

SAMPLE REPORT



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Results of Quantitative Volumetric and Diffusion Tensor Imaging Analyses

Client Name: [REDACTED]
Date of Birth: [REDACTED]
Gender: Male
Date of Data Collection: 09/25/2020
Date of Report: 10/05/2020

Background

The patient is a 41 year old male who self-reports that he was in an industrial accident 6 months prior to this test. He states that he has developed worsening headaches, fatigue, memory difficulties and personality changes. He indicates that he “does not feel right.” His doctor wanted him to have this test.

Technical Details

Magnetic resonance imaging was performed on 09/25/2020 at the MRI facility. Data were collected on an MRI machine using multiple imaging protocols including a 3D thin-slice T1-weighted sequence and a Diffusion Tensor Imaging (DTI) sequence with 30+ directions.

Images from the T1-weighted acquisition were used for quantitative volumetric analyses. The *Quantify Volumetric* process uses a standardized, objective, and automated SPM12-based processing pipeline to calculate regional brain volumes from 107 brain areas pre-selected from TDBrodmann and AAL Atlases. These volumetric measures were scaled by total brain volume to correct for brain size. As part of the *Quantify Volumetric* quality-assurance process, a senior neuroscientist evaluated the T1 data with respect to template alignment, motion, and artifacts that might substantially compromise the validity of results. The T1 data for the client were determined to be of acceptable quality for volumetric analyses.

Data from the Diffusion Tensor Imaging sequence were used for quantitative evaluation of white matter integrity, with regional average Fractional Anisotropy (FA) as the core metric. The *Quantify DTI* process uses a standardized, objective, and automated FSL-based processing pipeline to calculate FA values from 48 fiber tract regions pre-selected from the Johns Hopkins University white matter atlas. This pipeline includes automated motion and Eddy current corrections. As part of the *Quantify DTI* quality-assurance process, a senior neuroscientist evaluated the DTI data with respect to template alignment, motion, and other artifacts that might substantially compromise the validity of results. The DTI data for the client were determined to be of acceptable quality for analyses of FA values.

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The volumetric and FA metrics for the client were statistically evaluated with respect to average volumetric and FA metrics derived from all gender and age-range (+/- 10 years) matched control subjects within a large database of >1000 neuro-typical subjects that is maintained by the MINDSET Integrated in Albuquerque, New Mexico. For this statistical evaluation, a total of 106 matched MINDSET control datasets were identified for volumetric evaluation, with 104 datasets identified for FA evaluation. Each MINDSET database control subject was without any reported history of (1) traumatic brain injury, (2) neurological or psychiatric disease or injury, (3) substance use disorder, or (4) learning or developmental disability.

The MINDSET control data has been collected on an MRI system that belongs to the MIND Research Network (MRN) in Albuquerque. MRN and the MRI facility data acquisition parameters were identical. To account for any systematic differences between the scanners employed, several procedures were implemented. First, data acquisition parameters for the client were set to be comparable to those used in the collection of the MINDSET control datasets. Second, the data analysis procedures used to assess the data from the client were the same as those used in generation of the MINDSET control database. Third, prior to statistical analyses, data metrics for the client were scaled using brain-region specific, across-site matching factors derived from a traveling human phantom previously evaluated on both the MINDSET and MRI facility scanners. This scaling procedure provides for harmonization of datasets across the two scanners and allows the client's DI data to be validly evaluated with respect to the MINDSET normative database.

See the document [Appendix-I-MR-2020](#) for additional information of basic neuroscience and methodological details.

Volumetric Findings

As shown in Figure 1, volumetric analyses conducted individually on 107 brain regions showed 13 of the 107 evaluated regions to have an atypical volume ($p < 0.05$; 3 low; 10 high). Figure 2 provides demonstrative information on the spatial location of each evaluated brain region that showed an atypical volume on isolated individual testing.

Given the number of multiple regional brain volume comparisons ($N=107$), there is the possibility that even a subject without any history of neurological/psychiatric disease or injury may show a few brain regions with 'false positive' identification as statistically abnormal. To address this issue of false positives during multiple comparisons, a Benjamini-Hochberg correction was applied with a False Discovery Rate of 25%.

Following statistical correction for multiple comparisons, 6 of the brain regions with an atypical volume on isolated individual evaluation was still found to be abnormal from a statistical perspective (2 low; 4 high).

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DTI Findings

As shown in Figure 3, DTI analyses conducted individually on 48 fiber tract regions showed 10 of the 48 evaluated tract regions to have an atypical FA value ($p < 0.05$; 0 low; 10 high). Figure 4 provides demonstrative information on each of the evaluated fiber tract regions with an atypical FA value on isolated individual testing.

Given the number of multiple fiber tract FA comparisons ($N=48$), there is the possibility that even a subject without any history of neurological/psychiatric disease or injury may show a few fiber tract

regions with 'false positive' identification as statistically abnormal. To address this issue of false positives during multiple comparisons, a Benjamini-Hochberg correction was applied with a False Discovery Rate of 25%.

Following statistical correction for multiple comparisons, all 10 of the fiber tract regions with an atypical FA value on isolated individual evaluation was still found to be abnormal from a statistical perspective (0 low; 10 high).

Summary

The volumetric data for Mr. [REDACTED] were abnormal from a statistical perspective, with 6 brain regions (2 low and 4 high) showing deviant volumes on both isolated individual testing and after correction for multiple comparisons.

Low volumes may reflect perturbation of early developmental processes, malnutrition, toxic stress, toxic exposures, or more commonly neurological-psychiatric disease and/or brain injury. High volumes may reflect a perturbation of early developmental processes; baseline skill enhancement; or injury/disease related edema, astrogliosis, chronic inflammation, or reactive compensatory mechanisms triggered by damage to other brain regions.

The DTI data for Mr. [REDACTED] were abnormal from a statistical perspective, with 10 fiber tract regions (0 low and 10 high) showing deviant FA values on both isolated individual testing and after correction for multiple comparisons.

Possible causes for high FA include a developmental failure of axonal pruning, disease/injury related intracellular cytogenic edema, compaction of neurofilament, neuro-inflammation with microglial activation, astrogliosis, loss of crossing fibers, and/or compensatory mechanisms.

In summary, both the volumetric and DTI data for Mr. [REDACTED] were objectively abnormal from a statistical perspective.

