

Identification of Fibers at Risk for Degeneration By Diffusion Tractography in Patients at
High Risk for MS after a Clinically Isolated Syndrome

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This work supported by grants from the National MS Society (RG 3307-A-1), the
National Science Foundation (CCR-0086065), and General Electric Medical Systems

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Key Words: Multiple Sclerosis; Diffusion; tractography; neuronal degeneration; MRI;
inflammation; Wallerian degeneration

Abstract:

Focal inflammatory/demyelinating lesions are thought to be the source of Wallerian degeneration or other injury to local, transiting fiber tracts in the brain or spinal cord in MS. In theory, a methodology establishing connections between focal demyelinating lesions and intersecting fibers would permit explicit analyses of the pathology of secondary fiber injury. A strategy is described and feasibility demonstrated in three patients with a clinically isolated syndrome and positive MRI (at risk for MS). The strategy utilizes streamtube diffusion tractography to identify neuronal fibers that intersect focal lesion and pass through a region of interest, in this case corpus callosum, where distal (to focal lesion) interrogation can be accomplished. A new class of tissue is thereby defined called fibers-at-risk (FAR) through connectivity, which is distinct from the abnormal appearing white matter and comprises an impressive fraction of NAWM in the early stages of disease.

Introduction:

While multiple sclerosis is principally considered an inflammatory-demyelinating disease, acute focal inflammatory MS lesions can harbor substantial axonal injury that includes axonal transection [1, 2]. There is increasing literature that suggests these focal lesions are the source of Wallerian degeneration or other injury to local, transiting fiber tracts. Evidence for secondary degeneration includes visualization of high signal intensity bands on conventional T2-weighted imaging exit from inflammatory lesions [3, 4], in vivo MRI diffusion measures of corpus callosum injury [5-7], and direct measures of fiber loss at autopsy [8]. In addition, empty myelin cylinders were observed by confocal microscopy in the cervical spinal cord weeks after a focal brain stem inflammatory–demyelinating event [9].

In theory, a methodology establishing connections between focal demyelinating lesions and intersecting fibers would permit explicit analyses of the pathology of secondary fiber injury. Unfortunately, by conventional two and even three dimensional MR imaging, the relationship between focal MS lesions and fiber pathways cannot be accurately determined, and connectivity estimates based on imaging may be misleading. Here we report a strategy to establish connectivity between focal demyelinating lesions and distant fiber tracts applied in patients at high risk for MS after a clinically isolated syndrome (CIS). We demonstrate the feasibility of this approach and establish a new class of tissue we call fibers-at-risk (FAR) through connectivity that can be comprehensively interrogated by multiple quantitative MRI methodologies.

Methods

The feasibility of this approach is demonstrated in the first three consecutive patients recruited after informed consent into a prospective longitudinal study of neuronal tract degeneration in early MS. Entry requirements included a CIS involving the optic nerve, spinal cord, or brainstem/cerebellum, and a positive clinical MRI (two or more T2-hyperintense lesions, one periventricular and/or ovoid, with a minimum diameter of 3mm) placing these patients at high risk for ongoing demyelination [10]

A standardized baseline MRI was acquired at 3T with a whole body instrument. MRI sequences included: (i) axial 3mm thick non-gapped proton density/T2 fast spin echo (FSE) with TR 4000 TE_{1,2} 14,84, echo train (ET) 8, field of view (FOV) 22 x 22 cm, matrix 192 x 256 and 2 excitations (ii) sagittal T2-weighted FSE series with 3mm thick non-gapped slices with TR/TE 6000/110, ET 15, FOV 22x 22 cm, matrix 256 x192, 1 excitation (iii) diffusion tensor imaging in the axial plane using an echo-planar sequence with TR/TE 6075/69.8-minimum, FOV 28 x 28 cm, matrix 160 x160, 2 excitations. The maximal b-value was 1000 and 25 gradient directions were acquired. For the diffusion tensor sequence the slice thickness was 5.1 mm, run three times, each series shifted from the prior by 1.7 mm (iv) a 3D gradient-echo magnetization transfer (MT) pulse sequence was acquired in the sagittal plane with 3mm partitions, TR/TE/flip angle 50/2.4(fr)/15 degrees, FOV 22x22, matrix 256 x192, and one excitation. The MT pulse was a Fermipulse (used to minimize SAR) with pulse width of 8 ms and a peak B₁ field of 9.24 μ T, for a flip angle of 670°, applied 1200 Hz from the center frequency. The MT pulse

was applied for every slice encoding step, but only for the middle 30% of the phase encode steps to avoid exceeding SAR restrictions.

