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## Diffusion Tensor Imaging Based Tissue Segmentation: Validation and Application to the Developing Child and Adolescent Brain

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### Abstract

We present and validate a novel diffusion tensor imaging (DTI) approach for segmenting the human whole-brain into partitions representing grey matter (GM), white matter (WM) and cerebrospinal fluid (CSF). The approach utilizes the contrast among tissue types in the DTI anisotropy vs. diffusivity rotational invariant space. The DTI-based whole-brain GM and WM fractions (GMf and WMf) are contrasted with the fractions obtained from conventional magnetic resonance imaging (cMRI) tissue segmentation (or clustering) methods that utilized dual echo (proton density-weighted (PDw), and spin-spin relaxation-weighted (T2w) contrast, in addition to spin-lattice relaxation weighted (T1w) contrasts acquired in the same imaging session and covering the same volume. In addition to good correspondence with cMRI estimates of brain volume, the DTI-based accurately depicts expected age vs. WM and GM volume-to-total intracranial brain volume percentage trends on the rapidly developing brains of a cohort of 29 children (6–18 years). This approach promises to extend DTI utility to both micro and macrostructural aspects of tissue organization.

### Keywords

DTI; Segmentation; Icosa21; Child Brain Development; Meta Analysis

### Introduction

Diffusion tensor imaging (DTI) provides noninvasive orientation based tissue contrast mechanisms and complements relaxation based conventional MRI (cMRI) methods. DTI scalar and orientation maps can provide information about the tissue microstructural integrity in both health and disease. The DTI-derived rotational invariant metrics, including the local tissue eigenvalues and the corresponding anisotropy indices and mean diffusivity, have been utilized in several applications in clinical psychology, neuropsychiatry, neurology, and radiology (Pierpaoli et al, 1996; see Hasan, 2006 for extensive review of literature). In particular, DTI applications to the developing child brain in health and disease offers a unique opportunity to

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understand the organization of the brain during transition to adolescence and adulthood (Simon et al., 2005). Despite known limitations, at the current stage of DTI technology it seems that echo-planar imaging (EPI) is the mainstay in multidimensional MRI acquisitions that include diffusion-weighted imaging (DWI) and its DTI extension, relaxometry, functional and perfusion MRI (Le Bihan et al., 1995). EPI time-efficiency, motion insensitivity, reduced specific absorption rates (SAR) and recent improvements (Bammer et al., 2002; Wang et al., 2005; Jaerman et al., 2005) make it attractive for mapping brain-structure and function relations. A typical whole-brain high spatial and angular resolution DTI acquisition can be accomplished in under 7 minutes at 3.0 T. Advances in high field technology, parallel imaging, novel coil design and image quality improvements (gradient hardware) promise to make DTI extremely useful for imaging neonates, infants, children and the elderly in both health and disease where exam time is crucial (Dubois et al., 2006; Gupta et al., 2005; Maas et al., 2005; McKinstry et al., 2002).

The application of conventional MRI contrast methods to brain tissue segmentation is a well-established set of procedures that provides regional (Shaw et al., 2006; Sowell et al., 2003) or global surrogate markers of development (Castellanos, et al., 2002; Giedd et al., 1999; Matsuzawa et al., 2001; De Bellis, 2001), natural aging (Courchesne et al., 2000; Good et al., 2001; Sowell et al., 2003; see Table 1), and their interactions with neurodevelopmental and neurodegenerative pathologies (Bartzokis et al., 2002; Brunetti et al., 2000; Castellanos et al., 2002; Eliez et al., 2001; Prinster et al., 2006; Simon et al., 2005; Zhang and Sejnowski, 2005; Wilde et al., 2005). To attain soft brain tissue segmentation, multispectral data volumes are acquired in the same session and coregistered to maximize the information content and separability of tissue clusters (Pham et al., 2000; Good et al., 2001). Tissue segmentation using cMRI requires radio frequency field inhomogeneity correction of the masked and co-registered volumes (Rajapakse et al., 1996a; Good et al., 2001; see Table 1). MRI contrast-to-noise ratio per unit scanning time between tissue types reflects a complex interplay and compromise between intrinsic tissue and extrinsic parameters (Le Bihan, 1995; Conturro et al., 1995). However, confounding variables such as age can make cMRI protocols harder to maintain if a wide age span is included (Chard et al., 2002; Guttmann et al., 1998; Inglese and Ge 2003; Matsuzawa et al., 2001).

There are many published works on the natural aging of the human brain. Due to the use of continuously improving acquisition, spatio-temporal parameters and processing paradigms, it is noteworthy that there are discrepancies in the published works on the developing healthy brain using cMRI (see Table 1). Comparison across these studies can be difficult as the brain coverage may vary and many studies have used different slice thickness/skip and acquisition parameters. Nonetheless, these studies consistently document age-related changes in the regional tissue composition of the brain. In general, to account for variability in the healthy and normal human brain size due to skull size and gender, it is widely recommended to scale the absolute volume of tissues with respect to the intracranial volume (ICV) (Good et al., 2001; Rajapakse et al., 1996a,b). For example, there are at least two major cross-sectional studies that covered the entire human life span and provided consistent estimates of the trends observed in the developing human brain from childhood till late adulthood (Courchesne et al., 2000; Sowell et al., 2003). Consistent with other reports (see Table 1), the whole-brain grey matter fraction (GMf) decreases with age and white matter fractions (WMf) increases during childhood, reflecting the continued myelination and axonal elimination or caliber size changes with age (see Table 1 for more details). However, as Table 1 shows there is considerable variability in the tissue fractions yielded by the cMRI methods, which undoubtedly due to variations in segmentation procedures, imaging acquisition parameters and other factors (see Table 1).

The application of DTI contrast to soft brain tissue clustering was initially described by Pierpaoli et al. (1996). More recently, DTI methods have been applied to segment the thalamic nuclei (Wiegell et al. 2003), the thalamocortical projections (Behrens et al., 2003), the fetal brain (Maas et al. 2004), and the human corpus callosum (Hasan, 2006). Many recent DTI-based clustering strategies have also been described (Lenglet et al. 2006; Liu et al. 2006; Wang and Vemuri, 2005). The incorporation and fusion of cMRI contrasted maps, DWI, DTI-derived metrics “mean diffusivity and anisotropy” in tissue segmentation and applications have also been described on humans (Jones et al. 1999; Hadjiprocopis et al., 2005) and rodent brains (Ali et al., 2005; Carano et al. 2000).