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Quantify analyses provide an objective assessment of the structural integrity of the brain. They may reveal neurobiological insults that can be caused by head trauma, toxic exposures, or other conditions. However, quantitative volumetric and DTI analyses in isolation do not identify specific causes of brain compromise as multiple conditions can give rise to volumetric and FA abnormalities. The results should be interpreted by qualified experts with an understanding of the volumetric, DTI, and statistical methods used and knowledge of the foundational research and peer-reviewed literature linking imaging findings to neurobiological factors.

Determination of the most likely etiology of volumetric and DTI findings requires a comprehensive assessment of an individual's history and current status and use of the scientific method for differential evaluation of possible contributions from premorbid and co-morbid factors. Such a process is needed to elucidate the relationships between specific past events and present imaging findings and neurobiological status.

To request a free 15-minute consultation phone call to go over present results, please email info@quantifymri.com. Include your name and contact information and available dates and times for a call within the next week. You will then receive additional information on scheduling the call.



Edward L. Soll, M.D., Radiologist, Medical Director, The Concussion Group

Avery Knapp, MD, Radiologist, The Concussion Group

Jeffrey D. Lewine, Ph.D., Translational NeuroScience, Mindset

PATIENT

Patient: [REDACTED]

Date of Birth: [REDACTED]

Gender: Male

Date of Data Collection: 09/25/2020

Date of Report: 10/05/2020

Review and Comment:

I have supervised the DTI DiCOM data acquisition process performed onsite at Doctors Imaging and reviewed the Mindset Volumetric/DTI analysis; and confirm that the MRI protocols as well as the applied analytic tools were methodically utilized as designed. I accept the final DTI and Volumetric Z-Scores as statistically appropriate.

Edward L. Soll, M.D.
Radiologist and Medical Director
Certified American Board of Radiology, 1973
The Concussion Group

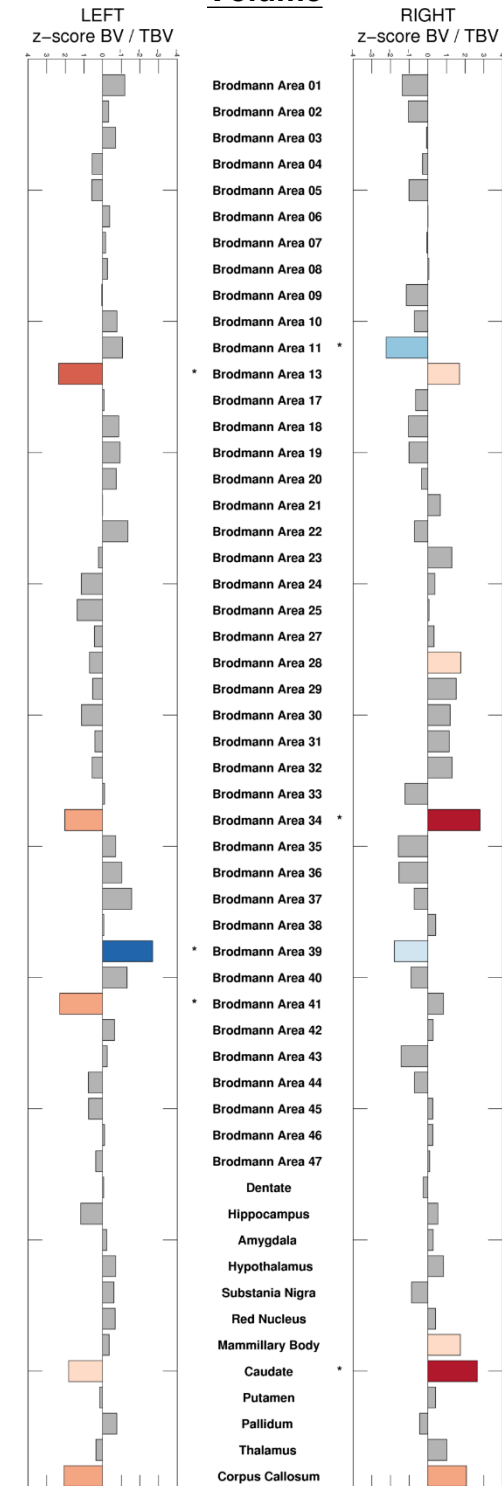
Review and Comment:

I have reviewed the Mindset Volumetric/DTI analysis and accept the final DTI and Volumetric Z-Scores as statistically appropriate.

Avery Knapp, M.D.
Radiologist
Certified American Board of Radiology, 2008
The Concussion Group

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Figure 1: Mr. [REDACTED]
Z-scores for Regional Brain
Volume Scaled by Total Brain
Volume



The corpus callosum is a midline structure.
 Identical values are indicated in left and right columns.

Region-of-interest volumetric analyses

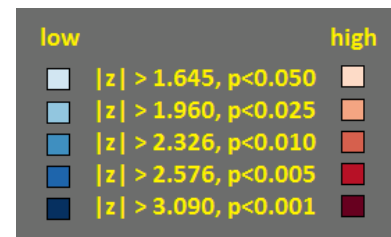
Data are shown as Z-score deviates based on comparison of client data with average data from a gender and age-range (+/- 10 years) matched group of **106** neuro-typical subjects.

At the level of individual isolated analyses

Gray bars show regions with volumes within normal limits.

Red bars show regions with atypically high volume (p<0.05).

Blue bars show regions with atypically low volumes (p<0.05).



For Mr. [REDACTED], isolated evaluation of individual brain regions revealed 13 of 107 brain regions to show atypical volumes: (p<0.05; 3 low; 10 high).

