reliability of ASL measures across four sites with three different 3 Tesla MRI platforms, using a larger sample and greater time interval between scans than a previous study that investigated inter scanner reliability of ASL measures across three different scanner platforms [27]. Furthermore, mean whole-brain gray matter aCBF for all subjects across sites was between 40 and 100 mL/100 g/min, which is considered to be within the normal range for healthy subjects. Despite similar precision across sites as discussed in the previous paragraph, direct comparison using repeated measures ANOVA among the four different sites and three different scanner platforms revealed significantly lower whole-brain aCBF in one of the sites (i.e. main effect of site but no effect of session nor session by site interaction), which could be explained by a within group difference rather than a scanner site difference since the subjects were not the same across sites. Furthermore, the large range of reliability values across different brain regions, underscores the need for further optimization of ASL acquisition parameters such as inversion efficiency, inversion time, post labeling delay time, background suppression, readout type, as well as SNR, and movement control [1,41,45].

The subgenual ACC and ventral striatum had the highest reliability values. These regions are implicated in the psychopathology of depression [36]. For example, a previous ASL study characterizing CBF in unipolar depression relative to healthy controls found decreased CBF in the subgenual ACC [12]. Moreover, CBF measures with ASL in subgenual ACC have been shown to differentiate unipolar and bipolar depression using pattern recognition analysis [9], and elevated ventral striatal activation has been implicated in bipolar disorder [58,59]. Differentiating between unipolar and bipolar depression is of key importance because it determines the choice of treatment [60]. Studies to ascertain the utility of pCASL for differentiating between these two disorders would thus be instructive.

SNR is a widely used measure to assess data imaging quality (for a review of advancements in ASL imaging parameters designed to optimize signal see (1,41,45)). The pCASL employed in this study has been shown to yield superior SNR compared with pulsed ASL, and is now becoming the sequence of choice to measure CBF [23,24,41,45]. In addition, earlier studies of BOLD sequences showed that movement and other imaging artifacts can significantly impact data quality [61,62]. We assessed data quality using two quantitative measures: gray/white matter SNR and framewise displacement.

Our sample size was relatively small relative to reliability studies of BOLD sequences; however, it is the largest test-retest reliability study of ASL across MRI platforms.

5. Conclusions

This is the first study to examine the test-retest reliability of ASL measures in healthy subjects across multiple sites and MRI platforms. We demonstrated good to excellent test-retest reliability of CBF measured with pCASL in a range of brain regions implicated in emotion and salience processing, emotion regulation, and the default mode network, which

have been previously linked to MDD symptomatology. Given its high reliability, ASL is a promising neuroimaging technique for longitudinal studies of psychiatric disorders and a useful tool to identify biomarkers reflecting neuropathophysiologic processes and neural predictors of treatment response.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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Conflict of interest statements

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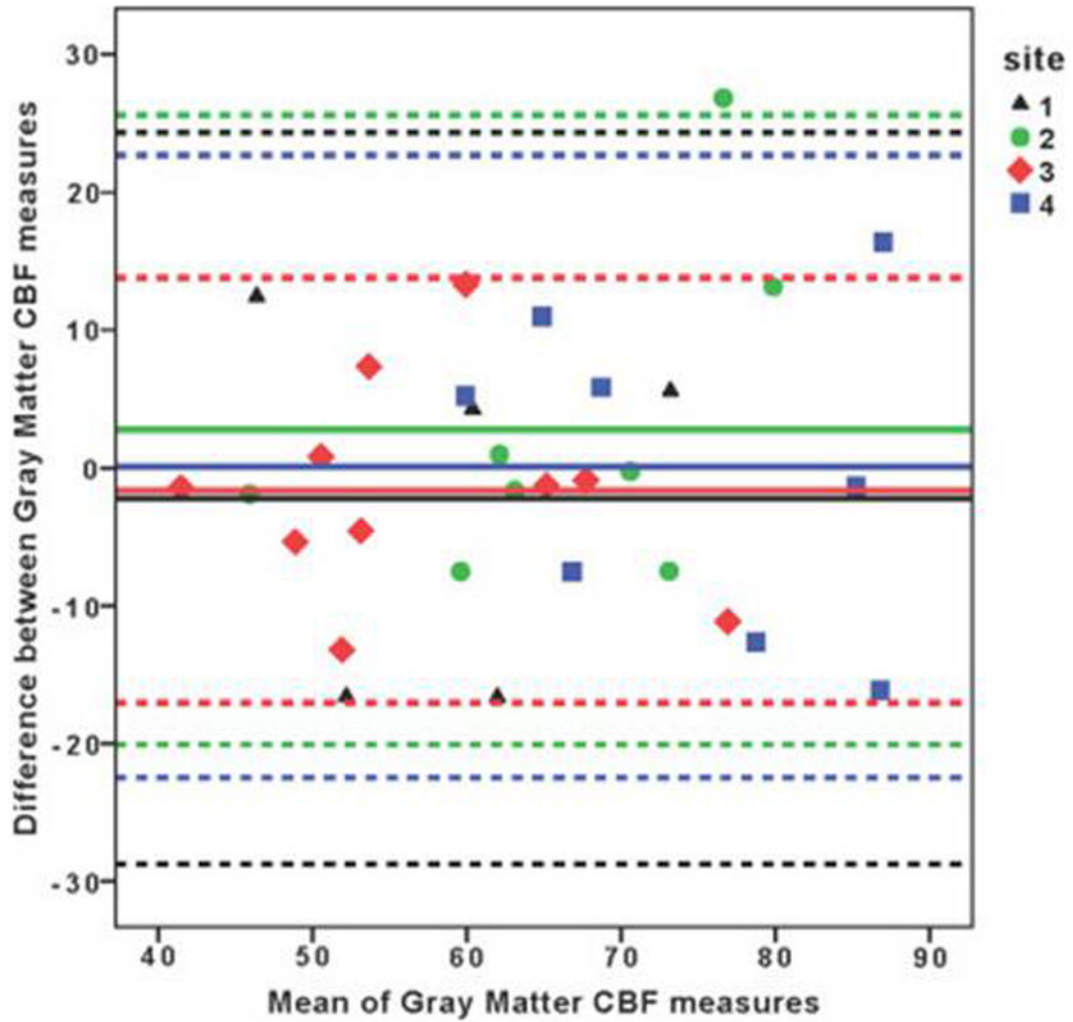


