Illustrative uncertainty visualization of DTI fiber pathways

Citation for published version (APA):
Brecheisen, R., Platel, B., Haar Romenij, ter, B. M., & Vilanova, A. (2013). Illustrative uncertainty visualization of DTI fiber pathways. The Visual Computer, 29(4), 297-309. https://doi.org/10.1007/s00371-012-0733-9

DOI:
10.1007/s00371-012-0733-9

Document status and date:
Published: 01/01/2013

Document Version:
Accepted manuscript including changes made at the peer-review stage

Please check the document version of this publication:
• A submitted manuscript is the version of the article upon submission and before peer-review. There can be important differences between the submitted version and the official published version of record. People interested in the research are advised to contact the author for the final version of the publication, or visit the DOI to the publisher's website.
• The final author version and the galley proof are versions of the publication after peer review.
• The final published version features the final layout of the paper including the volume, issue and page numbers.

Link to publication

General rights
Copyright and moral rights for the publications made accessible in the public portal are retained by the authors and/or other copyright owners and it is a condition of accessing publications that users recognise and abide by the legal requirements associated with these rights.

• Users may download and print one copy of any publication from the public portal for the purpose of private study or research.
• You may not further distribute the material or use it for any profit-making activity or commercial gain
• You may freely distribute the URL identifying the publication in the public portal.

If the publication is distributed under the terms of Article 25fa of the Dutch Copyright Act, indicated by the “Taverne” license above, please follow below link for the End User Agreement:
www.tue.nl/taverne

Take down policy
If you believe that this document breaches copyright please contact us at:
openaccess@tue.nl
providing details and we will investigate your claim.
Abstract Diffusion Tensor Imaging (DTI) and fiber tracking provide unique insight into the 3D structure of fibrous tissues in the brain. However, the output of fiber tracking contains a significant amount of uncertainty accumulated in the various steps of the processing pipeline. Existing DTI visualization methods do not present these uncertainties to the end-user. This creates a false impression of precision and accuracy that can have serious consequences in applications that rely heavily on risk assessment and decision-making, such as neurosurgery. On the other hand, adding uncertainty to an already complex visualization can easily lead to information overload and visual clutter. In this work, we propose Illustrative Confidence Intervals to reduce the complexity of the visualization and present only those aspects of uncertainty that are of interest to the user. We look specifically at the uncertainty in fiber shape due to noise and modeling errors. To demonstrate the flexibility of our framework, we compute this uncertainty in two different ways, based on (1) fiber distance and (2) the probability of a fiber connection between two brain regions. We provide the user with interactive tools to define multiple confidence intervals, specify visual styles and explore the uncertainty with a Focus+Context approach. Finally, we have conducted a user evaluation with three neurosurgeons to evaluate the added value of our visualization.

Keywords Diffusion Tensor Imaging · Fiber Tracking · Uncertainty Visualization · Illustrative Rendering · Neurosurgery Planning

1 Introduction

Diffusion is the process of random movement of water molecules over time, also called Brownian motion. Diffusion Tensor Imaging (DTI) is an imaging technique based on Magnetic Resonance (MR) that can measure diffusion of water in living tissues. By measuring the amount of diffusion in many different directions, the 3D shape of the diffusion profile can be approximated at each point in the tissue. In pure water, diffusion is unrestricted and has equal magnitude in all directions. This results in a spherical or isotropic diffusion profile. In fibrous tissues however, such as the brain white matter, the diffusion will be restricted in directions perpendicular to the fibers. This results in a more elongated or anisotropic diffusion profile. In DTI, the diffusion profile is modeled as a 3D Gaussian probability distribution using a 2nd-order tensor. In this model the tensor’s main eigenvector corresponds to the direction of greatest diffusion and is assumed to be aligned with the underlying fiber structure [1]. Based on this assumption it is possible to do streamline tracing in the main eigenvector field. In the context of DTI, this is called fiber tracking [20,26,31] and it allows reconstruction of the fibers in three dimensions. Diffusion-weighted MRI is the only imaging modality that allows to do this non-invasively and in-vivo. For this reason, it has great potential for applications that involve fibrous tissues, such as the brain white matter and muscle.

Despite its potential, the output of DTI fiber tracking contains a significant amount of uncertainty. This uncertainty is accumulated in the various stages of the DTI processing pipeline. A schematic overview of this pipeline is illustrated in Figure 1. The data acquisition stage suffers from noise, image distortion, movement artefacts, partial volume effects and scan parameters. The modelling stage can introduce approximation errors depending on the diffusion model that is used. For example, the 2nd-order tensor model produces un-
Brecheisen et al.