In this paper, we present and validate an application of whole-brain DTI to tissue segmentation of the developing healthy human child and adolescent brains; thus, extending the benefits of DTI to include the provision of regional and global atrophy micro- and macrostructural metrics that can be used in combination with region-of-interest and fiber based tracking methods to monitor the brain-behavior relations. We validate the DTI-based results against estimates from cMRI and against age to determine if the expected patterns of developmental change emerge.

## Materials and methods

### Participants

The participants included 15 girls and 14 boys (age range, mean  $\pm$  SE = 7–17,  $11.4 \pm 3.1$  years); 25/29 children were right-handed. The male and female groups did not differ in age ( $p=0.35$ ). All children were primarily English-speaking, were identified as neurologically normal by review of medical history and were medically stable at the time of the assessments. All scans were read as “normal” by a board certified radiologist (LK). Written informed consent was obtained from the guardians and adolescents and assent from the children participating in these studies per the University of Texas Health Science Center at Houston institutional review board regulations for the protection of human research subjects.

### MRI Data Acquisition

We acquired whole-brain data using a Philips 3.0 T Intera system with a SENSE parallel imaging receive head coil. The MRI protocol included (a) conventional MRI (3d spoiled gradient-echo, field-of-view= $240 \times 240$  mm<sup>2</sup> (isotropic voxel size = 0.9375 mm) and (b) 2d dual spin-echo images  $T_{E1}/T_{E2}/T_R=10/90/5000$  ms, in the axial plane (3mm slice thickness, square field-of-view= $240 \times 240$  mm<sup>2</sup> @ 44 sections) (c) phase-sensitive MRI in the sagittal and axial planes, in addition to a matching prescription of axial diffusion-encoded data as described below.

### Diffusion Tensor Acquisition

The diffusion-weighted data were acquired using a single-shot spin echo diffusion sensitized echo-planar imaging (EPI) sequence with the balanced *Icosa21* encoding scheme (Hasan, 2006; Hasan and Narayana 2003; 2006), a diffusion sensitization of  $b=1000$  sec.mm<sup>-2</sup>, a repetition and echo times of  $T_R=6.1$  s,  $T_E=84$  ms, respectively. EPI image distortion artifacts were reduced by using a SENSE acceleration factor or k-space undersampling of R of two (Bammer et al., 2002; Hasan and Narayana, 2006; Jaermann et al., 2004). The slice thickness was 3 mm with 44 axial slices covering the whole-brain (foramen magnum to vertex), a square field-of-view= $240 \times 240$  mm<sup>2</sup>, and an image matrix of  $256 \times 256$  that matched the 3d-SPGR (spoiled gradient or field echo) and 2d conventional MRI dual spin echo sequences described above. The number of non-diffusion weighted or  $b \sim 0$  magnitude image averages was 8; in addition, each encoding was repeated twice and magnitude-averaged to enhance the signal-to-noise ratio (SNR) (Conturo et al., 1995); thus, effectively 50 images were acquired for each of the 44 axial sections to cover the whole-brain. The total DTI acquisition time was

approximately 7 minutes and resulted in SNR-independent DTI-metric estimation (in brain parenchyma --grey and white matter tissue); SNR (b~0) ~ 50–60, SNR (DWI) ~ 20–30) which gave reproducible results.

### Conventional MRI Data Processing and Tissue Segmentation

The whole-brain cMRI data (i.e 3d SPGR and 2d dual-echo) were converted into ANALYZE formatted volumes (T1w, PDw & T2w) and skull stripped using MRICro software package which has the brain extraction tool. Using SPM99 (see Good et al., 2001), the PDw and T2w images were coregistered and resliced to their corresponding T1w volumes. Following recommendations by Simon et al. (2005), the Cincinnati Children's Hospital Medical Center, (CCHMC, Cincinnati, OH) pediatric brain template was obtained and the T1w images were spatially normalized to the same stereotactic space by registering each of the images to the CCHMC template. Average PDw and T1w brain templates were created in SPM2 as suggested by Good et al. (2001). Following the template formation, automated tissue segmentation was performed on the PDw and T2w images in addition to the T1w using their respected templates. Tissue volumes were calculated from the segmented volumes as suggested elsewhere (Good et al, 2001;Simon et al., 2005).

### DTI Data Processing

Although the DTI raw images were acquired with fat suppression, all image volumes were semi-automatically stripped to remove non-parenchymal tissue. Diffusion-weighted data were distortion-corrected using the mutual information maximization approach (Netsch and Van Muiswinkel, 2004). The details of the DTI processing are provided elsewhere (Hasan, 2006).

**DTI-derived Metrics**—The DTI-derived rotationally-invariant metrics used are the parallel ( $\lambda_{\parallel}=\lambda_1$ ) and transverse diffusivity ( $\lambda_t$ ). The transverse (perpendicular) diffusivity was defined as the mean of the minor (third) and medium (second) eigenvalues ( $\lambda_t = \lambda_{\perp} = (\lambda_2+\lambda_3)/2$ ) (Beaulieu, 2002;Hasan and Narayana, 2006;Song et al., 2005). The fractional anisotropy (FA) and mean diffusivity ( $D_{av}$ ) are computed using the eigenvalues as:

$$FA = \sqrt{\frac{1}{2} \frac{(\lambda_1 - \lambda_2)^2 + (\lambda_2 - \lambda_3)^2 + (\lambda_1 - \lambda_3)^2}{\lambda_1^2 + \lambda_2^2 + \lambda_3^2}} \quad [1]$$

$$D_{av} = \frac{\lambda_1 + \lambda_2 + \lambda_3}{3} = \frac{\lambda_{\parallel} + 2 \times \lambda_{\perp}}{3} \quad [2]$$

Notice that FA is scale-independent (i.e.  $FA(c\lambda_1; c\lambda_2 c\lambda_3) = FA$ ). The apparent diffusivities are generally estimated by log-linear algebraic transformations (Basser et al., 1994) or by the moments of the log-transformed apparent diffusion coefficients (Ackerman, 2003;Arfanakis et al, 2002;Frank, 2001;Hasan and Narayana, 2003). This log-linear decoupling of intensity makes anisotropy indices more immune to local intensity modulations due RF field excitation inhomogenities (Bammer et al., 2002).