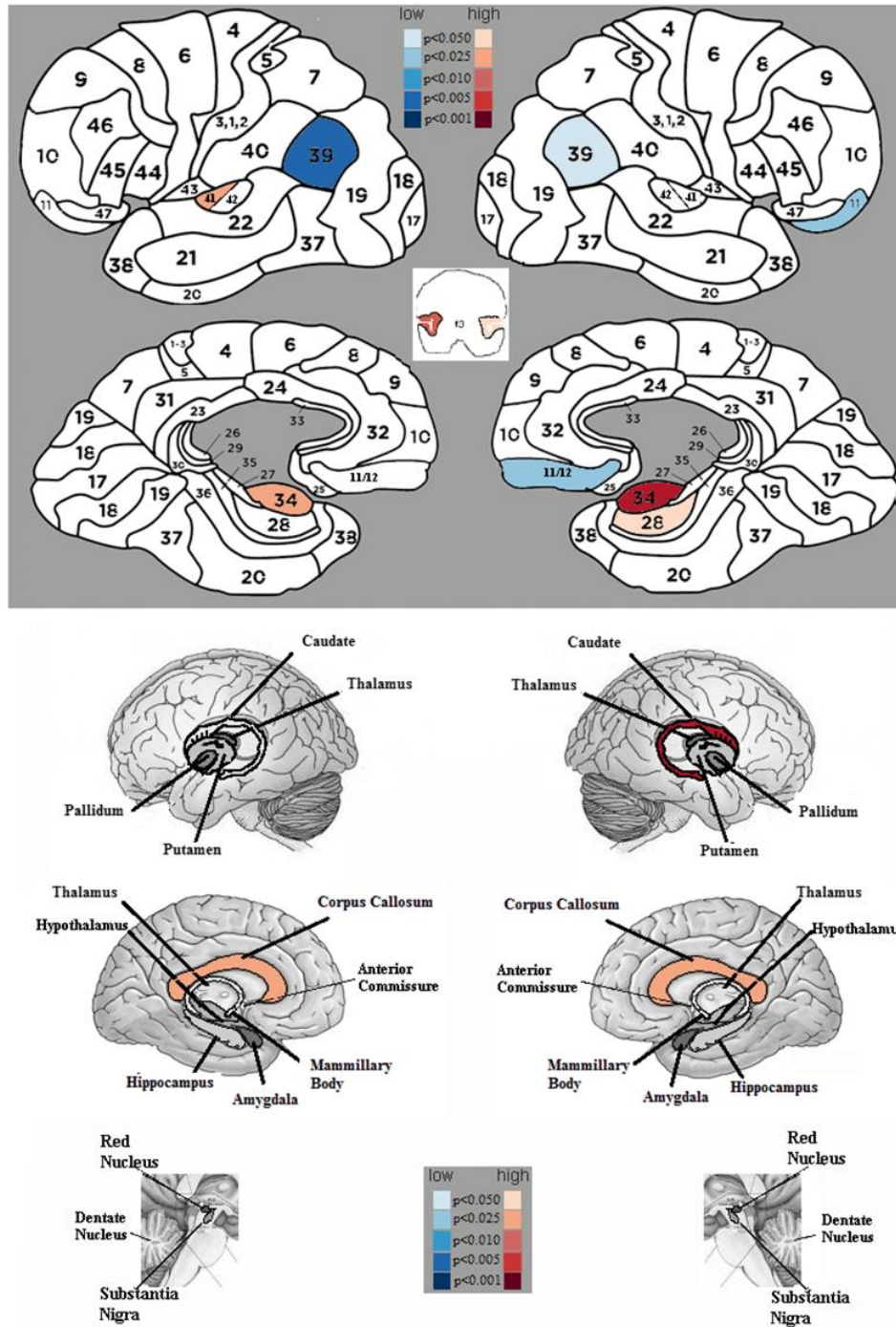
However, only those observations marked with an * survive additional correction for multiple comparisons (FDR=0.25). After correction for multiple comparisons, 6 of the brain regions with an abnormal volume on individual testing were still considered to be abnormal from a statistical perspective: (2 low, 4 high).

Overall, 6 of the 107 evaluated brain regions showed statistically abnormal volumes (2 low; 4 high) on both isolated individual testing (p<0.05) and following correction for multiple comparisons (FDR=0.25).

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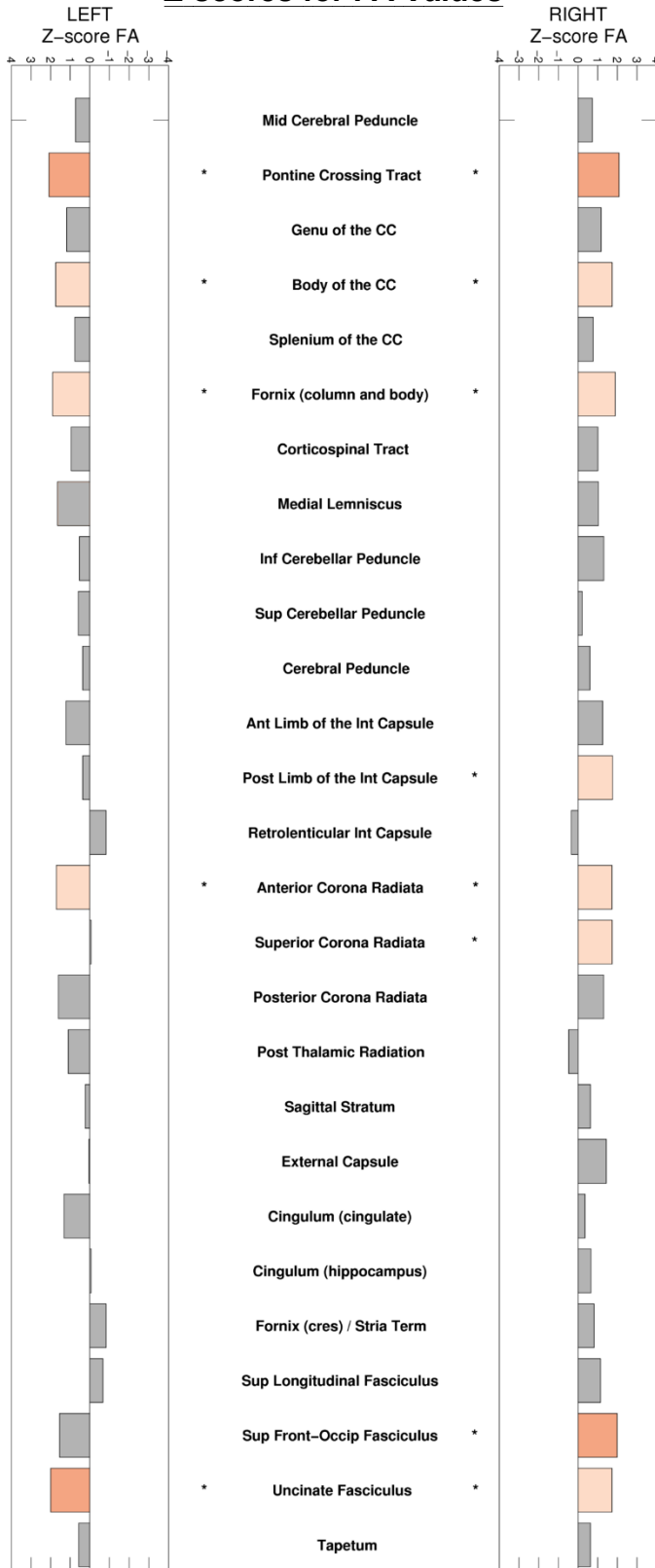
Figure 2: Mr. [REDACTED]
Shaded Brain Regions Show Atypical Volumes on Isolated Individual Testing