Total brain T2-hyperintense lesions were determined using the axial proton density/T2 image pairs using AAMI (Analysis Application for Medical Imaging), an in-house semi-automated analysis application written in IDL. The abnormal appearing white matter (AAWM) fraction of the corpus callosum was based on the central sagittal T2-weighted image using AAMI and expert confirmation. The normal appearing white matter of corpus callosum (NAWM) was defined on the same slice as the full corpus callosum minus the AAWM fraction on the same image. Corpus callosum area was determined based on lesion/edge-gradient segmentation routines from the same central slice. Segmented data was registered to B_0 space via affine transformations optimized by a mutual information algorithm.

The diffusion tensor was fitted from the acquired diffusion weighted images for each voxel in the brain using a non-linear approach [11]. Streamtubes and streamsurfaces were generated for visualizing regions of linear diffusion anisotropy area and planar diffusion anisotropy area, respectively [12, 13]. For analyzing affected neural fibers we used only the streamtubes model. A full set of streamtubes was first determined that represents coherent white matter tracts extending from a dense set of seed points following the direction of fastest diffusion (the principal eigenvector of the diffusion tensor). The extension was terminated when low linear diffusion or a data boundary was encountered, or the path curves excessively. This process resulted in about 500,000 streamtubes, which

were further culled based on length, average diffusion anisotropy, and redundancy [13], leaving about 2000 streamtubes (Fig 1). Next, the subset of streamtubes with seed points within the 3D coordinates of T2-hyperintense lesions was retained. Finally the set of streamtubes was further culled to retain only those which intersect the corpus callosum.

Streamtube tractography data (acquired using the echo-planar technique) and MTR data (acquired by 3D gradient echo) were registered to the T2 space via affine transformation optimized by a mutual information algorithm. MTR values for FAR regions in the corpus callosum were calculated based on $MTR = \frac{[Signal\ Intensity\ MT_{pre} - Signal\ Intensity\ MT_{post}]}{[Signal\ Intensity\ MT_{pre}]} * 100$, with MT pre and post the sequences prior to and after addition of the MT pulse.

Results:

Patient characteristics are summarized in Table 1. All three patients were treated after diagnosis of CIS and having a high-risk for MS based on a positive MRI (Patients 1-3) and oligoclonal bands (Patient 1). Over a 1.5 year follow-up, none have had a second clinical attack, but all three showed evidence for ongoing demyelinating activity based on follow-up MRI.

Figure 1 (top) provides representative T2-weighted images showing the number and distribution of focal T2-hyperintense lesions located within the central brain slices. The relationship of these lesions to fiber pathways is completely indeterminate. The culling process to obtain the set of limited neuronal FAR is summarized for patient 1 in Figure 1. The bottom row left image shows the still complex 3D set of ~ 2000 (simplified for visualization) streamtubes and streamsurfaces that are derived based on the full data set. The bottom row middle image is a considerably more interpretable visualization restricted to fibers intersecting with focal T2-hyperintense lesions, a broad group of fibers at risk for degeneration (FAR). The middle row right image shows that set of fibers intersecting T2-hyperintense lesions which pass through corpus callosum, a more restricted set of FAR.

Figure 2 shows the subset of FAR passing through corpus callosum in patients 2 and 3. A lateral view selected from the 3D visualization image set in patient 2 shows that the technique is capable of distinguishing fibers with a high degree of specificity based on

their anatomic course. This would not be possible by visualizing 2D or even 3D conventional images since the fibers of all pathways including corpus callosum follow complex and curving positions in space.

Figure 3 shows representation of the FAR in corpus callosum in the three patients relative to the NAWM and AAWM. In all cases, while focal MS lesions were common in parasagittal slices including corpus callosum (not shown), focal lesions were relatively rare in the true central slice of corpus callosum. FAR through corpus callosum accounted for 21 %, 10 %, and 30 % of the total corpus callosum mid-sagittal area in patients 1, 2, and 3, respectively. In contrast, AAWM accounted for a smaller fraction of central corpus callosum area at 4.1%, 0 % and 8.7 %, respectively, in patients 1, 2, and 3. These data were part of a large set of planned analyses in a prospective study and were not analyzed for statistical significance. However on preliminary examination, MTR values appear abnormally low in the AAWM fraction, and relatively normal for the FAR and NAWM fractions.