### Region-of-Interest and DTI Feature Space Formation

The regions-of-interest derived from the analysis represented 50 “normally appearing” white matter structures, grey matter structures and ventricular CSF. The structures included the right-left (contralateral) caudate head, putamen, internal capsule (anterior, external, posterior limb), corpus callosum segments (see Figure 1), corticospinal tract, forceps minor and major, inferior and superior longitudinal fasciculi and arcuate fasciculus and other grey (cortical, hippocampal) and white matter structures (cerebellar, peduncular). The compact white matter structures belonged to three major categories: commissural, association, and projection fibers

that are involved in inter and intra-hemispheric communication. The ROI placement and fiber tracking procedures were done by trained raters and used anatomical landmarks from the conventional MRI sets (sagittal phase-sensitive inversion recovery). The ROI placement procedure was supervised by a radiologist and used a sophisticated three-dimensional system that fused DTI maps with conventional high resolution anatomical MRI (Hasan, 2006); this procedure was shown to help reduce partial averaging artifacts due to CSF. From these ROIs, a feature space was created to determine the tissue classification thresholds (see Figure 1).

### **DTI based Tissue Segmentation**

The collected ROI DTI metrics and their corresponding errors were further used in a multidimensional supervised and trained feature space initiated clustering procedure to segment the whole-brain (cerebrum, brain stem and cerebellum) tissue into three tissue types: white matter, grey matter and cerebrospinal fluid (CSF). The method uses the contrast in FA maps between CSF, white matter, and grey matter and the cluster separability and discriminability of white matter and grey matter based on the principal diffusivity indices. The CSF was segmented based on its high diffusivity and low anisotropy (Pierpaoli et al., 1996; Hasan, 2006).

### **Statistical analysis**

Group mean comparisons were made using the t-test for unpaired groups: within group comparisons were conducted using paired t-tests. Alpha levels were adjusted for the number of dependent univariate tests  $M=3$  (GM, WM & CSF) ( $p < 0.05/M$ ). Correlations with age were based on the Pearson coefficient (Glantz, 2002; Howell, 2006). For reproducibility studies on the same rater, same subjects, and SNR measurements we have used the intraclass indices (ANOVA), paired student t-test and the Bland-Altman bias analysis as described elsewhere (Glantz 2002; Hasan, 2006). Meta statistical analyses of published reports were conducted as described elsewhere (Hedges, 1985; Hunter and Schmidt, 2004)

### **Data Quality and Reproducibility**

Since the data from all healthy control children were collected over a one-year span at high SNR, we collected, using identical protocols a database of water phantom measurements, healthy adult controls to assure the field uniformity and stability of the MRI scanner. The diffusion encoding (Icosa21b) provided three levels of SNR and hence the SNR dependence of the DTI-metrics and estimates (fiber tracks, group comparisons, correlations) reported in this work were also studied on all subjects. The DTI feature space generated by carefully placed ROIs was invariant to rating as judged by the intraclass correlation coefficient.

## **Results**

### **Illustration of the DTI based feature Space**

Figure 1 illustrates the formation of the tissue clustering borders using the FA vs. mean diffusivity space. We have also included results from the entire spherical water phantom measurements acquired using the same DTI protocol over the time span of the brain studies. Notice the stability and reproducibility of our studies as judged by the insignificant spread of water phantom DTI-metric measurements over a one year span. The GM, WM and CSF are separable using this two dimensional feature space at the selected high SNR and spatial resolution working levels.

### **Comparison of DTI segmentation with cMRI methods**

Figure 2 shows a group mean comparison of the DTI results and cMRI results on all healthy controls. Notice that although the approaches yielded roughly comparable estimates, the

differences are statistically significant ( $p < 0.01$ ) using post-hoc paired comparisons (see Table 2 for more quantitative details). However, both the DTI and cMRI methods gave group means comparable consistent with published results on children (see Table 1). The DTI measured volume tend to be higher for GM and lower for WM but are largely consistent with many of the studies on children listed in Table 1, yielding a GMf of  $\sim 0.6$ . Our results are particularly convergent with more recent studies (e.g. Sowell et al. 2003; Castellanos et al. 2002; De Bellis, 2001).

The cMRI methods were generally less sensitive to age variation compared to the DTI approach (see Table 2 for detailed summary and comparisons). To investigate the origin of the group variance in the DTI based method we plotted the GMf, WMf, total intracranial volume and GM-to-WM ratio as function of age (Figure 3). The DTI based group mean whole brain volume partitions in  $\text{cm}^3$  (mL) were found to be ( $\text{GM}_v = 913 \pm 81$ ,  $\text{WM}_v = 494 \pm 68$ ,  $\text{CSF}_v = 97 \pm 27$   $\text{cm}^3$ ;  $N=29$ ), group fraction means  $\pm$  SE: ( $\text{GM}_f = 60.7 \pm 3.1$ ,  $\text{WM}_f = 32.9 \pm 3.3$ ,  $\text{CSF}_f = 6.4 \pm 1.7$ ) and age trends:  $r(\text{age}, \text{GM}_f; p) = (-0.48; 0.008)$ ,  $r(\text{WM}_f, \text{age}; p) = 0.35; 0.06$ ,  $r(\text{CSF}_f, \text{age}; p) = (0.145; 0.47)$ . Notice the ability of the DTI approach to capture the predicted trends (see Table 1 for an extensive comparison). The DTI-segmented whole-brain fractions (GMf, WMf & CSf) and age trends were not significantly different between the age-matched males and females in our sample. The cMRI published or non-DTI based (Table 1) GM-to-WM volume ratios are shown in Figure 4 as function of the population mean age (see Table 1), and contrasted with our own results. A meta analysis (Hedges 1985; Hunter and Schmidt 2004) of the 22 cMRI published quantitative reports incorporating age of the population (Table 1) on the gray matter fractions provided the regression equation  $\text{GM}_f = 0.67 - 0.00187 * \text{Average Age}$ ;  $r = -0.245$ ,  $p < 0.03$ ). Our DTI segmentation results are consistent with published results on both mean values and rate of change of GMf and WMf (Rajapakse et al., 1996; see Table 1).