Fig. 1 DTI processing pipeline with sources of uncertainties at each stage

reliable results when fibers are crossing, kissing or diverging within a voxel. The fiber tracking stage attempts to reconstruct 3D pathways from the tensor field. However, different numerical integration schemes produce different results and each fiber tracking algorithm has several user-defined parameters that can significantly affect fiber length and/or shape [5]. Finally, the visualization stage may introduce uncertainty due to the use of different illumination models or simplification of the fiber geometry.

Existing DTI visualizations largely ignore the above mentioned sources of uncertainty and their effect on the reconstructed fiber pathways. This gives an impression of certainty that can be misleading. This is unacceptable in applications such as neurosurgery, where fiber tracking may be used for surgical risk assessment and decision-making. If false positives or false negatives in the fiber tracking output are not correctly identified this can result in sub-optimal tumor resection or damage to healthy brain tissue. To deal with risks of healthy tissue damage neurosurgeons may take into account safety margins around critical brain structures [32]. However, such safety margins are mostly fixed and have no relation to the uncertainty in the underlying data. Furthermore, safety margins enclose only one out of many possible fiber configurations. Figure 2 illustrates that, due to noise and modelling errors, there are many such configurations possible, each one slightly different. It is clear from this example that a fixed safety margin may not adequately cover the possible variations in fiber shape.

With the visualization framework described in this paper we attempt to visually communicate to the neurosurgeon that the fiber tracking algorithm they use may produce a sub-optimal reconstruction of the tracts of interest. We are specifically focusing on variations in the output due to noise and modelling errors. Such information can be captured by different probabilistic algorithms but intuitively showing this information is not a trivial task. Even without considering uncertainty, diffusion tensor data presents considerable visualization challenges. For this reason, each tensor is often reduced to a single vector describing the principal direction of diffusion. Streamline visualization can then be used to show pathways through the tensor field. This gives a good impression of the global structure of fiber tracts. However, if there are too many streamlines this approach can lead to highly cluttered visualizations, such as illustrated in Figure 2 on the right. Probabilistic fiber tracking algorithms generally sample thousands of potential pathways in order to compute connection probabilities between different regions of the brain. In that case, it is no longer feasible, nor informative, to render each individual streamline.

To deal with this issue, we propose Illustrative Confidence Intervals, a rendering technique based on illustrative silhouettes and outlines that provides information about the variation in fiber pathways while at the same time reducing the visual clutter associated with standard streamline visualizations. We define a silhouette to be the interior area of an

Fig. 2 Left: single fiber obtained from original tensor volume. The yellow silhouette represents fixed safety margin. Right: 100 variations of same fiber based on wild bootstrap method [13] with same safety margin. It is clear that it does not cover the possible variations in the tensor data.
object’s shape when projected to the viewing plane. An outline is defined as the border of a silhouette. Our framework does not depend on a specific algorithm to compute the variation in fiber pathways. It only requires a set of streamlines, each associated with a so-called confidence value. A confidence value is a scalar-valued, fiber-specific measure that can be computed in different ways depending on the application. The specific measure used in our framework can be freely chosen but to demonstrate the flexibility of our approach we have selected two such measures, (1) based on fiber distance and (2) based on the connection probability between different brain regions. Our contributions are the following:

1. A processing framework that allows different methods for constructing sets of streamlines with associated confidence values. We implemented two example methods but other methods can be added easily (see Section 3).
2. A visualization and interaction framework that uses illustrative silhouettes and outlines to analyze variations in the output of a selected fiber tracking algorithm (see Section 4). A set of interaction widgets that allows easy specification of intervals and their visual styles. A Focus+Context uncertainty lens shows confidence intervals only within a user-defined region of interest. Outside this region standard fiber visualization can be used allowing for easy comparison between the two visualizations (see Sections 4.3 and 4.4).
3. A quantitative and qualitative user study based on video demonstrations, questionaires and interviews with three neurosurgeons.

2 Related Work

Visualization of uncertainty is closely related to and dependent on the method of measuring or computing uncertainty. As Figure 1 illustrates there are many different sources of uncertainty present in the DTI processing pipeline. In our current work, we focus on the uncertainty in the DTI (tensor) data caused by noise and modelling errors. In the next subsections we discuss related work in the areas of DTI uncertainty analysis, uncertainty visualization and illustrative rendering for DTI applications.