Fig. 1. Bland-Altman plot of repeated CBF measurements (baseline and week 1) across four sites in absolute whole-brain gray matter CBF (mL/100 g/min). The Bland-Altman plot represents intrascan aCBF differences plotted against mean aCBF. Intrascan aCBF differences are randomly distributed and are not dependent on mean aCBF. Multiple linear regression demonstrated no similarity difference between sites. The solid middle line denotes the bias from zero, which represents absolute no difference between baseline and week 1 measures. The dotted top and bottom lines represents limits of agreement (i.e. bias \pm 1.96 \times standard deviation). Different colors and symbols represent each of the four sites. (For interpretation of the references to color in this figure legend, the reader is referred to the web version of this article.)

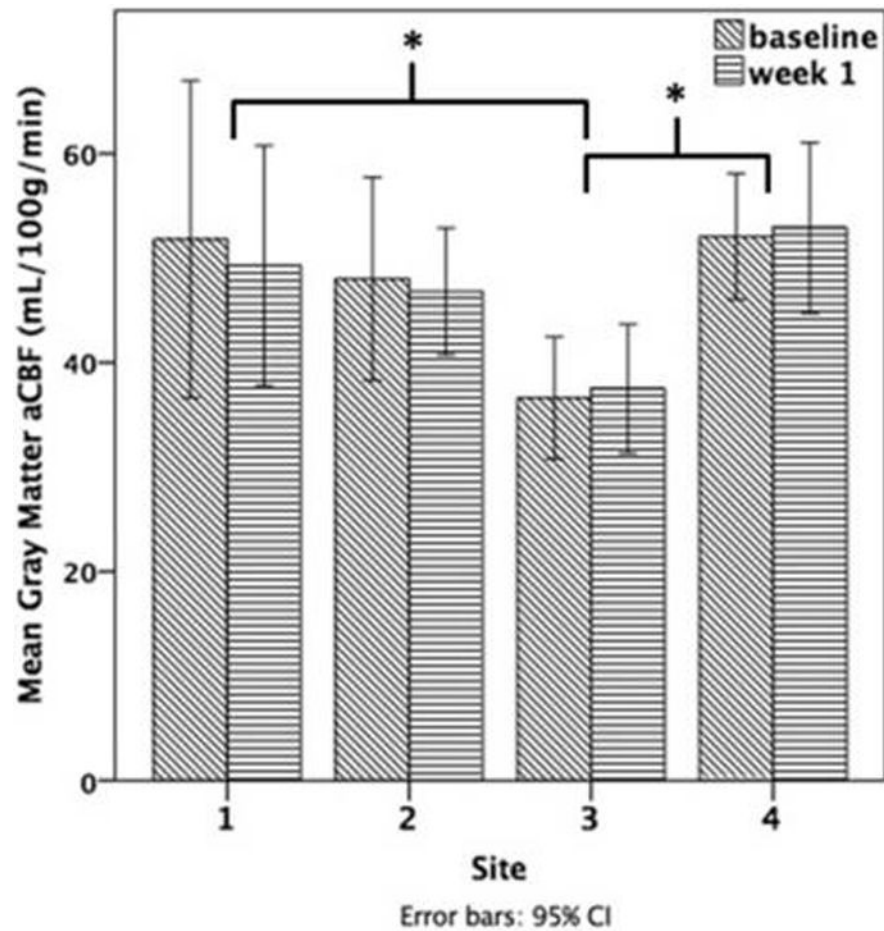


Fig. 2. Bar graph representing mean absolute whole-brain gray matter CBF (mL/100 g/min) at baseline and week 1 for each site. Each cluster represents a specific site. Each bar constitutes either baseline (diagonal lines) or week 1 (horizontal lines) mean whole-brain gray matter aCBF. Asterisks denote a significant difference ($p < 0.05$) between sites resulting from a mean effect of site in the repeated measures ANOVA.