## Discussion

In this report, we described and applied for the first time a novel automated approach for tissue segmentation that is based on intrinsic rotationally-invariant and coregistered metrics derived from the DTI data. The approach is based on deterministic and trained feature space (Figure 1) derived on a population of subjects using the same high SNR protocol. The simple approach presented and used can also be adapted for more regionally driven segmentation and to incorporate more established statistical and probabilistic segmentation methods (Ali et al., 2005; Lenglet et al. 2006; Liu et al. 2006; Maas et al. 2004; Wang et al. 2005; Wiegell et al. 2003). We have validated the approach using a cohort of developing child and adolescent brains and found that the whole brain GM and WM fractions derived and the corresponding rate of change per unit year are comparable to published trends (see meta analysis results above, Table 1, 2; Figs 2–4), and to estimates based on two alternative methods based on cMRI. The two cMRI methods also yielded significantly different estimates of tissue fractions even though they were based on the same brains, image acquisition sequences, and normalized to the same template (see Table 2). Thus, the differences across methods cannot be attributed to the DTI method alone. This DTI approach can be fused with cMRI methods or used, for example, as a first pass segmentation to evaluate the needed thresholds for compact white matter fiber tracking. We have tested this approach using optimally designed encoding schemes that were shown to have minimal rotational variance and biases in the estimation of grey and white matter DTI metrics (Hasan, 2006; Hasan and Narayana 2006). Our approach used the DTI-derived eigenvalues and the corresponding scalar maps such as mean diffusivity and FA.

Since DTI requires high signal-to-noise ratios (SNR) and high spatial accuracy to distinguish white-grey matter tissue boundaries, the estimation of anisotropy in grey matter, in particular, is expected to be problematic (Conturo et al., 1995; Hasan and Narayana, 2003; Le Bihan, 1995; Pierpaoli et al., 1996). This problem is mitigated by the use of rotationally-invariant

encoding schemes at high SNR as implemented in this paper. The entire DTI scan took 7 minutes, which is tolerable by most children, young and older adults. In special situations when imaging sedated children or uncooperative subjects, the optimized DTI data set would provide both microstructural and macrostructural information, so that the approach presented here offers an efficient use of available scan time. The DTI data acquired did not undergo any further special steps as all data were coregistered as part of the DTI processing pipeline. Additional advantages of the DTI approach over multi-modal cMRI include (I) the use of high SNR isotropically distributed and rotationally invariant diffusion encoding scheme (Hasan and Narayana 2003; Pierpaoli et al., 1996), (II) the use of reduced image distortions following the adoption of parallel imaging (Bammer et al., 2002; Jaerman et al., 2005), (III) the utility of intrinsic tissue diffusion based contrast in the self-registered DTI maps, and (IV) the fact that DTI processing involves decoupling of radio frequency inhomogeneity field related intensity variations which are common in conventional MRI based approaches (Bammer et al., 2002; Rajapakse et al., 1996a,b; Table 1 references). The whole-brain GM, WM and CSF fractions found using this approach are consistent with published reports using validated multispectral cMRI approaches on children, adolescents (Castellanos et al., 2002; Corchesne et al. 2000; Giedd et al., 1999; Giedd et al, 2004; Simon et al., 2005; Sowell et al. 2003; Wilde et al. 2005; Table 1) and the smooth transitions from birth to childhood (Matsuzawa et al., 2001) and the universal transition to adulthood (Ge et al., 2002; Ingelese and Ge 2004; Corchesne et al. 2000; Sowell et al., 2003; see also Table 1).

GM/WM ratios across cMRI studies of children and adolescents show marked variability, ranging from 1.23 to 2.65. Despite this variability, the DTI-based GM/WM ratio of our normally developing sample was 1.88, which closely fit the age-adjusted regression line for all studies and provided further support for the validity of our approach (see Fig. 4 and Table 2). In contrast to the cMRI segmentation solutions, the DTI volumetry metrics were sensitive to age-related variation that reflects the dynamic progressive and regressive neural processes regulating axonal and dendritic development and connectivity. This sensitivity represents a significant advantage for studies of developmental changes in tissue microstructure and macrostructure. DTI-based volumetry also has potential to enhance understanding of changes in tissue composition in clinical populations due to interactions of developmental changes with injury or disease processes

It is also noteworthy that the partial averaging in the voxel and SNR related biases in DTI can be studied further as function of spin history preparation (CSF fluid, GM or WM attenuation), encoding, slice thickness, or the b-factor. For example, the encoding scheme adopted in our work (Icosa21b) can provide independent rotationally-invariant data sets (Icosa6 and Icosa15) at different SNR values using the same data (Hasan and Narayana, 2003), and hence some sensitivity analysis can be conducted to show the SNR performance of this approach both regionally and globally. These studies will be reported in a future communication.