Discussion:

The streamtube diffusion tractography strategy described here provides a mechanism for identifying voxels in the brain that contain neuronal fibers that are at risk for injury through secondary neuronal degeneration. FAR are distinct from tissue at risk by focal inflammatory/demyelinating pathology, and distinct from the NAWM fraction of tissue. Alternative approaches to relate focal MS lesions to related fiber pathways are limited by the necessity to assume anatomical relationships. Our studies show that even for the relatively well oriented inter-hemispheric fibers of the corpus callosum, fibers gracefully curve in 3D space across and through slices which makes assumptions about spatial relationships inaccurate at best.

This pilot study provides proof-of-concept that the tractography approach provides a unique perspective into the early MS pathology. For example, we can visualize the full extent of potentially injured FAR in the corpus callosum. The FAR are distinct from the AAWM, although not unexpectedly there is some overlap. FAR in these “at-risk for” MS cases accounted for 10-30 % (mean 20%) of the central sagittal corpus callosum area. Based on estimates of about 160 million fibers crossing the adult corpus callosum [14], our series suggests a potential mechanism that places about 32 million fibers at risk already at the time of diagnosis or time of first recognized risk for MS. As the corpus callosum undergoes extensive volume loss early in the disease [15,16], ultimately associated with a striking decrease in neuronal fiber count[8], the large number of FAR if proven to be injured, may provide insight into the early injury process.

While we do not know at this time if and how the FAR in corpus callosum are injured, there are several lines of evidence suggesting that corpus callosum fiber injury is initiated very early in the disease. Transcallosal bands have been observed based on signal change on T2-weighted imaging in patients at high risk for MS at the time of a CIS [3]. These changes are similar in signal pattern to fiber injury in the corticospinal tract [4]. While these conventional T2-evident hyperintensities may represent the most severe of a broad range of injuries, those lesions already evident by T2-weighted imaging are likely to be but a small fraction of the total abnormality that would be detected by quantitative MRI methodologies [5-7, 17, 18].

How much of this injury in the NAWM of the corpus callosum or other NAWM is due to focal microscopic pathology as opposed to secondary degeneration is not known as the quantitative MRI techniques are not specific for the pathology of neuronal degeneration. Loss of diffusion anisotropy, and particularly increased diffusivity perpendicular to the principal fiber direction, findings which are detected in corpus callosum, are believed to be “signatures” of Wallerian degeneration [7, 19, 20]. In vivo, a regional relationship is found between reduced fractional anisotropy and increased mean diffusivity in the corpus callosum and the separate lesion load in the cerebral lobes [5,6], and atrophy of corpus callosum is correlated with brain T2 lesion load [21,16]. In studies of fixed autopsy specimens, there is a correlation between the regional load of macroscopic lesion in white matter and axonal density and number in the corresponding projection area in corpus callosum [8].

While we do not see a major decrease in MTR in the FAR in these three cases this small pilot study was not designed for that purpose, and it is interesting to note that the MTR of the NAWM was likewise not obviously reduced. The relatively normal MTR in FAR and NAWM may be due to a combination of evaluation at an early stage in disease and the small sample size. Future analyses including evaluation based on the longitudinal series will address the sensitivity of MTR and measures based on diffusion tensor (fractional anisotropy and diffusivity) in this population. It is apparent that noise in the image data and the small size of the tissue fractions of interest may require further optimization of the MRI acquisition, even for studies at 3T where the added signal-to-noise is a benefit compared to conventional imaging ($\leq 1.5T$). With regard to the FAR, a potential source of error in our analyses is the possibility for artifactual drop-out of visualized fibers related to the known decrease in fractional anisotropy that accompanies the focal pathology in MS [22,23]. It is encouraging however to note that fibers can be readily traced through and from the surfaces of the focal pathology. Future studies will address in quantitative manner issues of fiber dropout and the characteristics of the focal pathology associated with more severe fiber dropout.

This FAR strategy is applicable to any neuronal tract that can be imaged with high quality diffusion tensor and structural MRI. In theory, a methodology establishing connections between focal inflammatory and/or demyelinating lesions and intersecting fibers should benefit explicit analyses of the pathology of fiber injury related to individual focal lesions or a composite of all lesions. Such studies could improve our

understanding of the temporal relationships of injury and its consequences, the heterogeneity of injury and disease, and potentially provide a basis for assay of neuroprotective therapy.