Table 1

Demographic information across sites.

Site	1 (N = 5)		2 (N = 8)		3 (N = 10)		4 (N = 8)		Total (N = 31)	
	Mean	SD	Mean	SD	Mean	SD	Mean	SD	Mean	SD
Age	30.0	8.9	45.6	19.1	31.9	11.0	43.4	16.8	38.1	15.6
Year of education	13.6	3.0	17.8	8.7	14.9	2.4	16.4	2.3	15.8	4.9
Gender	N = 5	%	N = 8	%	N = 10	%	N = 8	%	N = 31	%
Male	1	20.0%	1	12.5%	5	50.0%	3	25.0%	9	29.0%
Female	4	80.0%	7	87.5%	5	50.0%	6	75.0%	22	71.0%
Race	2	40.0%	4	50.0%	7	70.0%	6	75.0%	19	61.3%
African American	0	0.0%	2	25.0%	3	30.0%	2	25.0%	7	22.6%
Asian	1	20.0%	2	25.0%	0	0.0%	0	0.0%	3	9.7%
Other	2	40.0%	0	0.0%	0	0.0%	0	0.0%	2	6.5%
Hispanic	3	60.0%	7	87.5%	10	100.0%	8	100.0%	28	90.3%
Yes	2	40.0%	1	12.5%	0	0.0%	0	0.0%	3	9.7%
Marital Status	5	100.0%	4	50.0%	8	80.0%	3	37.5%	20	64.5%
Single	0	0.0%	1	12.5%	2	20.0%	4	50.0%	7	22.6%
Married	0	0.0%	3	37.5%	0	0.0%	1	12.5%	4	12.9%
Divorced	0	0.0%	0	0.0%	0	0.0%	0	0.0%	0	0.0%

Age F = 2.1; $p = 0.1$; Years of Education F = 0.9; $p = 0.5$; Gender $\chi^2 = 3.5$; $p = 0.3$; Race $\chi^2 = 17$; $p = 0.05$; Hispanic $\chi^2 = 7.3$; $p = 0.06$; Marital Status $\chi^2 = 12.4$; $p = 0.054$.

Table 2

Absolute and Relative Cerebral Blood Flow Reliability Across All Sites.

	Absolute CBF			Relative CBF		
	ICC	95% CI		ICC	95% CI	
		Lower bound	Upper bound		Lower bound	Upper bound
sgACC	0.81	0.64	0.90	0.81	0.64	0.90
rACC	0.76	0.56	0.88	0.76	0.56	0.88
dACC	0.78	0.59	0.89	0.78	0.58	0.89
mCC	0.69	0.45	0.84	0.69	0.45	0.84
pCC	0.67	0.42	0.83	0.67	0.42	0.83
Left amygdala	0.59	0.30	0.78	0.59	0.30	0.78
Right amygdala	0.42	0.09	0.67	0.42	0.09	0.67
Left dlPFC	0.68	0.43	0.83	0.68	0.43	0.83
Right dlPFC	0.69	0.45	0.84	0.69	0.45	0.84
Left vlPFC	0.57	0.28	0.77	0.57	0.28	0.77
Right vlPFC	0.54	0.24	0.75	0.54	0.23	0.75
Left caudate body	0.66	0.41	0.82	0.46	0.14	0.70
Right caudate body	0.49	0.18	0.72	0.68	0.44	0.83
Left insula	0.68	0.43	0.83	0.68	0.43	0.83
Right insula	0.59	0.30	0.78	0.59	0.30	0.78
Left mPFC	0.71	0.47	0.85	0.71	0.47	0.85
Right mPFC	0.69	0.45	0.84	0.69	0.45	0.84
Left putamen	0.64	0.37	0.81	0.64	0.37	0.81
Right putamen	0.64	0.37	0.81	0.65	0.39	0.81
Left VS	0.80	0.63	0.90	0.72	0.50	0.86
Right VS	0.72	0.50	0.86	0.80	0.63	0.90

ICC: intraclass correlation coefficient; CI: confidence interval; no color = poor reliability (<0.40); red color = fair reliability (0.40 < ICC < 0.60); blue color = good reliability (0.60 < ICC < 0.75); green color = excellent reliability (>0.75); ACC: anterior cingulate cortex; sgACC: subgenual ACC; rACC: rostral ACC; dACC: dorsal ACC; CC: cingulate cortex; mCC: middle CC; pCC: posterior CC; PFC: prefrontal cortex; dlPFC: dorsolateral PFC; vlPFC: ventrolateral PFC; mPFC: medial PFC; VS: ventral striatum.