Following correction for multiple comparisons, 6 of the brain regions with atypical volumes on isolated individual evaluation were still considered to be abnormal from a statistical perspective.



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**Figure 3: Mr. [REDACTED]
Z-scores for FA Values**



The first 6 tracts are at the midline and without separable left/right components. Identical values are plotted in left and right columns.

FA Values for Fiber Tract Regions

Data are shown as Z-score deviates based on comparison of client data with average data from a gender and age-range (+/- 10 years) matched group of **104** neuro-typical subjects.

At the level of individual isolated analyses

Gray bars show tracts where FA values are within normal limits.

Red bars show tracts with atypically high FA values (p<0.05).

Blue bars show tracts with atypically low FA values (p<0.05).

| low | | high | |
|-----|---------------------|------|--|
| | z > 1.645, p<0.050 | | |
| | z > 1.960, p<0.025 | | |
| | z > 2.326, p<0.010 | | |
| | z > 2.576, p<0.005 | | |
| | z > 3.090, p<0.001 | | |

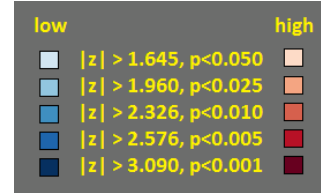
For Mr. [REDACTED], isolated evaluation of individual fiber tracts revealed 10 of 48 regions to show atypical FA values: (p<0.05; 0 low; 10 high).

However, only those observations marked with an * survive additional correction for multiple comparisons (FDR=0.25). After correction for multiple comparisons all 10 of the fiber tract regions with abnormal FA values on individual testing were still considered to be abnormal from a statistical perspective: (0 low, 10 high).

Overall, 10 of the 48 evaluated fiber tract regions showed statistically abnormal FA values (0 low; 10 high) on both isolated individual testing (p<0.05) and following correction for multiple comparisons (FDR=0.25).

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Figure 4: Mr. [REDACTED]
Shaded Tracts Show Atypical FA Values on
Individual Isolated Testing ($p < 0.05$).



| | | | | | | | |
|---|---|--|--|--|--|--------------------------------------|--------------------------------------|
| Middle Cerebellar Peduncle | Pontine Crossing Tract * | Genu of the Corpus Callosum | Body of the Corpus Callosum * | Splenium of the Corpus Callosum | Fornix * | R Corticospinal Tract | L Corticospinal Tract |
| | | | | | | | |
| R Medial Lemniscus | L Medial Lemniscus | R Inferior Cerebellar Peduncle | L Inferior Cerebellar Peduncle | R Superior Cerebellar Peduncle | L Superior Cerebellar Peduncle | R Cerebral Peduncle | L Cerebral Peduncle |
| | | | | | | | |
| R Anterior Limb of the Internal Capsule | L Anterior Limb of the Internal Capsule | R Posterior Limb of the Internal Capsule * | L Posterior Limb of the Internal Capsule | R Retrolenticular part of the Intern Capsule | L Retrolenticular part of the Intern Capsule | R Anterior Corona Radiata * | L Anterior Corona Radiata * |
| | | | | | | | |
| R Superior Corona Radiata * | L Superior Corona Radiata | R Posterior Corona Radiata | L Posterior Corona Radiata | R Posterior Thalamic Radiation | L Posterior Thalamic Radiation | R Sagittal Stratum | L Sagittal Stratum |
| | | | | | | | |
| R External Capsule | L External Capsule | R Cingulum (cingulate gyrus) | L Cingulum (cingulate gyrus) | R Cingulum (hippocampus) | L Cingulum (hippocampus) | R Fornix (cres) and Stria Terminalis | L Fornix (cres) and Stria Terminalis |
| | | | | | | | |
| R Superior Longitudinal Fasciculus | L Superior Longitudinal Fasciculus | R Superior Frontal-Occipital Fasciculus * | L Superior Frontal-Occipital Fasciculus | R Uncinate Fasciculus * | L Uncinate Fasciculus * | R Tapetum | L Tapetum |
| | | | | | | | |

Table 1: Mr. [REDACTED]
Regions-of-Interest with Abnormal Volumes on Individual Evaluation