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Figures:

Fig. 1. Case 1. Top. Five representative T2-weighted slices show multiple small T2 hyperintense lesions in the vicinity of corpus callosum. Bottom. Culling procedure. Left. Streamtube visualization (view from above) including the full set of streamsurfaces (green) and streamtubes (red) in relation to focal T2-hyperintense lesions (yellow) and ventricle (blue). Middle. Visualization culled to include streamtubes that intersect focal T2-hyperintense lesions. Right. Set of streamtubes fully culled to include fibers intersecting focal T2-hyperintense lesion and crossing corpus callosum. Saturation of colors are used to represent strength of linear anisotropy (redness).

Fig. 2. Fully culled images for Case 2 (left) and Case 3 (right). Bottom is an intermediate culled lateral visualization from Case 2. Note the complex curvature of callosal fibers in both the superior-inferior directions and anterior-posterior direction, which can be distinguished from non-callosal fiber tracts running anterior-posterior.

Fig. 3. Co-registered data on sagittal T2 –weighted mid-sagittal images show the tissue classes corpus callosum (yellow), abnormal appearing white matter (green) and fiber-at-risk (FAR) in red. Overlapping pixels represented by intensity variation. Note that most of the FAR is not associated with the AAWM fraction. In chart (bottom), the relative fraction of FAR and AAWM are shown as well as MTR values for the three cases.

Table 1 . Patient Characteristics

| Patient | Age Gender | CIS Presentation | DIS | Brain T2 lesion Volume (cc) | CDMS | New MR lesions on follow-up? | Treated |
|---------|---------------|-----------------------------------|-----|--------------------------------------|------|------------------------------------|---------|
| 1 | 38 M | Optic neuritis OCB positive | + | 5.83 | No | At 3m | Yes |
| 2 | 33 F | Partial transverse myelitis | + | 2.99 | No | At 1m,3m | Yes |
| 3 | 26 F | Optic neuritis | + | 10.69 | No | At 1 y | Yes |