2.1 Uncertainty Analysis

Several approaches are possible to characterize the effects of noise and modelling errors on DTI fiber tracking. They can be roughly subdivided into two types: empirical methods and mathematical modelling methods. There also exist different fiber tracking methods. We divide them into deterministic algorithms, i.e. they give the same result given the same input (such as streamline tracing), and probabilistic algorithms that involve a random process. To characterize the uncertainty, an obvious, empirical approach is to repeat a diffusion-weighted imaging (DWI) scan multiple times. Due to random noise each scan will be slightly different. Repeating this often enough will give a good approximation of the uncertainty in the data. Of course, this would require hours or days of scanning. A variation of this method, called bootstrap, is to generate new datasets, by taking random samples with replacement from a much smaller set (4 to 5) of DWI scans [23]. However, this approach is still too time-consuming for clinical purposes. Wild bootstrap [13, 33, 16] is a method that generates random variations based on a single DWI scan. This method is often combined with deterministic fiber tracking to obtain information about fiber shape variations (see Section 3.1). The second type of approach to characterize uncertainty is based on mathematical modelling [14,10,2]. Klein et al. [14] generate variations of a single DWI dataset by adding increasing levels of complex Gaussian noise. A drawback of such methods is that they assume a noise model (e.g., Gaussian) which generally is a simplification of the real noise characteristics. Others take a predictive model, such as the 2nd-order tensor, and use Bayesian statistics to estimate a posterior PDF for the model parameters, including fiber orientation [10, 2]. These methods generally do not use deterministic fiber tracking but instead apply probabilistic algorithms to reconstruct fiber pathways (e.g. based on Monte Carlo simulations). Still, these methods assume a noise model and have to deal with the complexity of having to propagate this noise model through the many, often non-linear, transformations in the DTI processing pipeline. On the other hand, the explicit mathematical representation of assumptions allows a more formal way to compute uncertainty and can be applied to a wide range of diffusion models. Some authors use a combination of mathematical modelling and empirical techniques to characterize uncertainty in the data [28,3].

In this paper we make no statements about which approach for computing uncertainty is superior. Our main focus is to visualize variations in fiber pathways due to noise and modelling errors. To demonstrate our approach we have chosen the Wild Bootstrap method by Jones et al. [13] and the ConTrack algorithm by Sherbondy et al. [28]. However, our method does not depend in any way on the particular choice of algorithm used to characterize uncertainty.

2.2 Uncertainty and Illustrative Fiber Visualization

The last decade has shown a slow but growing interest in the visualization of uncertainty, especially for scalar and vector fields. Pang et al. [25] provide an extensive overview of techniques, as well as a classification scheme for describing data together with its associated uncertainty. Lodha et al.
[17] present a method for visualizing variations in streamline tracing due to different integration schemes. They provide different options for representing the uncertainty, e.g. glyphs, ribbons and motion-blurred animation. Pang [24] and Wittenbrink et al. [34] use different types of glyphs to display the effect of data uncertainty on height field measurements and the direction and magnitude of 2D vector fields.

The literature concerning uncertainty visualization for tensor fields is very limited. Variations in the tensor main eigenvector were visualized by Jones et al. [13] using cone glyphs and a modification of hyperstreamlines [7]. They compute these variations using the Wild Bootstrap method. However, they render all fibers in the same way (as in Figure 2) without giving an impression of which fibers are more reliable or reproducible than others. Furthermore, their method cannot easily be adapted to more complex fiber shapes. Enders et al. [8] compute a ‘mean’ fiber to create a closed surface hull wrapped around the fiber bundle. However, similar to Jones et al., their method cannot deal with complex fiber shapes and concave bundle cross-sections without significant simplification of the hull geometry. The wrapped hull geometry described by Chen et al. [6] suffers from similar problems. Merhof et al. [18] propose a hull generation algorithm based on smoothed isosurfaces that does not suffer from these limitations.

The visualization methods discussed so far primarily deal with noise and modelling errors. The effect of user-defined parameters on the output of DTI fiber tracking was addressed by Brecheisen et al. [5] who use a combination of coloring, linked views and interactive parameter space exploration to visualize the threshold sensitivity of streamline tracing. To improve the visualization of fiber structures, either to reduce visual clutter or provide surrounding context, a number of researchers have used illustrative techniques [9, 29, 4, 11, 27]. Weiler et al. [32] attempt to highlight fiber tract uncertainty by means of a fixed, user-defined safety margin. However, this safety margin does not take data uncertainty into account.

