Connectivity-aware sectional visualization of 3D DTI volumes using perceptual flat-torus coloring and edge rendering

Abstract

*We present two new methods for visualizing cross-sections of 3D diffusion tensor magnetic resonance imaging (DTI) volumes. For each of the methods we show examples of visualizations of the corpus callosum in the mid-sagittal plane of several normal volunteers. In both methods, we start from points sampled on a regular grid on the cross-section and, from each point, generate integral curves in both directions following the principal eigenvector of the underlying diffusion tensor field. We compute an anatomically motivated pairwise distance measure between each pair of integral curves and assemble the measures to create a distance matrix. We next find a set of points in a plane that best preserves the calculated distances that are small—each point in this plane represents one of the original integral curves. Our first visualization method wraps this planar representation onto a flat-torus and then projects that torus into a visible portion of a perceptually uniform color space (L*a*b*). The colors for the paths are used to color the corresponding grid points on the original cross-section. The resulting image shows larger changes in color where neighboring integral curves differ more. Our second visualization method lays out the grid points on the cross section and connects the neighboring points with edges that are rendered according to the distances between curves generated from these points. Both methods provide a way to visually segment 2D cross sections of DTI data. Also, a particular contribution of the coloring technique used in our first visualization method is to give a continuous 2D color mapping that provides approximate perceptual uniformity and can be repeated an arbitrary number of times in both directions to increase sensitivity.*

1. Introduction

Diffusion-Tensor Magnetic Resonance Imaging (DTI) enables the exploration of fibrous tissues such as brain white matter and muscles non-invasively *in-vivo*. It exploits the fact that water in these tissues diffuses at faster rates along the fibers than orthogonal to them fibers. However, the multivalued nature of DTI data poses challenges in visualizing and understanding the underlying structures. Integral curves that represents neural pathways by showing paths of fastest diffusion are among the most common information derived from DTI volumes. They are generated tracking the principal eigenvector of the underlying diffusion tensor field in both directions. They are often visualized with streamlines or variations of streamlines (streamtubes and hyperstreamlines) in 3D. In this paper, we present two new methods for visualizing cross sections of DTI volumes that preserve the 3D out-of-plane connectivity information typically conveyed by the integral curves (see 1). Slice-based 2D visualizations of scientific data are generally effective, fast and synoptic [CM02, SWD04]. Also, looking at 2D cross-sections is still the most common practice by far among scientists and physicians for data exploration. Furthermore, there is some evidence that incorporation of 2D cross-sections in 3D visualizations of medical data sets data is preferred [DJK*06].

For each of the visualization methods we show examples of visualizations of the corpus callosum in the mid-sagittal plane of one normal volunteers. The corpus callosum is the largest white matter fiber bundle in the brain and a target for clinical and neuroscience research into normal developmental vs. pathological changes in white matter integrity across the lifespan and the functional correlates of those changes. Distinct cross-sectional regions of the corpus callosum may contain fibers that subserve specific cognitive or behavioral functions mediated by the cortical regions to which they project. Proxy measures (e.g., thickness, volume, area, shape) of the health of these cross-sectional regions may correlate with measures of the cognitive and behavioral functions they subserve. In fact, the corpus callosum has been shown to differ on such measures by handedness , gender, and age as well as in diseases such as Alzheimer's disease and schizophrenia [HJL*04].

2. Related Work

Mapping colors to data values is a fundamental operation in scientific visualization. Previous work based on empirical studies addressed the problem of generating perceptually effective colormaps [War88, Hea96, LH92, KRC02]. We use the CIE L*a*b* color space that is a perceptually uniform (approximately) color space proposed by the Commission Internationale de l'Eclairage (CIE) in 1976. A color space is said to be perceptually uniform if the perceptual difference between any two colors in just noticeable difference (JND) units is equal to the Euclidean distance between the two colors in that color space. Several different geometric models, including line, plane, cone, cylinder, and B-spline surfaces have been proposed for univariate, bivariate or trivariate colormapping [Pha90, Rob88]. We extend the earlier models