OCB = oligoclonal bands

DIS= fulfill McDonald criteria for dissemination in space

CDMS =clinically definite MS based on second attack by 1.5 year follow-up

Treated= Treatment with immunomodulatory therapy

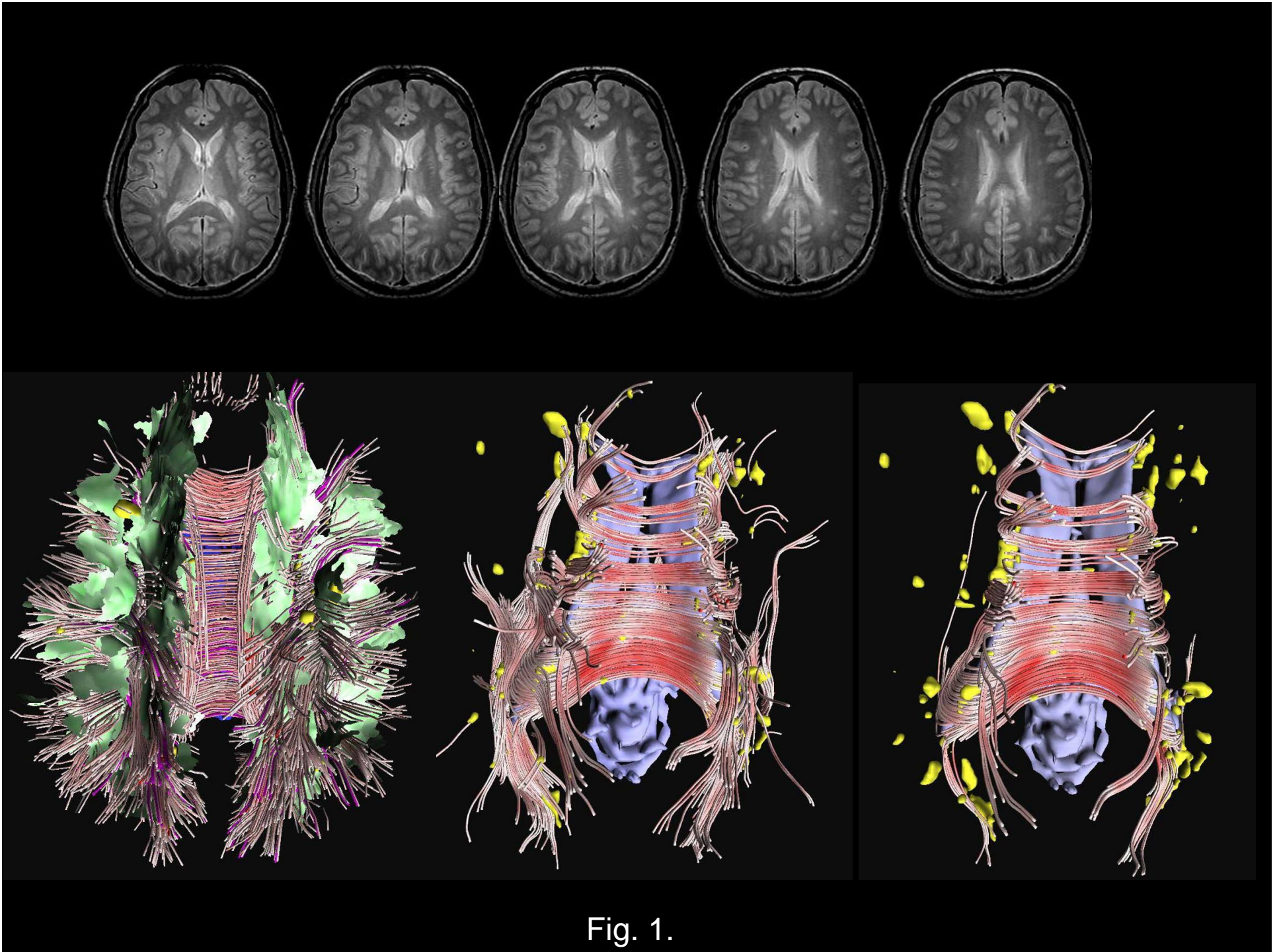
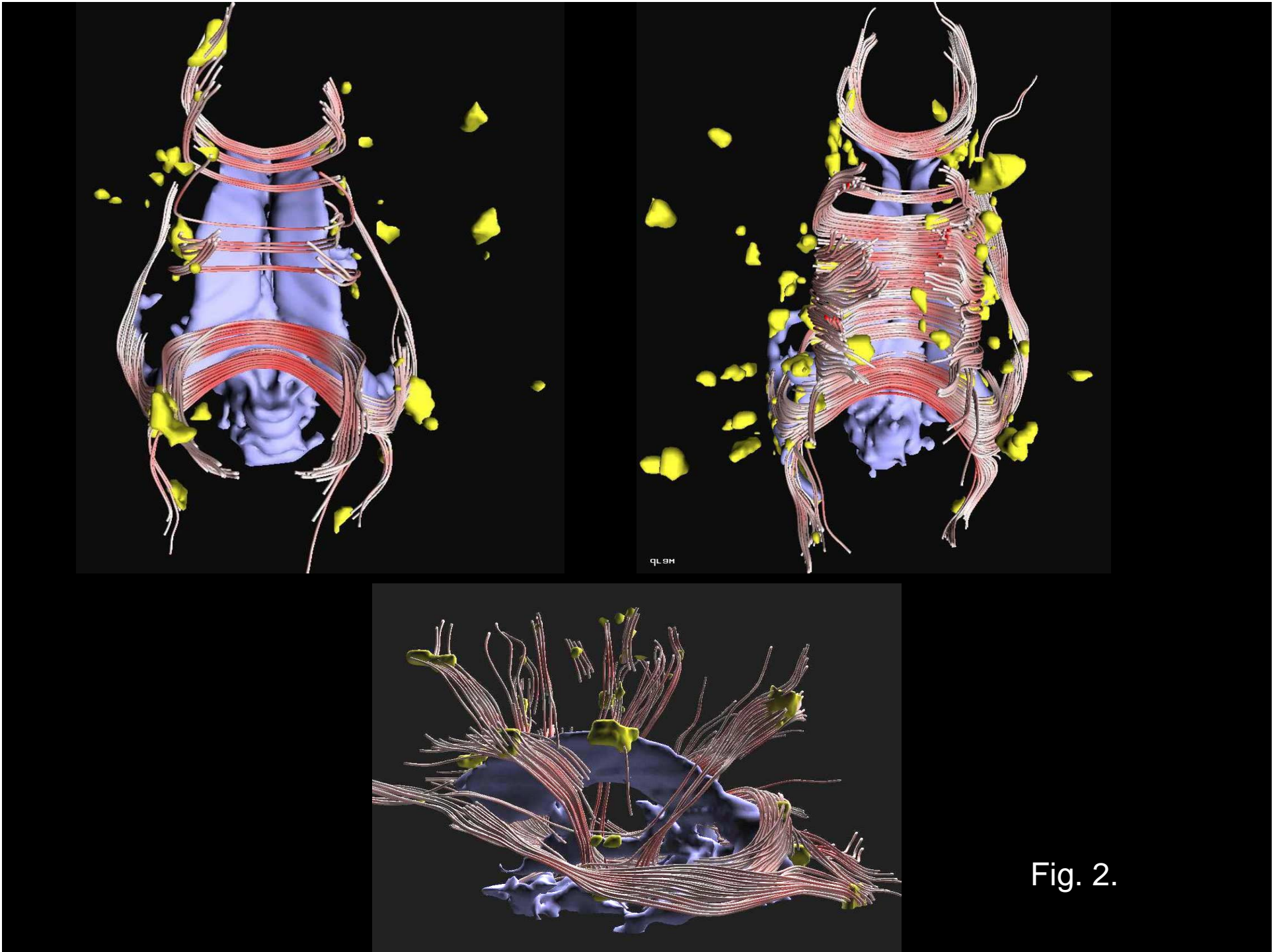
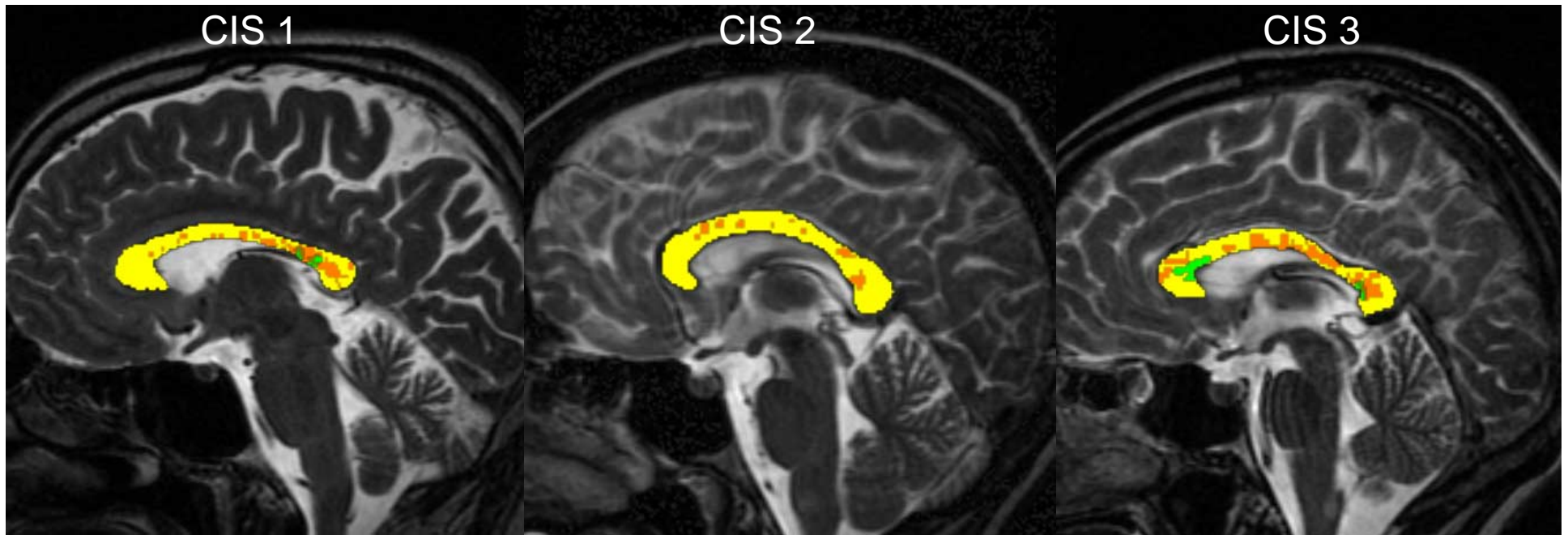


Fig. 1.





| ID | Corpus Callosum Area | | Mean MTR | | | |
|-------|----------------------|--------|---------------------|---------------|---------------|--------------|
| | FAR % | AAWM % | All Corpus Callosum | NAWM Fraction | AAWM Fraction | FAR Fraction |
| CIS 1 | 21.0 | 4.1 | 48.94 | 49.06 | 45.98 | 50.57 |
| CIS 2 | 10.0 | 0 | 49.55 | 49.55 | | 49.78 |
| CIS 3 | 30.0 | 8.7 | 48.53 | 48.55 | 48.36 | 50.63 |

Fig. 3.