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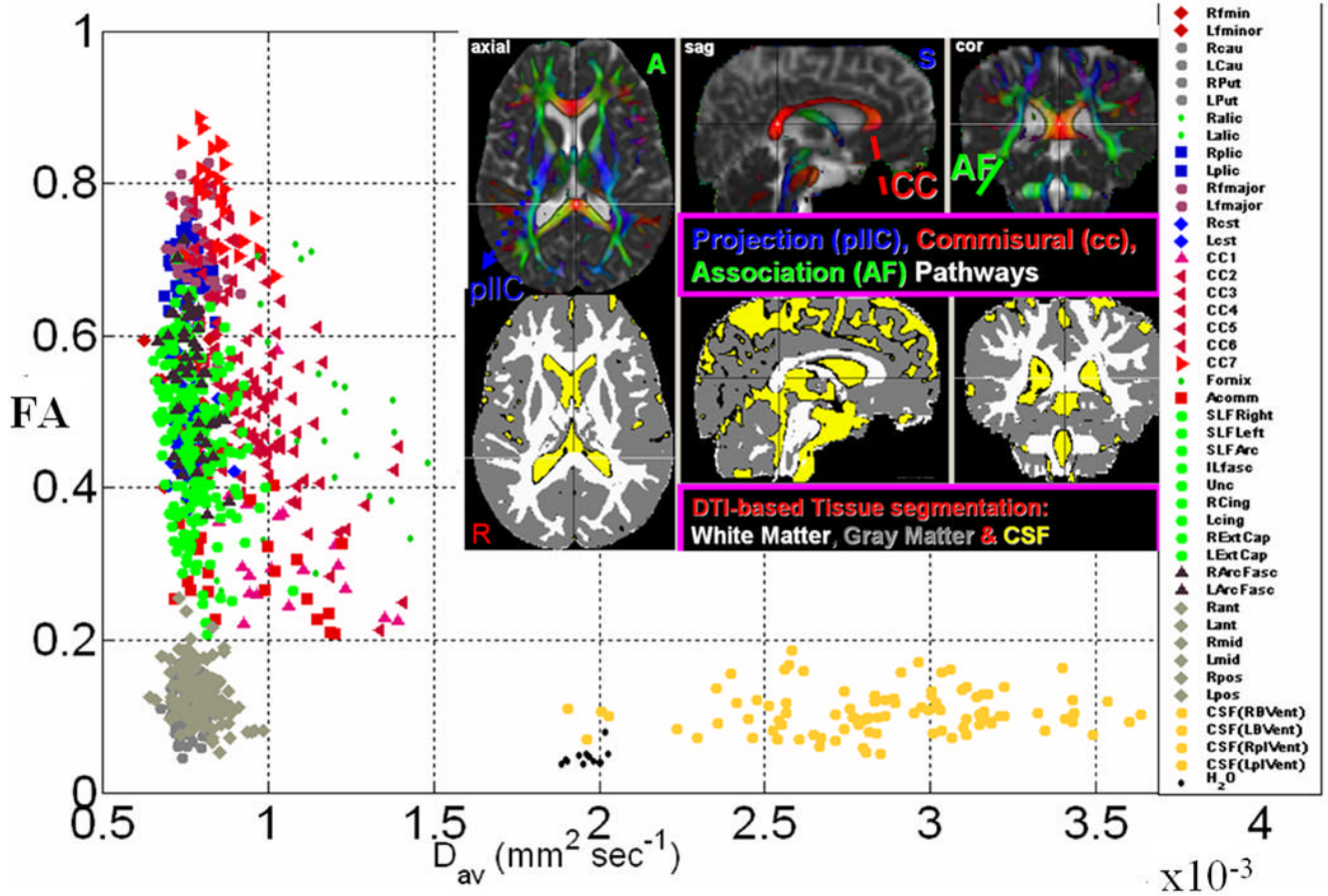
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