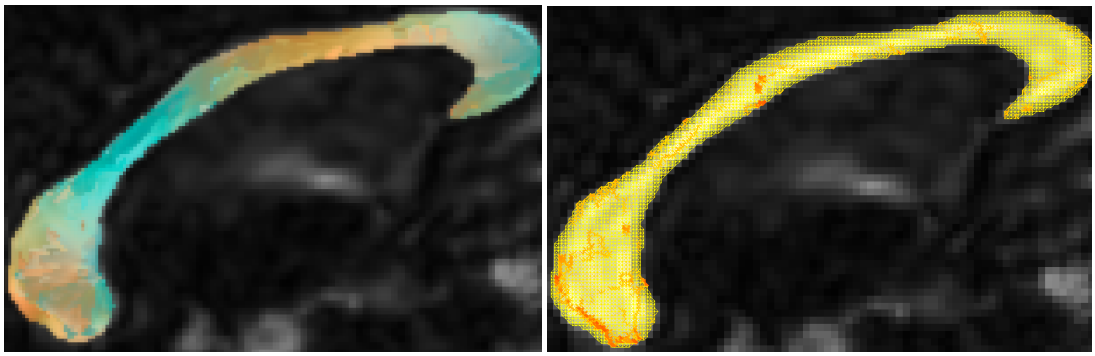

Figure 1: *Flat-torus coloring (left) and edge rendering (right) visualizations of the mid-saggital plane of the corpus callosum in a normal person's brain*

by introducing the flat-torus model to give a continuous 2D color mapping that is approximately uniform and that can be repeated an arbitrary number of times in both directions to increase sensitivity. Integral curves generated from DTI volumes have been visualized generally with streamlines in 3D with different geometric (i.e., hyperstreamlines, streamtubes, etc.) and coloring combinations. It is also common practice to juxtapose streamtubes with 2D cross-sections of the volume data. Volume visualization of DTI data included isosurface extraction and volume rendering. Previous cross-sectional visualizations of DTI included mapping glyphs (box, ellipsoid and superquadratic) and colors to tensor voxels [ZKL04]. Pajevic *et al.* proposed methods to colormap principle eigenvectors of tensor voxels on DTI cross-sections using different color spaces, including perceptually uniform CIE L*u*v color space. The authors point the potential limitations due to irregularity of L*u*v space. Our flat-torus model addresses some of the limitations discussed in this work [PP99].

3. Methods

In both visualization methods presented here, we start from points (seeds) sampled on a regular grid on the cross-section and, from each point, generate integral curves in both directions following the principal eigenvector of the underlying diffusion tensor field. We compute an anatomically motivated pairwise distance measure between each pair of integral curves and assemble the measures to create a distance matrix. The distance matrix is utilized by the both methods to convey the out-of-plane connectivity information. We explain how we measure distances between the integral curves to construct the distance matrix in the next section.

3.1. Distance Measure Between Integral Curves

Integral curves generated from DTI volumes are solutions to the following first-order differential equation:

$$\frac{dC}{ds} = \vec{v}_1(C(s)) \tag{1}$$

where s parameterizes the curve and v_1 is the principal eigenvector at the point $C(s) = (x(s), y(s), z(s))$. We compute the integral curve $C(s)$ passing through a given seed point $C(0)$ (initial conditions) by integrating the above equation for $s > 0$ and $s < 0$ (i.e., both directions from the seed point). There have been different distance measures proposed for integral curves generated from DTI volumes. In the current work we adapt a measure proposed by Zhang *et al.* with a slight modification [ZDL03]. Note that our measure does not necessarily satisfy triangle inequality therefore it is not a metric. Given any two integral curves C_i and C_j that are represented as polylines with m and n vertices respectively (like the ones shown in Figure 2), we first find mean distances d_{ij} and d_{ji} then, determine the maximum of these two distances as the distance D_{ij} between the two curves:

$$d_{ij} = \frac{\sum_{k=1}^{m} dist(C_i^k, C_j)}{m} \tag{2}$$

$$d_{ji} = \frac{\sum_{k=1}^{n} dist(C_j^k, C_i)}{n} \tag{3}$$

$$D_{ij} = D_{ji} = max(d_{ij}, d_{ji}) \tag{4}$$

The function $dist(p, C)$ returns the shortest Euclidean distance between the point p and curve C. We compute distance between each pair of integral curves as we denoted and assemble the measures to create a distance matrix. The distance matrix is a real positive symmetric matrix with zeros along the diagonal.