| Area | | General Location | Supported Functions |
|------------------------|---------|--|---|
| Brodman 1,2,3 combined | L/R | Primary somatosensory cortex | Tactile sensation |
| Brodman 4 | L/R | Primary motor cortex | Motor control |
| Brodman 5 | L/R | Somatosensory association cortex | Tactile object recognition |
| Brodman 6 | L/R | Premotor and Supplementary motor cortex | Control of proximal and trunk muscles; Motor sequencing |
| Brodman 7 | L/R | Somatosensory association cortex | Visuo-spatial processing; Praxic abilities |
| Brodman 8 | L/R | Frontal eye fields | Planning of complex movements, control of eye movements |
| Brodman 9 | L/R | Dorsolateral Prefrontal Cortex | Executive Function, Working Memory |
| Brodman 10 | L/R | Anterior Prefrontal Cortex | Strategic Planning, Cognitive Branching |
| Brodman 11 | R* | Orbital Frontal Cortex | Behavioral/Emotional Regulation, Behavioral Inhibition |
| Brodman 13 | L*/R | Insula | Social emotions, multimodal sensory processing, salience |
| Brodman 17 | L/R | Primary Visual Cortex | Basic Vision |
| Brodman 18 | L/R | Secondary Visual Cortex | Shape recognition, visual attention |
| Brodman 19 | L/R | Association Visual Cortex | Visual-spatial processing, motion, face/ word processing (L/R) |
| Brodman 20 | L/R | Inferior Temporal Gyrus | High-level visual information processing and recognition memory |
| Brodman 21 | L/R | Middle Temporal Gyrus | Complex auditory processing and language |
| Brodman 22 | L/R | Superior Temporal Gyrus | Auditory processing, Receptive Language (Wernicke's area) |
| Brodman 23 | L/R | Posterior Cingulate Cortex | Emotion and Memory, Intrinsic Control |
| Brodman 24 | L/R | Anterior Cingulate Cortex | Behavioral Control, Decision Making, Social Evaluation |
| Brodman 25 | L/R | Subgenual Ventromedial Prefrontal Cortex | Decision making, emotional processing, social behavior |
| Brodman 27 | L/R | Periform Cortex | Olfaction |
| Brodman 28 | L/R | Ventral Entorhinal Cortex | Short-term memory |
| Brodman 29 | R | Retrosplenial Cingulate Cortex | Episodic memory, navigation, imagination, and future planning |
| Brodman 30 | L/R | Part of Cingulate Cortex | Episodic memory, navigation, imagination and future planning |
| Brodman 31 | L/R | Dorsal Posterior Cingulate Cortex | Emotion and Memory, Intrinsic Control |
| Brodman 32 | L/R | Dorsal Anterior Cingulate Cortex | Behavioral Control, Decision Making, Social Evaluation |
| Brodman 33 | L/R | Part of the Anterior Cingulate Cortex | Behavioral Control, Decision Making, Social Evaluation |
| Brodman 34 | L/R* | Dorsal Entorhinal Cortex/Parahippocampal Gyrus | Olfaction, Short-term memory |
| Brodman 35 | L/R | Perirhinal Cortex | Memory, Emotion, |
| Brodman 36 | L/R | Ectorhinal Area | Short-term memory |
| Brodman 37 | L/R | Fusiform Gyrus | Word recognition (L) / Face Processing (R) |
| Brodman 38 | L/R | Temporopolar Regions | Memory, Language |
| Brodman 39 | L*/R | Angular Gyrus | Language, reading, mathematics, attention |
| Brodman 40 | L/R | Supramarginal Gyrus | Spatial perception, phonological choices |
| Brodman 41 | L* | Primary Auditory Cortex | Basic Hearing |
| Brodman 42 | L/R | Auditory Cortex | Auditory processing |
| Brodman 43 | L/R | Gustatory Cortex | Taste |
| Brodman 44 | L/R | Pars opercularis, part of Broca's area | Expressive Language |
| Brodman 45 | L/R | Pars triangularis, part of Broca's area | Expressive Language |
| Brodman 46 | L/R | Dorsolateral Prefrontal Cortex | Executive Function, Working Memory |
| Brodman 47 | L/R | Pars orbitalis, inferior frontal gyrus | Syntax |
| Dentate Nucleus | L/R | Cerebellum | Planning and initiation of voluntary movements |
| Hippocampus | L/R | Medial Temporal Lobe | Short-term Memory |
| Amygdala | L/R | Medial Temporal Lobe | Emotion |
| Hypothalamus | L/R | Just left/right of Midline | Regulates autonomic functions, pituitary, hunger, sleep |
| Substantia Nigra | L/R | Brainstem | Dopaminergic motor control |
| Red Nucleus | L/R | Brainstem | Motor Coordination |
| Mammillary Body | R | Just left/right of Midline | Memory |
| Caudate | L/R* | Part of the Basal Ganglia | Regulates movement, cognition |
| Putamen | L/R | Part of the Basal Ganglia | Regulates movement |
| Pallidum | L/R | Part of the Basal Ganglia | Regulates movement |
| Thalamus | L/R | Just left/right of Midline | Sensory, Motor, Emotional and Cognitive Functioning |
| Corpus Callosum | midline | Midline | Interconnects L/R hemisphere |

Client volume abnormally high, p<0.05 (isolated, individual testing)

High regional brain volumes may reflect perturbation of early developmental pruning, enhanced skill development, or brain injury related edema, astrogliosis, inflammation, and/or compensatory mechanisms.

Client volume abnormally low, p<0.05 (isolated, individual testing).

Low regional brain volumes may reflect perturbation of early developmental processes, malnutrition, toxic stress, toxic exposures, or more commonly neurological/psychiatric disease and/or brain injury.

* indicates survival of correction for multiple comparisons

Table 2: Mr. [REDACTED]
Fiber Tract Regions of Interest with Abnormal FA Values on Individual Isolated Evaluation

Following correction for multiple comparisons, **ALL 10** fiber tract regions with atypical FA values on isolated individual evaluation are still considered to be abnormal from a statistical perspective.