In this paper we propose a method that uses the output of probabilistic fiber tracking methods to generate Illustrative Confidence Intervals that do take uncertainty of the data into account. Each fiber is assigned a confidence value based on a chosen confidence measure (to be explained in the next section). A confidence interval \( [C_i, C_{i+1}] \) contains a subset of all fibers with a confidence between \( C_i \) and \( C_{i+1} \). An illustrative confidence interval is the visual representation of these fibers and shows the variation in fiber pathways due to noise and modelling error. The rendering algorithm uses a silhouette and outline representation that is based on the work of Otten et al. [22] but we extend it with additional visual styles, interaction features, Focus+Context views and anatomical context. Furthermore, we apply and evaluate our approach in the context of neurosurgical risk assessment and decision-making.

3 Computing Fiber Confidence

The visualization framework we propose requires a set of fibers (represented by streamlines) where each fiber is associated with a confidence value. Figure 3 gives an overview of the processing pipeline to compute fiber confidence from a set of diffusion-weighted images. Confidence values are stored in a table, normalized and sorted from high (1.0) to low (0.0) confidence. How we use the table to visualize fiber confidence intervals is described in the next section. As described in the related work section, there are different ways to obtain a confidence value for each streamline. Each approach has its advantages and disadvantages. Our framework is general and does not depend on a specific approach for computing fiber confidence. To demonstrate this flexibility we have chosen two methods, (1) the Wild Bootstrap algorithm [13] combined with a fiber distance measure and (2) the ConTrack fiber tracking algorithm [28] which assigns a connection probability to each pathway. Both methods rely on a random process, either in the fiber tracking itself (ConTrack) or in the generation of the underlying tensor field from which the fibers are reconstructed (Wild Bootstrap). For this reason, we consider both algorithms to be probabilistic in nature. The output of both algorithms is a set of fiber pathways. Often these pathways are converted to a probability volume by counting the number of times each voxel is intersected by a pathway. This approach however is dependent on the chosen resolution of the data volume. We prefer to work with the streamlines directly because it preserves the visual continuity of the fiber tract of interest. This is especially important for the neurosurgeon who expects a certain correspondence between the visualization of a fiber tract and his or her knowledge of white matter anatomy.

3.1 Wild Bootstrap and Fiber Distance

The Wild Bootstrap method has been described by different authors [33, 16] but was first combined with fiber tracking by Jones et al. [13]. It attempts to simulate repeated diffusion-weighted imaging (DWI) scans by generating random variations of the tensor data from a single DWI dataset. For this, the diffusion tensor model is fit to the data in each voxel. For more than 6 diffusion directions, this results in a set of model errors or residuals, one for each measurement direction. The residuals are randomly flipped (by multiplying them with either -1 or 1), after which the tensor model is fit again. By repeating this procedure many times a large number of tensor volumes is generated. For details on the exact procedure to compute the Wild Bootstrap we refer the reader to Jones et
al. [13]. In each generated tensor volume we perform deterministic streamline tracing [30] starting from a pre-selected seed region. The same seed region is used for all tensor volumes and tracing is terminated when the anisotropy drops below a pre-defined anisotropy threshold or the angle of two consecutive main eigenvectors exceeds a given angular threshold. For each seed point this results in a set of fibers, one for each tensor volume. Each fiber describes one possible result of the streamline tracing algorithm. Because we keep the seed region fixed, we consider fiber variations for each seed point separately. To be able to assign a confidence value to each fiber we assume that the distribution of fibers in each seed point is uni-modal. In that case, we can choose a distance measure between pairs of fibers and use it to define a ‘mean’ fiber which is most central, given the distance measure, in the set of fibers at each seed point. We subsequently use each fiber’s distance to the mean fiber as a representation of confidence. A confidence interval can then be expressed as: (1) a percentage (e.g. 50%) of the most confident fibers or (2) all fibers with a confidence higher than a given threshold confidence (e.g. 0.95). Which option is most appropriate depends on the user task and the distribution of confidence values in the total set of fibers. Our framework supports both options. Fiber distance measures have been extensively described in earlier work on fiber clustering [19] and comparison of fiber tracking algorithms [12].