3.2. Flat-Torus Coloring

The goal of our first method is to reflect the boundaries in distance differences in the data as perceptual boundaries. For

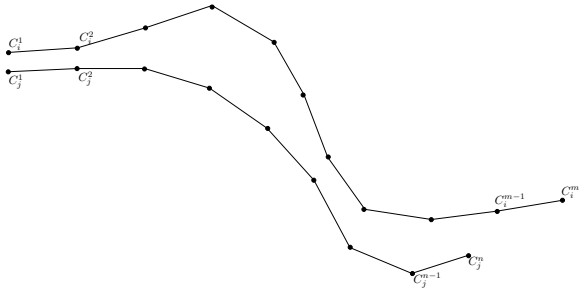

Figure 2: *Polyline representations of two integral curves C_i and C_j*

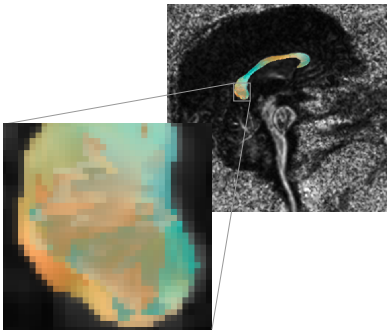

Figure 3: *Flat-torus coloring of the mid-sagittal of the corpus callosum in a normal person's brain*

this, we lay out the seed points on the plane and adjust their positions using a simple mass-spring based optimization algorithm so that the calculated distances between their associated integral curves are locally preserved best. Figure 4 illustrates how seed point coordinates change after running the optimization algorithm. We coordinate-transform the adjusted points using principle component analysis (PCA) to have a succinct representation. Finally, we wrap this planar representation onto a flat torus and then projects that torus into a visible portion of the CIE L*a*b* space. A flat-torus in 4-space is a Cartesian product of two circles in R^2. It can be obtained by a mapping $W : R^2 \rightarrow R^4$ such that

$$W(x,y) = (u,v,z,t) = (r_1 \cos x,\ r_1 \sin x,\ r_2 \cos y,\ r_2 \sin y) \tag{5}$$

—where $r1$ and $r2$ are the radii of the circles. The flat-torus has 0 Gaussian curvature everywhere (i.e., is a developable surface), therefore a plane can be wrapped around it without distortion [dC76].

We project the flat-torus to the visible partition of L*a*b* color space, centered at (L_o, a_o, b_o) as follows.

$$L* = L_o + t \tag{6}$$

$$a* = a_o + r_1 + u + z \tag{7}$$

$$b* = b_o + r_1 + v + z \tag{8}$$

The resulting images show larger changes in color where neighboring integral curves differ more. One of the advantages of using this flat-torus projection is that we can adjust the sensitivity of the color mapping by rescaling point plane and wrapping around the two circles continuously.

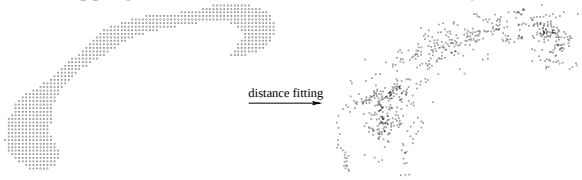

Figure 4: *Seed points are adjusted so that Euclidian distances between the points on the plane reflect the distances between their associated integral curves.*