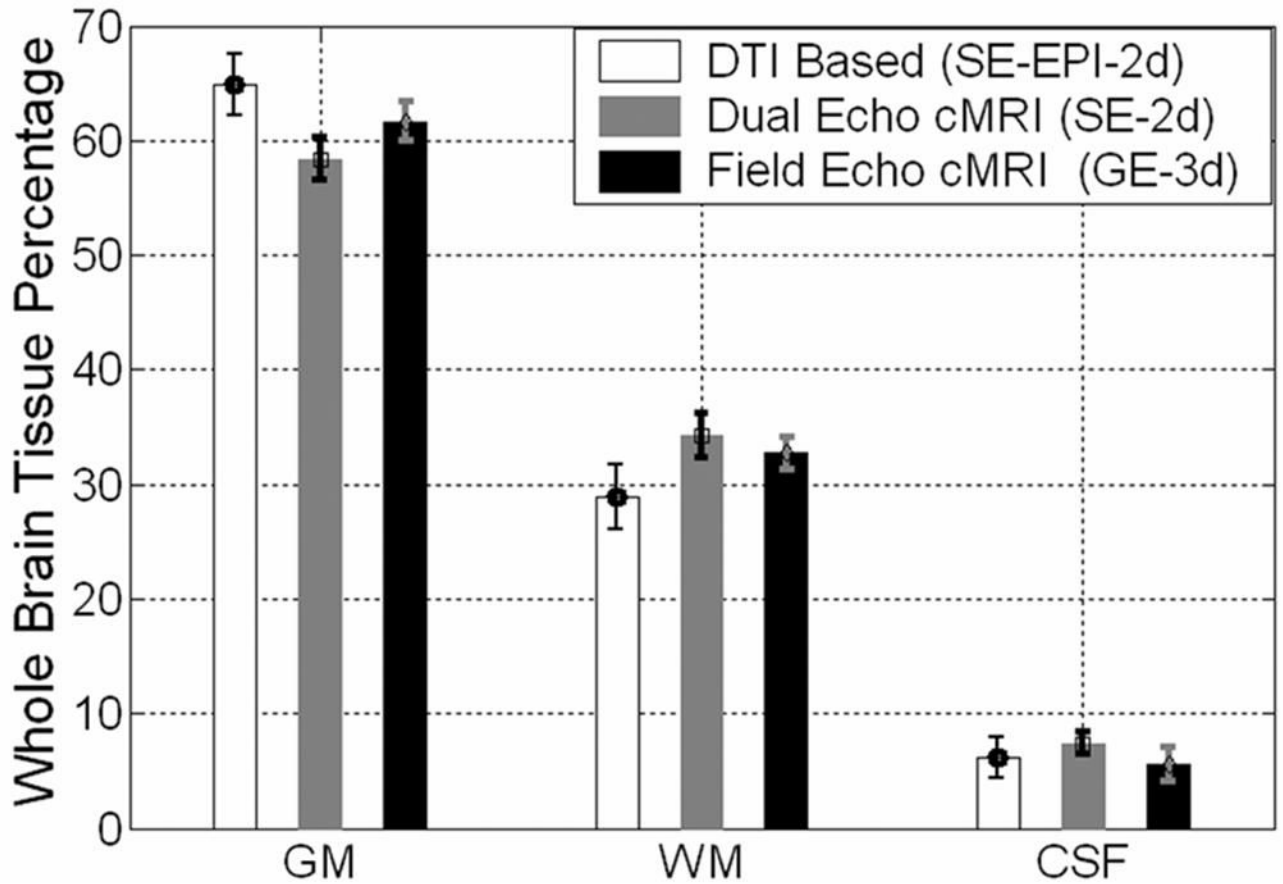
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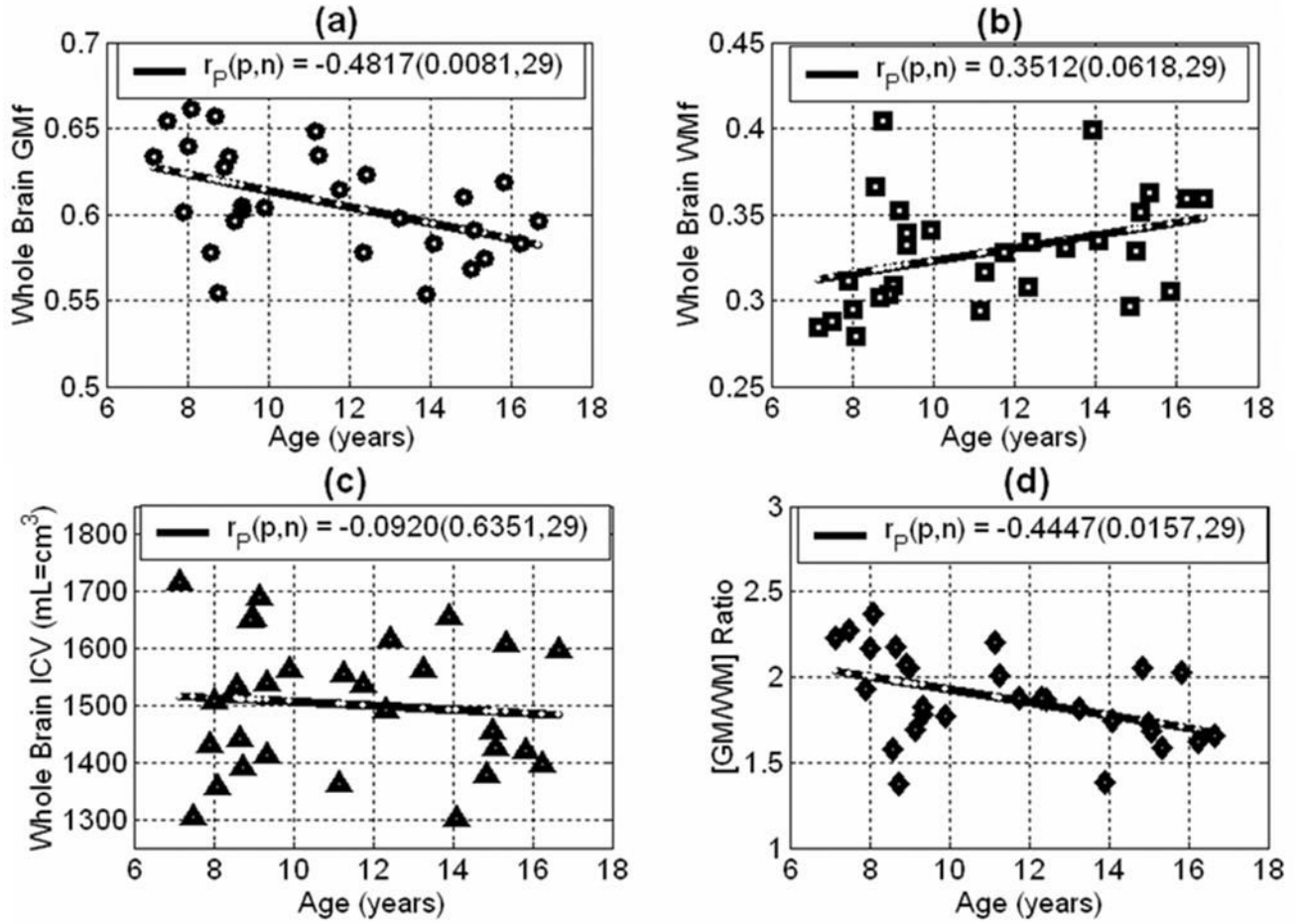