| JHU White Matter Atlas Fiber Tract Regions | | Connections | Supported Functions |
|---|----------|--|---|
| Middle Cerebral Peduncle | Midline | Interconnects Cerebellum and Pons | Initiation and Timing of Volitional Movement |
| Pontine Crossing Tracts | Midline* | Interconnects Pons and Contralateral Cerebellum | Coordination of Movement |
| Genu of the Corpus Callosum | Midline | Interconnects Right and Left Anterior Frontal Lobes | Interhemispheric Integration of Executive Functions |
| Body of the Corpus Callosum | Midline* | Interconnects Right and Left Posterior Frontal Lobes Interconnects Right and Left Parietal Lobes | Interhemispheric integration of Motor and Somatosensory Functions |
| Splenium of the Corpus Callosum | Midline | Interconnects Right and Left Occipital Lobes | Interhemispheric Integration of Visual Functions |
| Fornix | Midline* | Interconnects the Hippocampus and Mammillary Bodies | Short-Term Memory |
| Corticospinal Tracts | L/R | Connects Primary Motor Cortex with Contralateral Spinal Motor Neurons | Motor Control of the Contralateral Side of the Body |
| Medial Lemniscus | L/R | Connects Dorsal Column Nuclei with the Contralateral Thalamus (VPL) | Somatosensory Perception of the Contralateral Side of the Body |
| Inferior Cerebellar Peduncles | L/R | Connects Spinal Cord and the Medulla to the Cerebellum | Posture, Balance, and Coordination |
| Superior Cerebellar Peduncles | L/R | Interconnects Cerebellum to Pons and Midbrain | Motor Coordination and Balance |
| Cerebral Peduncles | L/R | Interconnects Cerebellum with the Thalamus and Motor Cortex | Motor Control |
| Anterior Limb of the Internal Capsule | L/R | Contains Fibers Interconnecting the Thalamus and Frontal Lobe; Lentiform and Caudate Nuclei; Cortex and Corpus Striatum | Motor Control; Higher Cognitive Function |
| Posterior Limb of the Internal Capsule | R* | Contains Fibers Interconnecting Motor Areas with the Brainstem; Midbrain and the Thalamus, Occipital Lobes, and Temporal Lobes | Visual-Spatial Processing, Visual Motion, Face and Word Processing (L/R) |
| Retrolenticular Part of the Internal Capsule | L/R | Interconnects Thalamus and Occipital Cortex | Visual Processing |
| Anterior Corona Radiata | L*/R* | Contains Descending and Ascending Fibers Related to Cortex – especially for the Frontal Lobes | Executive Function, Emotional Control |
| Superior Corona Radiata | R* | Contains Descending and Ascending Fibers Related to Cortex – especially for the Motor Cortex | Motor Control |
| Posterior Corona Radiata | L/R | Contains Descending and Ascending Fibers Related to Cortex – especially for the Parietal Lobes | Attentional Control, Somatosensory Function |
| Posterior Thalamic Radiation | L/R | Interconnects Thalamus and Cortex | Visual and Auditory Function |
| Sagittal Stratum | L/R | Interconnects Thalamus with Occipital, Parietal, Temporal and Cingulate Cortices | Visual, Auditory, and Cognitive Function |
| External Capsule | L/R | Contains Cortico-Cortical Association Fibers for Occipital, Temporal, Parietal, and Cingulate Cortices | Cognitive Processing |
| Cingulum (cingulate cortex) | L/R | Interconnects Cingulate and Pre-Frontal Cortices | Cognitive Processing and Decision Making |
| Cingulum (hippocampus) | L/R | Interconnects Cingulate and Entorhinal Cortices | Memory, Emotional Processing |
| Fornix (cres) and Stria Terminalis | L/R | Interconnects Hippocampus and Mammillary Bodies; Amygdala with the Septal Region and Hypothalamus | Memory, Emotional Processing, Fear Response |
| Superior Longitudinal Fasciculus | L/R | Interconnects the Front and Back of the Cerebrum, Including Frontal, Parietal, Occipital, and Cingulate Areas | Higher Cortical Functions including Language, Attention, Motor Control, and Spatial Processing. |
| Superior Fronto-Occipital Fasciculus | R* | Interconnects the Frontal Lobe with the Occipital and Parietal Lobes | Spatial Awareness |
| Uncinate Fasciculus | L*/R* | Interconnects Hippocampus and Amygdala with Orbital Frontal Cortex | Memory, Emotional Processing, Language |
| Tapetum | L/R | Contains Commissural Fibers Interconnecting Right and Left Temporal Lobes | Interhemispheric Integration for Auditory Processing |

Client FA value abnormally high, $p < 0.05$ (isolated, individual testing).

High FA values may reflect a developmental failure of axonal pruning, disease/injury related intracellular edema, compaction of neurofilament, neuro-inflammation with micro-glial activation, astrogliosis, loss of crossing fibers, and/or compensatory mechanisms.

Client FA value abnormally low, $p < 0.05$ (isolated, individual testing)

Low FA values may reflect perturbed development or more commonly, disease/injury related extracellular edema, demyelination, and/or loss of axonal fibers.

* indicates survival of correction for multiple comparisons

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Appendix-I-MRI-2020

**Basic Concepts in Brain Organization, Magnetic Resonance Imaging, and
Quantitative Assessment of Regional Brain Volumes and Diffusion Tensor Imaging
Data.**

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BASIC CONCEPTS IN BRAIN ORGANIZATION, MAGNETIC RESONANCE IMAGING AND TRAUMATIC BRAIN INJURY

Brain Organization:

The human brain is composed of more than 100 billion cells, including neurons and supporting glial cells. Electrochemical signals are used to encode and transmit information within brain cells, with neurotransmitters used to transfer information from one neuron to another across synapses. Neurons have three key parts: (1) the dendrites, which bring information into the cell, (2) the cell body, which integrates the information, and (3) the axon, which takes information to the next cell. Within the brain, cell bodies often cluster together in what is known as gray matter. The outermost gray matter region of the brain is known as the cerebral cortex. Axons often travel together in tracts, in what is known as the white matter, where they connect different parts of the brain. Cells may be arranged differently in different parts of the cortex. This, along with the specific local and distant interconnections between cells gives rise to cortical specialization and networks – that is, the brain is partly organized into functional modules that are interconnected to support different calculations and behavioral abilities.

Magnetic Resonance Imaging

Magnetic Resonance Imaging (MRI) is a non-invasive technique for evaluating structure and function. By examining how the hydrogen protons of water molecules behave when placed within a steady magnetic field and bombarded by pulses of radio-wave energy, the MRI system can reconstruct a picture of the brain's anatomy. By altering the parameters and sequence of radio-wave pulses, it is also possible to explore the white matter interconnections between brain regions, brain biochemistry, metabolism, and hemodynamics.

Imaging evaluations in patients with brain injury often include T1-weighted, T2-weighted, FLAIR, SWI and DTI sequences. T1-weighted imaging provides exceptional gray-white matter differentiation and a detailed picture of brain anatomy. T2 and FLAIR imaging are especially useful for identifying edema and lesions in the white matter. SWI – susceptibility weighted imaging – is very sensitive to iron deposition associated with bleeding caused by stroke or small hemorrhages that may be caused by traumatic brain injury and other conditions. DTI – Diffusion Tensor Imaging -- examines the diffusion properties of water within the brain. DTI provides insight into the integrity of axonal tracts (which normally restrict the direction of water diffusion to be along [rather than across] the direction of the tracts).

Clinical, visual inspection of MR data can often reveal major pathologies like tumors, multiple sclerosis, and severe head trauma. Milder injuries, however, such as those associated with a concussion, rarely produce MR changes that can be seen on routine visual inspection. Quantitative analyses of regional brain volume (that is, the measurement of the size of various brain regions as visualized on T1-weighted images), and a quantitative assessment of the integrity of white matter pathways (as indexed by Fractional Anisotropy (FA) on DTI), offer more sensitive, objective evaluation of brain compromise.