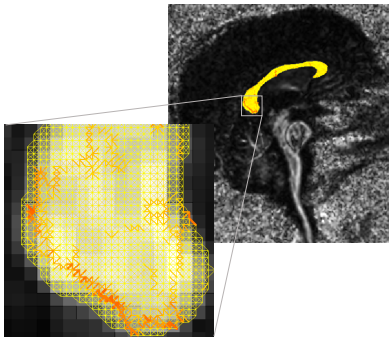

Figure 5: *Edge rendering visualization of the mid-sagittal plane of the corpus callosum in a normal person's brain.*

3.3. Edge Rendering

Our second visualization method lays out the grid points on the cross section and connects the neighboring points with edges that are rendered according to the distances between curves generated from these points. Note that we sample seed points on a rectilinear grid where the vertical and horizontal distances between the grid points are equal to δ. We define the seed points X_i and X_j to be neighbors if $||X_i - X_j||_2 = \delta$ or $||X_i - X_j||_2 = \delta\sqrt{2}$ (i.e., a seed point can have maximum 8 neighbors). Edges are drawn redder in color and thicker where neighboring seed points' integral curves differ more.

4. Results and Discussion

Figure 3 and 5 show the visualizations of the same normal person's corpus callosum with close-up views of the same region. Notice the correspondance between regions in flat-torus coloring and edges in edge rendering. More results from two different persons' DTI brain data sets are shown in Figure 6. It is important to note that the perceptual uniformity in our color mapping is an approximation due to the flat-torus to the L*a*b* space projection distortions. While flat-torus cannot be mapped to the L*a*b* space isometrically, it could be possible to have a projection that preserves perceptual uniformity better than the one presented in this paper.

5. Conclusions

We have presented two new cross-sectional visualization methods. The primary strength of both methods is providing a compact and contextual visualization by bringing higher dimensional connectivity information onto 2D plane which is effective and familiar to practitioners. We have applied them to visually segment the mid-saggital cross-section of the corpus callosum in the brain. Feedback from neuroscientist collaborators suggests that our visualization methods can be useful in identification of smaller caliber anatomically or functionally related white-matter structures, particularly those that are contained within large bundles or fasciculi that project to multiple areas. Flat-torus coloring is a new geometric model for bivariate color mapping allowing approximately uniform and that can be repeated an arbitrary

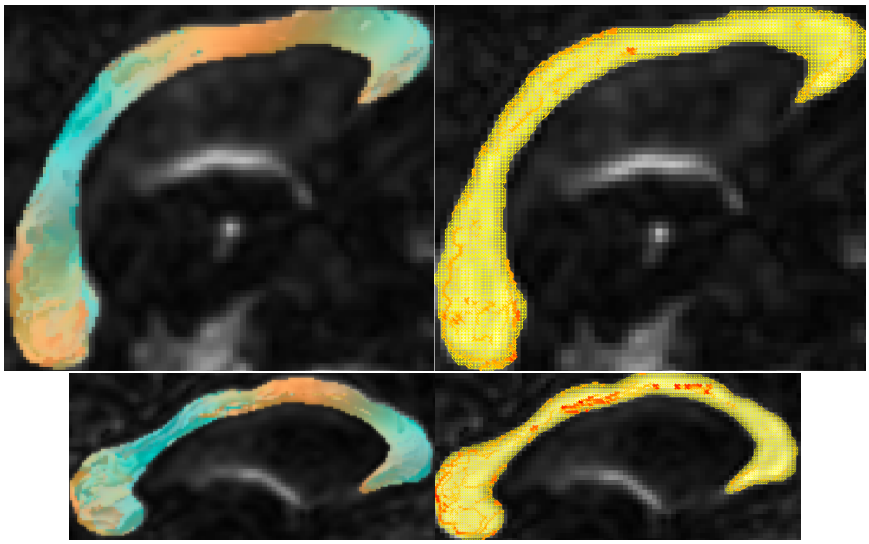

Figure 6: *Flat-torus coloring (left) and edge rendering (right) visualizations of the mid-saggital plane of the corpus callosum in two normal persons' brains.*

number of times in both directions. The underlying of idea of this work can be extended to visualization of any other vector, tensor or multi-scalar data volumes.

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