The choice of distance measure is entirely dependent on the application within which it is used. Each measure has its advantages and disadvantages. For this reason, our framework is not dependent on the particular choice of distance measure in any way. To demonstrate its flexibility in this respect, we implemented two widely used measures: (1) the Minimum End-Point Distance and (2) the Mean of Closest-Point Distances [19]. The minimum end-point distance \( d_E \) between two fibers \( F_i \) and \( F_j \) is defined as follows,

\[
d_E(F_i,F_j) = \min(d_1,d_2)
\]

where,

\[
d_1 = |F_{i,1} - F_{j,1}| + |F_{i,end} - F_{j,end}|
\]

\[
d_2 = |F_{i,1} - F_{j,end}| + |F_{i,end} - F_{j,1}|
\]

where \( F_{i,1} \) and \( F_{i,end} \) refer to the first and last points on fiber \( F_i \). The mean of closest-point distances \( d_M \) is defined as follows,

\[
d_M(F_i,F_j) = \frac{\min(d_m(F_i,F_j),d_m(F_j,F_i))}{2}
\]

where, given that \( p_i \) and \( p_j \) are the points on each fiber,

\[
d_m(F_i,F_j) = \min_{p_i \in F_i,p_j \in F_j}||p_i - p_j||
\]

Given the assumptions described previously, we define, for each seed point, the mean fiber to be the fiber with the smallest sum of distances to all other fibers at that seed point. This definition will result in an existing fiber and, given the chosen distance measure, this will be the most central fiber of the set. Our approach is similar to O’Donnell et al. [21] in that they also select an existing fiber instead of trying to compute an exact center line. The latter approach has been attempted in previous work [13,8,6], however, this can lead to ill-defined results when streamlines strongly diverge or even turn back on themselves. Merhrhof et al. [18] circumvent this problem by reverting to a smoothed, voxel-based representation of the fibers which allows them to generate a tight fitting surface mesh around the tracts. However, their approach does not take uncertainty into account.

Our definition of fiber distance assumes a uni-modal distribution of fibers originating from a seed point. Although this may be a valid assumption for some fiber tracts, it may be incorrect for others. In that case a different approach to compute fiber confidence must be chosen. Our framework allows easy integration of alternative methods. One such method will be described in the next section.
3.2 ConTrack

ConTrack is a probabilistic fiber tracking algorithm proposed by Sherbondy et al. [28]. It is specifically designed to take into account knowledge that a fiber connection exists between two regions. For example, in a normally sighted individual we know that there exists a functional connection between the lateral geniculate nucleus (LGN) and the primary visual cortex (V1 and V2). However, most probabilistic FT algorithms would estimate the connection probability between LGN and V1/V2 to be substantially less than one. The reason for this is that the probability of a connection between A and B, that is \( P(A \rightarrow B) \), depends on the probability of connections between A and all other points. A consequence of this dependence is that in most cases connections are not symmetric, that is \( P(A \rightarrow B) \neq P(B \rightarrow A) \). In contrast, ConTrack ensures both independence and symmetry. The algorithm outputs a large set of streamlines, each associated with a pathway score representing the probability of connection between the source and target regions. This fits the requirements of our visualization framework very well because the pathway score can be used directly as a measure of confidence. We do not claim that ConTrack is superior to other probabilistic methods such as proposed by Behrens et al. [2] or Friman et al. [10]. These methods also sample many pathways but do not always explicitly store the pathways as streamlines. Instead, the pathways are used to compute a probability density volume where each voxel value represents the number of pathways intersecting it.

4 Visualizing Fiber Confidence

In the current work we have chosen to describe the uncertainty in fiber pathways in terms of discrete confidence intervals and represent the fibers in each interval by means of silhouettes and outlines. Confidence intervals provide a high-level overview of the distribution of fibers in the set, similar to confidence intervals used in descriptive statistics. Detail is lost by aggregating multiple fibers in a single interval but if multiple confidence intervals are shown, which our framework allows, the user can obtain a good impression of which parts of the fiber tract are most reliable or reproducible and which parts are highly variable. Showing the fibers by means of streamlines, even if each confidence interval would be colored differently, will result in a large amount of visual clutter. Also, users are often interested in individual fibers but rather in fiber bundles. A silhouette and outline representation fulfills these requirements by reducing visual clutter while preserving the overall shape of the fiber bundle. As explained before, we define a silhouette as the interior area of an object’s shape when projected to the viewing plane. An outline is defined as the border of a silhouette. Otten et al. [22] presented an approach to render silhouettes and outlines for sets of clustered fibers using a GPU-accelerated algorithm. We extend this approach in the following ways:

- Extending the set of visual styles (e.g. transparency, color schemes, blurring, etc.)
- Making each visual style adaptive to interval confidence
- Combining an arbitrary number of confidence intervals in multiple render passes
- Adding anatomical context by means of 3D orthogonal slices, volume rendering and surface models (e.g. tumor, brain ventricles and cortical surface) and other types of streamline visualizations

Figure 4 gives an overview of the different stages in our visualization pipeline.