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Volumetric Analyses

The brain is a 3-Dimensional structure that can be divided into small cubes, known as voxels. For MRI images, an automated computer algorithm can, in each voxel, measure the amounts of gray matter, white matter, and cerebrospinal fluid (CSF). The computer can also divide the brain into regions-of-interest, based on known structural-functional relationships. For example, the computer can determine which voxels comprise brain regions like the hippocampus or primary motor cortex and thereby calculate the total amount of brain tissue in these regions for each individual client. These methods provide quantitative measures of the actual volume of brain structures for each client, and they allow for a statistical comparison of the client's data with respect to data from a large group of neuro-typical subjects.

For volumetric analyses, Quantify uses SPM12 to correct for bias-field inhomogeneities, transform the data into MNI space, segment the data, and extract volume-related metrics. For region of interest volumetric analyses, MATLAB scripts are then used to divide the brain into 107 cortical and subcortical areas derived from TD-Brodmann (Lancaster, 2000) and AAL atlases (Tzourio-Mazoyer, et al., 2002). Figure 6 shows cortical regions from the TD-Brodmann atlas. Additional AAL atlas areas of interest include the basal ganglia and other subcortical motor nuclei, the thalamus, hippocampus, amygdala, and corpus callosum. To correct for variability in overall head/brain size, values are scaled for each individual prior to statistical evaluation.

The Quantify normative database is made up of volumetric and DTI data from more than 1,000 neurotypical subjects collected as control subjects across 36 research studies at the Mind Research Network (MRN) in Albuquerque, NM.

Different physical scanners introduce some variability into volumetric measures, even when they are nominally of the same configuration. Therefore, to compare client data obtained on one scanner with normative control data obtained on a different scanner, there is a need to harmonize data metrics across sites, prior to statistical analyses. Several procedures are implemented to accomplish this. First, data acquisition parameters at the client scanner are set to be comparable to those used in the collection of the Quantify control datasets. Second, data analysis software algorithms, procedures, and processing pipeline used to assess client data and the Quantify control data are the same. Third, prior to statistical analyses, data metrics from the client scanner are harmonized to those of the MRN scanner using a human physical phantom scaling procedure (see Palacios et al., 2017; Venkatraman et al., 2015). Briefly, the same human subject was scanned on both the client and MRN machines. For volumetric scaling, a single multiplicative factor is derived for each ROI. These procedures allow client data to be validly evaluated with respect to the Quantify normative database.

MATLAB scripts are then used to perform statistical analyses. Statistical evaluation of client data is based upon comparison of client metrics with sex and age-range matched control subjects drawn from the Quantify database, typically with an age range of +/- 10 years. Standard univariate procedures are used to generate Z-scores for each individual metric.

To account for multiple comparisons, a Benjamini-Hochberg procedure with a false discovery rate of 25% is additionally applied. All procedures are performed in a completely automated and objective manner free of any bias or client information other than gender and age. Table 1, below, provides specific data processing steps:

$$(\Omega \mu^A_N \tau i f \gamma)$$

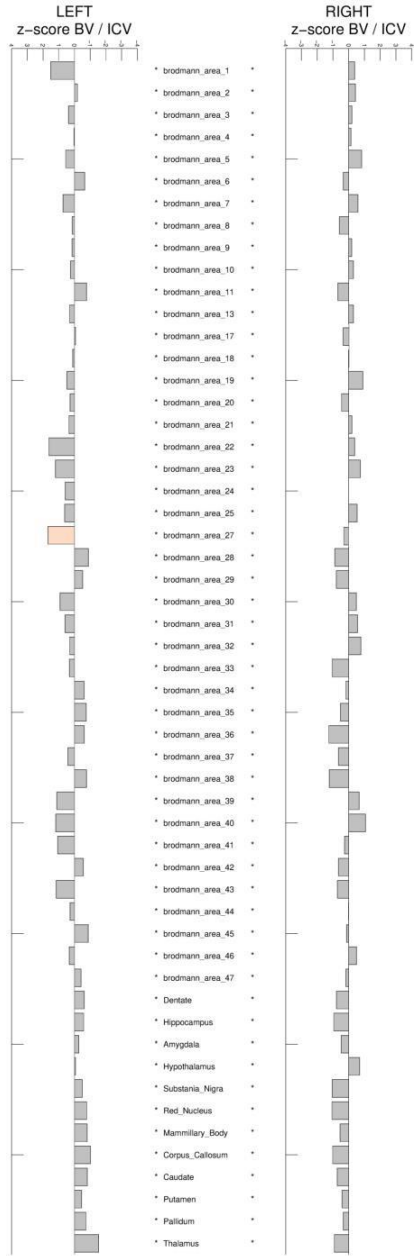
Table 1:

| VOLUMETRIC STEPS | SOFTWARE | COMMAND/ACTIVITY |
|-----------------------------------|-----------------------|---------------------------------------|
| 1. Convert Dicom data to NIFTI | MRICRON | DicomConvert |
| 2. Reorient to AC-PC | SPM12 | Reorient-to |
| 3. Grey/White/CSF Segmentation | SPM12 | Segmentation-NativeSpace |
| 4. Grey/White/CSF Segmentation | SPM12 | Segmentation-StandardSpace |
| 5. Extraction of ROI volumes | MATLAB | AAL and TD-Brodmann Templates |
| 6. Application of scaling factors | MATLAB | Site and region specific factors |
| 7. Database Comparison Statistics | MATLAB | Gender/age-range match, BH correction |
| 8. Plot Generation | MATLAB | Bar plots and scatter plots. |
| 9. Table Generation | MATLAB/Excel | Data tables |
| 10. Quality Control Check | Senior Neuroscientist | Review of data and QC metrics |
| 11. Generation of spatial maps | Senior Neuroscientist | Presently done by hand in GIMP |

Figure 1 provides an example of the bar plot outputs which show statistical results (steps 7 and 8).