**Figure 1.** Illustration of the formation of the DTI metric (mean diffusivity vs. fractional anisotropy) training feature space on all children. The scatter plot shows the FA vs.  $D_{av}$  two dimensional space on 50 manually selected ROIs that span the whole-brain white matter fiber networks (association, projection, and commissural), grey matter (cortical and basal ganglia) and ventricular CSF. For scanner temporal stability and quality assurance, the scatter plot also includes the DTI (FA vs.  $D_{av}$ ) results on the entire isotropic water phantom collected using the same protocol over the time span of acquisition of the data on controls. Notice the ability to classify the different tissue types in this space

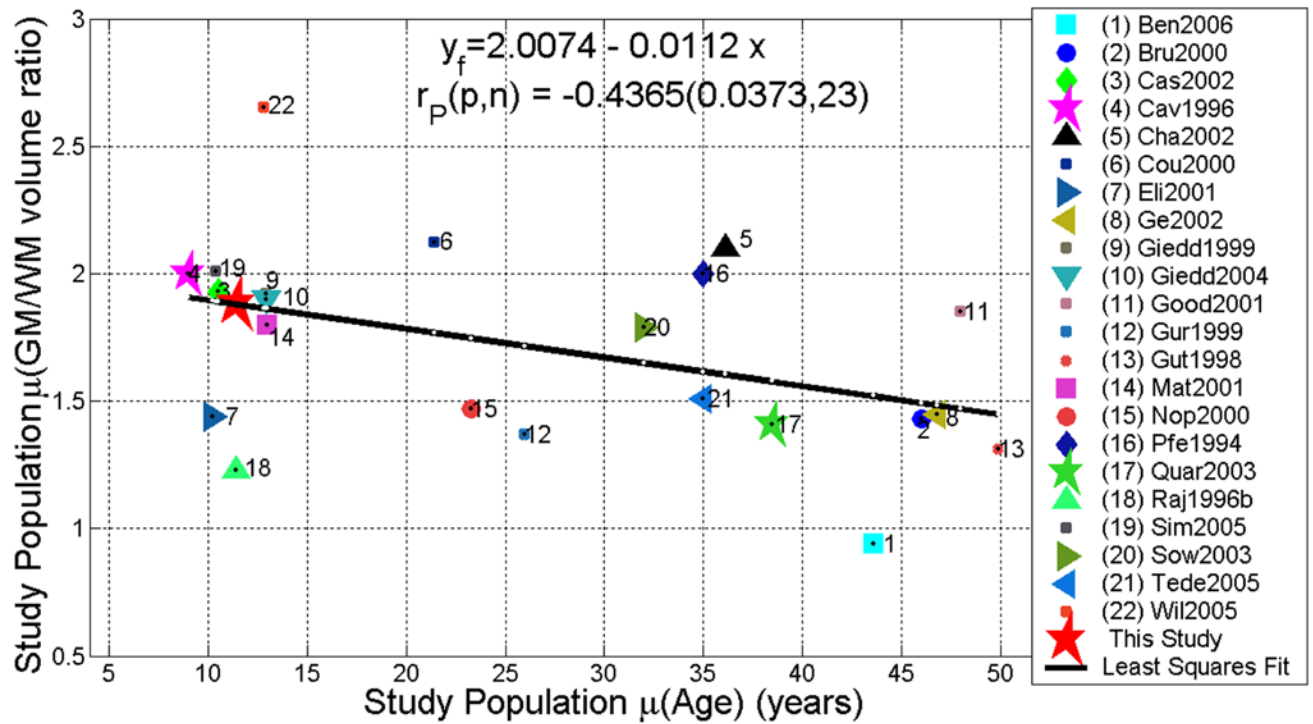


**Figure 2.**

Group comparison of three different approaches used in tissue segmentation of the whole-brain (a) DTI based and (b) cMRI based using the 2d dual echo and (c) using the 3d spgr data sets. The ANOVA tests and *ad hoc* follow-up paired comparisons indicated that the three methods are comparable but differed significantly ( $p < 0.01$ ). The sources of the variability in the DTI approach as function of age is analyzed in Figure 3.



**Figure 3.** The scatter plots and least squares linear regression fits of age vs. (a) GMf and (b) WMf as obtained using the DTI-based tissue segmentation approach. The age vs. total intra cranial volume and grey-to-white matter volume ratios are shown in c and d respectively. Notice (a) the decrease in GMf, (b) increase in WMf, (c) the variability of the ICV due to variations in head size and gender and (d) the global decrease in GM/WM ratio (or corresponding whole-brain fractions) as function of age. The correlation coefficient and corresponding p values are also shown.



**Figure 4.** A scatter plot showing the mean population age in years vs. the whole-brain GM/WM volume ratio reported in 22 quantitative studies summarized in Table 1 on both children, young and older adults. The least-squares linear regression fit predicts a steady decrease in the GM/WM ratio vs. age relation (our study result is also shown; see also Figure 3d and the underlined entire life span studies in Table 1).

**Table 1**  
**Key Publications using MRI volumetry of the normal human whole-brain (WB)**

Study	Pop. N, Gender	Age Range MN±SE yrs	Method	Regions	GMf WMf CSFf *100 (%)	GM WM CSF (mL)	GM/ WM ratio ~	Age vs. Tissue Type Trend (r (p) / slope)
Ben2006	89 39M 50F	11–76 43.6	1.5T 5mm×24 axial	WB	NA	730 780	0.94 (NI)	GM ↓ 2.8 ml/yr WM ↓ 1.1 ml/yr CSF NA
Bru2000	55 24M 31F	16–76	1.5T 4mm×30 axial	WB	56 39 4	NA	1.43	GMf –0.68 (0.00001) ↓ WMf not significant CSFf 0.66 (0.00001) ↑
Cas2002	139 56F 83M	4.5–19 yrs (10.5 ±3.2)	1.5T 1.5mm axial	Cereb	66 34	728±74 377±50	1.93	NA
Cav1996	30 15M 15F	7–11 9	1.5T 3.1mm × 60 axial	WB/Cereb	58 29 13	761 380 171	2.0	GM NA WM ↑ CSF NA
Cha2002	27 13M 14F	23.2–55.2 36.1	1.5T 1.5mm×124 T1w axial	WB	55.5 26 17.5	750±56 356±32 237	2.1	NA
Cou2000	116 79M 37F	19m–80 yrs 21.4±20	1.5T 3mm axial	WB	59 29 12	805 400 150	2.12±0.5	GM ↓ –0.75 (0.001) WM ↑↓ (peak ~ 40–50 yrs) CSF ↑ 0.85 (0.001)
Eli2001	85 21M 64F	4–19 10.2±3.8	1.5T 5mm axial	WB	54 38 8	655±74 455±73 100±26	1.44	GM ↓ WM ↑ CSF ↓
Ge2002	54 22M 32F	20–86 46.8±19.3	1.5T 3mm 50sl axial	WB	50 35 14	680±102 478±72 188	1.45	GMf ↓ –0.57 (0.0001) WMf ↑↓ –0.35 (0.009) CSFf NA
Giedd1999	145 89M 56F	4.2–21.6	1.5T 1.5mm3daxial	WB	55 28 17	730 380 225	1.92	GM NA WM ↑ 12.4% CSF NA
Giedd2004	161 95M 66F	4.2–21.6	n/a	Cereb	NA	727 383	1.90	GM ↓ WM ↑ CSF NA
Good2001	465 265M 200F	17–79	2.0T 1.5mm×120 sagmp-rage	WB	48 26 25	788 424 399	1.85	GM ↓ R <sup>2</sup> =0.48 WM ↑↓ R <sup>2</sup> =0.33 CSF ↑ R <sup>2</sup> =0.38
Gur1999	80 40M 40F	18–45 26±5.6	1.5T 5mm axial	WB	53 39 8	663 483 102	1.37	GM ↓ –0.26 (0.01) WM NA CSF NA
Gut1998	72 22M 50F	18–81	1.5T 3mm Dual axial	WB	47 36 11		1.31	GM ↓ –0.44 (0.46) WM ↓ –0.084 (0.07)