4.1 Generating Silhouettes and Outlines

To create a silhouette and outline representation we render the streamlines directly into an off-screen buffer image. Next, we apply a dilation operator to each non-empty pixel of the streamline image. The dilation operator is based on a circular structuring element with a user-defined silhouette radius \( R_s \). This widens the area covered by the streamlines and closes holes between neighboring streamlines. A second, larger structuring element with a user-defined outline radius \( R_o \) is used to add an outline of a given thickness to the silhouette. The general mechanism is illustrated in Figure 5.

4.2 Rendering Confidence Intervals

To visualize the illustrative confidence intervals we transfer the streamlines and associated confidence table to our rendering pipeline (see Figure 4). We then specify a confidence interval \( \Delta = [a, b] \), where \( [a, b] \subseteq [0, 1] \), to control which streamlines are rendered to screen as silhouettes and outlines. If the confidence interval is \( [0, 1] \) we render all streamlines at once and the resulting silhouette represents all possible variations of streamlines. If we have \( N \) intervals, then each interval \( \Delta_i \) where \( 0 \leq i < N \), can be rendered with a different visual style. Figure 4 illustrates the different steps...
in our visualization pipeline. For each interval $\Delta_i$, these steps are executed as follows:

**Step 1:** We start with the interval of highest confidence, that is $\Delta_{N-1}$ and render the corresponding streamlines to a color buffer $C_{STR}$ and depth buffer $D_{STR}$ using OpenGL framebuffer objects. Depth testing is set to $\text{GL}\_\text{LESS}$.

**Step 2:** In this step, stencil testing is enabled (function: $(\text{GL}\_\text{EQUAL}, 0, 1)$, operation: $\text{GL}\_\text{INCR}$) to ensure that subsequent intervals (with lower confidence) do not overwrite the current one (with higher confidence). We could have solved this by setting the depth test to $\text{GL}\_\text{ALWAYS}$ and rendering the low confidence intervals first. However, reducing the opacity of a given silhouette would then result in unwanted color mixing of the different intervals. If the opacity is reduced, we wish to see the anatomy lying behind it and not other confidence intervals. The buffers $C_{STR}$ and $D_{STR}$ from the previous step are passed to a GPU shader program that creates the silhouette and outline representation described previously. The output result of the shader program is written to the color buffer $C_{SILH}$ and depth buffer $D_{SILH}$. In this step the visual style parameters are applied (see also sub-section 4.3).

**Step 3 (Optional):** This step is only performed if silhouette blurring is enabled. This allows the user to show intervals of lower confidence with increased blurring. The stencil test is now performed at this step instead of Step 2. The buffers $C_{SILH}$ and $D_{SILH}$ from Step 2 are passed to a second GPU shader program that applies Gaussian blurring to the color buffer. The shader program takes a blurring radius and brightness offset as parameters (see also sub-section 4.3). The brightness offset allows the user to adjust brightness in case it is too low after the blurring operation.

Fig. 4 Pipeline for visualizing our illustrative confidence intervals

Fig. 6 Left: Confidence Histogram Widget with confidence values increase from left to right. White arrows indicate possible manipulation direction. Selected active property is ‘Opacity’. Right: Corresponding confidence intervals of optic radiation fibers looping around ventricles.

4.3 Confidence Histogram Widget

Depending on the distribution of confidence values within the fiber set, the user may wish to choose the confidence intervals differently. To make this possible we introduce the **Confidence Histogram Widget** that shows a histogram of confidence values and allows the user to define and manipulate confidence intervals graphically. Additionally, it is possible to identify deviations from a uni-modal distribution in the fibers. In Section 3 we explained how we use distance to compute fiber confidence. This approach assumes that the distribution of fibers originating from a seed point is unimodal. If this is not the case and the fibers split up into two, equally-sized bundles, the mean fiber may be assigned randomly to one or the other bundle. All fibers *not* in the same bundle as the mean fiber will have a larger distance and are therefore assigned a lower confidence. In that case, the histogram is likely to show two peaks. Figure 6 on the left illustrates an example histogram computed from output scores of the ConTrack algorithm. The confidence intervals are displayed as a set of semi-transparent rectangles overlaid on
Uncertainty in DTI fiber tracking algorithms is an important issue in neurosurgical applications. However, some fiber tracts can be more reliably reconstructed than others, even with deterministic approaches. Also, not all regions of the brain are affected by the surgical approach. In such regions, showing uncertainty may be unnecessary or even confusing. For this reason, we provide the Uncertainty Lens as a Focus+Context approach to show uncertainty only within a user-defined region-of-interest (ROI). Outside the lens standard streamline visualization can be used. For example, the surgeon can place the uncertainty lens over the tumor and its immediate surroundings and see the fiber variations due to the uncertainty within a relevant context. If necessary, the user can interactively move and resize the uncertainty lens. Figure 7(c) gives an example visualization.