$$(\Omega \mu^A_N \tau i f \gamma)$$

Neurotypical Control Subject



Example Client Dataset

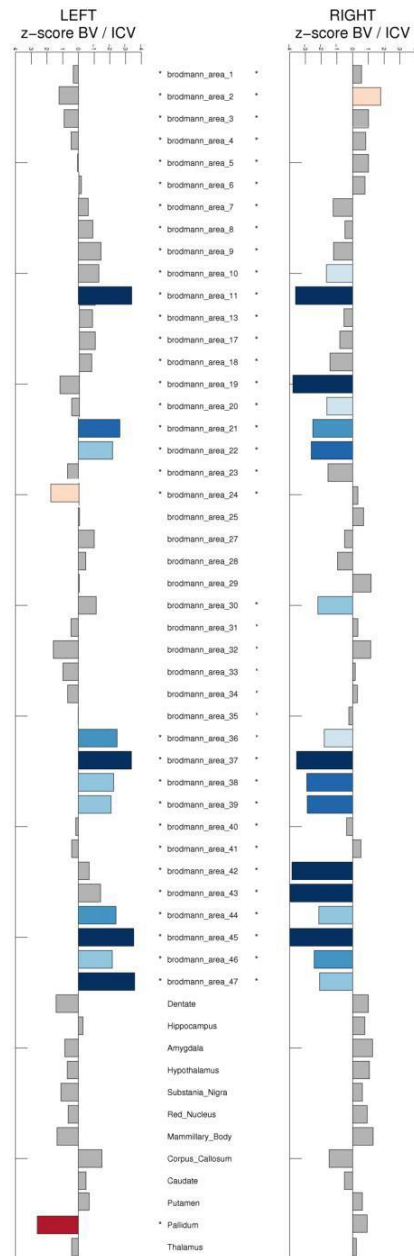


Figure 1: Main Bar Plot Outputs. Bar plots show z-score deviations from the normative database, in the indicated region. Examples are shown for a neurotypical control subject and an example client dataset. Gray indicates regions that are not statistically significant ($-1.645 < z < 1.645$; $p > 0.05$). Red indicates that a region has an atypically large volume ($z > 1.645$; $p < 0.05$). Blue indicates that a region has an atypically low volume ($z < -1.645$; $p < 0.05$). To correct for multiple comparisons, a Benjamini-Hochberg procedure is used with a False Discovery Rate of 25%. A region is ultimately considered to show a statistically significant deviation from normal if, and only if, it is abnormal on individual, isolated evaluation ($p < 0.05$) and it survives correction for multiple comparisons. Such regions are designated by an *. For the example control subject, left Brodmann area 27 has a z-score of ~ 1.71 , which is above of $p < 0.05$ threshold of $z = 1.645$, so the bar is colored red. However, this observation does not survive correction for multiple comparisons, so it is not ultimately considered to be significant (so no * is indicated). In contrast, the example client has 27 regions of significantly low volumes (blue bars with z-scores < -1.64) and three regions with abnormally high regional volumes. All of the client regions that are significant on individual isolated evaluation survive correction for multiple comparisons. That is, each of these 30 regions (27 low; 3 high) is ultimately considered to show statistically significant deviation from normal (as indicated by the *).

(Ω μ^A N τ i f γ)

Diffusion Tensor Imaging (DTI)

Diffusion tensor imaging provides insight into the integrity of white matter tracts by measuring the diffusion properties of water within the brain. For example, if you put a drop of ink in the middle of the ocean, the ink would spread (diffuse) in all directions equally. This is called isotropic diffusion. Now imagine that the ocean was filled with a bunch of pipes running in a common direction. In this case, the ink would spread in a preferred direction along the pipes, since it cannot easily penetrate into/across the pipes. When the ink diffuses in a preferential direction, this is called anisotropic diffusion. The brain is like the ocean filled with pipes – axons running within oriented fiber tracts. So, within the brain, water diffuses in preferred directions based on the local white matter anatomy. Using Diffusion Tensor Imaging the preferred direction of water movement within each voxel can be characterized by Fractional Anisotropy (FA), a value from 0 to 1, where 0 indicates isotropic diffusion and 1 indicates fully restricted diffusion in a single direction.

Typical diffusion tensor images color code the preferred directions of tracts, with color intensity indicating FA values. A process known as tractography can be used to reconstruct tract trajectories based upon FA values.

Just as was described for region-of-interest analyses in volumetric assessments, a computer algorithm can identify white matter tracts within the brain of each client and determine the average FA value for each tract. This client-specific value can then be statistically assessed with respect to a normative dataset.

For DTI, the FNIRT algorithm in combination with the DTIFIT tool in FSL is used to compute FA maps with SPM alignment to the MNI FA template and subsequent extraction of tract-based FA values within the 48 fiber tracts defined by the Johns Hopkins University MRI Atlas of Human White Matter (Mori et al., 2009). Table 2, below, provides the processing steps. For FA harmonization across scanners, a two-step procedure is used. As was the case for volumetric data, human phantoms are evaluated at both client and MRN sites. A global linear scaling factor is then derived by calculating the average FA value across all brain regions-of-interest. Then, a set of additional region-specific multiplicative factors is derived, with both the linear and multiplicative factors applied to the client data to achieve harmonization with the Quantify database. The Quantify analysis outputs include bar graphs and scatter plots similar to those provided for volumetric analyses, plus a demonstrative figure showing which tracts are abnormal.

Table 2:

| DTI STEPS | SOFTWARE | COMMAND/ACTIVITY |
|---|-----------------------|---------------------------------------|
| 1. Convert Dicom data to NIFTI | MRICRON | DicomConvert |
| 2. Eddy Current and Motion Correction | FSL | eddy.qc |
| 3. Calculation of Fractional Anisotropy | FSL | dtifit |
| 4. TBSS normalization to standard space | FSL | FSL_reg/FNIRT |
| 5. Extraction of ROI FA average FA | MATLAB | JHU template |
| 6. Application of scaling factors | MATLAB | Site and region specific factors |
| 7. Database Comparison Statistics | MATLAB | Gender/age-range match, BH correction |
| 8. Plot Generation | MATLAB | Bar plots and scatter plots. |
| 9. Table Generation | MATLAB/Excel | Data tables |
| 10. Coloring of abnormal tract ROIs | Custom Code | |
| 11. Quality Control Check | Senior Neuroscientist | Review of data and QC metrics |

$$(\Omega \mu^A_N \tau if\gamma)$$

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