Study	Pop. N, Gender	Age Range MN±SE yrs	Method	Regions	GMf WMf CSFf *100 (%)	GM WM CSF (mL)	GM/ WM ratio ~	Age vs. Tissue Type Trend (r (p) / slope)
								CSF ↑ 0.996 (0.0003)
<b>Jer2001</b>	78 37M 41F	30–99 64±17.4	1.5T 4mm coronal	Cereb	NA	NA	NA	GM r = –0.53 (0.001) WM r = – 0.63 (0.001) CSF r = 0.74 (0.001)
<b>Mat2001</b>	34 17M17F	1m–26yrs	1.5T 1mm×155 axial	WB	55 31 14	675 375 175	1.8	GM ~↑ WM ↑ CSF~↑
<b>Nop2000</b>	84 42M 42F	19–31 23.3±2.2	1.5T 2mm coronal	WB (no cerebellum)	55 38 7	708 483 84.5	1.47	NA
<b>Pfe1994</b>	161	3m–70yrs	NA	WB	NA	NA	2.0	GM ↓ 0.7 (mL/yr) WM ↑, CSF ↑ 0.6
<b>Quar2003</b>	54 22M 32F	16–70 38.5 ±13.1	1.5T 4mm axial	WB	53% 38% 9%	NA	1.41	GM ↓ –0.43 (0.001) WM no change CSF ↑ 0.47 (0.001)
<b>Raj1996b</b>	105 56M 49F	4–18 11.4±3.5	1.5T 1.5mm×124 axial	WB Shelling No CS F	54 44	NA	1.23	GM ↓ WM ↑
<b>Sim2005</b>	18 11M 7F	7.5–14.2 10.4±2	1.5T 1mm×160 sag;mp-rage	WB	56 28 16	763 379 223	2.01	NA
<b>Sow2003</b>	176 86F 90M	7–87 yrs 32±22	1.5T 1.2mm×124 axial	WB	58 32 10	825 460 150	1.79	GM ↓ WM ↓↓ peak ~ 40–50 yrs CSF ↑
<b>Tede2005</b>	104	20–50	1 T 4mm×30 axial		53 35 12		1.51	NA
<b>Wil2005</b>	16 8M 8F	9–16.4 12.8±2.4	1.5T 1mm T1w 1.5mm T2w sagittal	WB	69 25 6	884±77 333±71 72±33	2.65	NA
<b>This Study</b>	29 14M,15F 25R/4L	7.2–16.7 11.4±3.1 med=11.4	3.0T 3mm×44 DTI-epi	WB	61 ±3.1 33 ±3.3 6 ±1.7	913±81.4 494±68 97±27	1.88±0.26	GM ↓ –0.48 (0.008) WM ↑ 0.35 (0.06) CSF 0.17 (0.38)

\* Data reported from healthy control human brains. Studies are listed alphabetically by leading author last name and year. Underlined studies covered the entire human life span. The up and down arrows summarize the reported trends; the up-down arrows indicate a parabolic (rise-plateau and decay trends).

M=males; F=females, m=month, yrs=years; GM=grey matter, WM=white matter; WB=whole brain, mL= cm<sup>3</sup>; NA=not available; NI=not included in the group mean vs. age GM/WM regression in Fig 4.)

**Table 2**  
**Detailed group mean, age correlations and rate of change comparisons between the three methods used**

N=28	cMRI-2d Dual Echo Method 1	cMRI-3d SPGR Method 2	DTI-segmentation Method 3
$\mu \pm \sigma$ (GMf) p Group Means paired t-test	0.5838 $\pm$ 0.0178 p(1,3)=0.001	0.6169 $\pm$ 0.0174 p(1,2)<1e-5	0.6074 $\pm$ 0.0305 p(2,3)=0.10
$\mu \pm \sigma$ (WMf)	0.3424 $\pm$ 0.0198 p(1,3)=0.07	0.3281 $\pm$ 0.0131 p(1,2)=0.0005	0.3291 $\pm$ 0.0323 p(2,3)=0.87
$\mu \pm \sigma$ (GM/WM)	1.7131 $\pm$ 0.1447 p(1,3)=0.009	1.8847 $\pm$ 0.1130 p(1,2) < 1e-5	1.8696 $\pm$ 0.2609 p(2,3)=0.76
r (GMf, age) & p	-0.1916 (0.33) p(1,3)=0.21	-0.2234(0.25) p(1,2)=0.91	-0.4969(0.007) p(2,3)=0.26
r (WMf, age) & p	0.1294(0.51) p(1,3)=0.28	0.2076(0.29) p(1,2)=0.78	0.4088(0.03) p(2,3)=0.43
r (GM/WM) & p	-0.1532(0.44) p(1,3)=0.18	-0.2424(0.21) p(1,2)=0.74	-0.4899(0.008) p(2,3)=0.31
$\beta \pm \sigma(\beta)$ (GMf, age)	-0.0011 $\pm$ 0.0011 p(1,3)=0.06	-0.0013 $\pm$ 0.0011 p(1,2)=0.90	-0.005 $\pm$ 0.0017 p(2,3)=0.07
$\beta \pm \sigma(\beta)$ (WMf, age)	0.0008 $\pm$ 0.0013 p(1,3)=0.28	0.0009 $\pm$ 0.0008 p(1,2)=0.95	0.0043 $\pm$ 0.0029 p(2,3)=0.26
$\beta \pm \sigma(\beta)$ (GM/WM)	-0.0073 $\pm$ 0.0092 p(1,3)=0.05	-0.0090 $\pm$ 0.0071 p(1,2)=0.88	-0.0421 $\pm$ 0.0147 p(2,3)=0.05

- The subjects (N=28) are the same included in Figure 1 (one female subject 1/29 was excluded as the cMRI segmentation results were not feasible)
- GMf=GM/ICV & WMf=WM/ICV (volume fractions to total intracranial volume are used to account for variability due to gender and brain size at equal age).
- The correlation ( $r$ ) is the Pearson coefficient. The slope  $\beta$  for the least squares regression (rate of change of  $y =$  GMf, WMf or GM/WM per unit  $x =$  age in years can also be computed from  $r$  as  $\beta = r * \frac{\sigma_y}{\sigma_x}$  ( $x = \text{age}$  ( $\mu \pm \sigma$ ) = 11.2  $\pm$  3.0; N=28)
- All possible post hoc paired comparisons between the three methods (1, 2&3) are two-tailed. The paired t-test is used for group comparisons (1,3; 2,1 & 3,2).
- The age-correlation ( $r$ ) and rate ( $\beta$ ) comparisons are done as described in Glantz (2002) and Howell (2006).