5 Results and Discussion

Figure 7 illustrates uncertainty visualizations created with our framework. Figure 7(a) depicts a transversal view of the optic radiation (running from the thalamus to the visual cortex) with brain ventricles rendered using DVR. Figure 7(b) provides a sagittal view of the corticospinal tract surrounding a tumor. Figure 7(c) illustrates our Uncertainty Lens together with a standard streamline visualization, tumor and semi-transparent cortical surface. Figures 7(d-i) illustrate a close-up of the optic radiation with most of the visual styles we support (warm-to-cool outlines, warm-to-cool surfaces, decreasing opacity, light-to-dark, increasing dilation, increasing blur).

5.1 Datasets and Performance

These visualizations were created with datasets from three suppliers. Kempenhaeghe Epilepsy Center: DTI (112 × 112 × 60, 2 × 2 × 2mm$^3$, b-value 1000, 30 grad. dirs.). St. Elisabeth hospital: DTI (128 × 128 × 60, 1.75 × 1.75 × 2 mm$^3$, b-value 800, 30 grad. dirs.), T1 MRI (288 × 288 × 175, 1 × 1 × 1mm$^3$). VisContest 2010: DTI (128 × 128 × 72, 1.8 × 1.8 × 1.8mm$^3$, b-value 1000, 30 grad. dirs.), T1 MRI (512 × 512 × 176, 0.488 × 0.488 × 1mm$^3$). We measured the framerate performance of our rendering pipeline using a GeForce 8800 GTX graphics card from NVIDIA and a screensize of 720 × 760 pixels: Figure 7(a) = 6.6 fps, Figure 7(b) = 10.2 fps and Figure 7(c) = 17.5 fps.

5.2 User Evaluation

To evaluate our visualization approach we conducted an informal user study with three neurosurgeons. First, we wanted to know their general opinion about the need and benefits of uncertainty visualization for their work. Second, we wanted
to evaluate the specific rendering options that our visualization framework provides. We presented the neurosurgeons with a set of screen-shots, videos and live demonstrations. Furthermore, we asked them to fill out a questionnaire with respect to the visual material. Three clinical scenarios were presented: (1) anterior temporal lobe resection (ATLR) for the treatment of focal epilepsy, (2) resection of a low-grade glioma close to motor cortex and (3) resection of an intracerebral metastasis with surrounding edema (fluid), also close to motor cortex. For each scenario we showed a standard fiber visualization (stream tubes) and our illustrative confidence intervals. We used three intervals each containing 10%, 25% and 100% of the most confident fibers. Table 2 shows the general opinion of the neurosurgeons on the questions listed below. Table 3 shows the specific ratings for each question. Ratings are given on a [1–5] Likert scale where 1 means either ‘not useful at all’ or ‘low’ and 5 means ‘highly useful’ or ‘high’, depending on the context of the question.

Q1.1 What is the risk of a visual field deficit in anterior temporal lobe resection?
Q1.2 How important is uncertainty visualization for anterior temporal lobe resections?
Q1.3 What is the risk of resecting this particular low-grade glioma?
Q1.4 How important is uncertainty visualization for low-grade glioma resection?
Q1.5 What is the risk of resecting this particular intracerebral metastasis?
Q1.6 To what extent would you discuss the uncertainty with the patient?

Table 2 General opinion about DTI uncertainty visualization. Ranking of visual styles rating is done on a [1–5] Likert scale.

|        | User A | User B | User C |
|--------|--------|--------|--------|
| Q1.1   | 2      | 2      | 2      |
| Q1.2   | 4      | 4      | 3      |
| Q1.3   | 4      | 1      | 1      |
| Q1.4   | 4      | 4      | 5      |
| Q1.5   | 2      | 2      | 2      |
| Q1.6   | 5      | 4      | 4      |

Q2.1 To what extent does the standard fiber visualization give you more confidence in assessing risk of anterior temporal lobe resections?
Q2.2 To what extent does the standard visualization give more confidence in assessing risk of resecting the given low-grade glioma?
Q2.3 To what extent does the standard visualization give more confidence in assessing risk of resecting the given intracerebral metastasis?
Q2.4 Which two visual styles do you prefer?
Q2.5 Does our representation of uncertainty give you more
5.3 General Discussion

As noted in the user evaluation, the neurosurgeons were skeptical about the fiber tracking results for the intra-cerebral metastasis. The edematous fluid surrounding the tumor causes the tracking algorithm to prematurely terminate thereby resulting in false negatives. Our particular implementation of the Wild Bootstrap method will also suffer from such false negatives (failing to show something that is actually there) because it uses the same fiber tracking algorithm with the same termination criteria. If diffusion in a voxel, because of edematous fluid, is almost isotropic, the tensor model will fit the data well and have very small residuals. Random perturbation of the residuals will not result in large shape variations, so on average the tensors will remain isotropic. Because the Wild Bootstrap method uses the same thresholds as standard streamline tracing, none of the generated streamlines will get past these isotropic tensors. With crossing fibers however, the tensor can become disk-like while the underlying diffusion profile might actually be cross-shaped. In this case, the fractional anisotropy of the tensor remains high which may cause standard algorithms to continue tracking even though the model fit is bad (large residuals). The Wild Bootstrap will expose this problem by showing a large amount of variation in fiber pathways originating from the crossing fiber region. This indicates that the standard tracking algorithm may be generating random results. In order to deal with false negatives, alternative methods for probabilistic fiber tracking can be used. These are often based on Monte Carlo sampling of pathways and less dependent on stopping criteria [10,2]. Even in regions of low anisotropy these algorithms can detect the presence of fiber pathways.

In this paper we have applied the Wild Bootstrap approach using the 2nd-order tensor model and deterministic streamline tracing. However, these choices are not specific to the Wild Bootstrap. It is a general technique that can be applied to any model that tries to fit data and results in non-zero residuals. Even with higher-order models that are able to detect crossing fibers, you can still use the Wild Bootstrap to investigate the sensitivity of these models to noise in the data. Our choice to use streamline tracing is also not the only choice. Other fiber tracking algorithms, such as tensor deflection (TEND) [15], can be used in combination with the Wild Bootstrap as well.

With respect to the computation of confidence based on fiber distance we wish to point out that the choice of distance measure may have a large effect on the confidence intervals. As Jiao et al. [12] point out distance measures that are based on averages, such as the mean of closest-point distances, may over- or under-estimate the true distance due to streamline discretization problems or complex

| Q2.1 | 4 | 2 | 3 |
| Q2.2 | 3 | 2 | 2 |
| Q2.3 | 1 | 1 | 1 |
| Q2.4 | Opacity | Outline | Opacity |
| Lightness | - | Lightness |
| Q2.5 | 4 | 4 | 4 |
| Q2.6 | 4 | 4 | 3 |
| Q2.7 | 4 | 4 | 4 |
streamline configurations. The closest end-points distance may surely over-simplify the situation. However, for certain cases it may provide relevant information. For example, if apriori knowledge is available that fibers start and end in the same anatomical regions, they could still be considered ‘close’ even though they may take wildly different routes to get to their destination. In general we regard distance measures to be application-specific without any particular measure being the best one for all situations.

The user evaluation, although not conclusive, has provided us with useful information. Whether DTI fiber tracking, and therefore DTI uncertainty, is relevant for neurosurgical planning depends on many factors, such as tumor type. High-grade gliomas have a bad prognosis and are commonly associated with functional deficits in the patient. If there is any risk of additional damage due to resection, tumor tissue is simply left in place. Low-grade gliomas, on the other hand, have a relatively good prognosis. The tumor grows slowly allowing brain functions to reorganize if they get invaded (brain plasticity). In these cases neurosurgeons are willing to use all the information available in order to maximize resection while at the same time minimizing damage to healthy tissue. DTI fiber tracking and uncertainty visualization are especially relevant here. For anterior temporal lobe resections DTI uncertainty information is important because part of the optic radiation, Meyer’s Loop, is often damaged during the procedure, resulting in a partial loss of vision. The extent of Meyer’s Loop varies significantly between patients (between 3 and 6 cm’s from the temporal pole). This makes it difficult to predict in advance whether visual defects will result or not. Meyer’s Loop however is a difficult fiber tract to reconstruct due to strong curvature and proximity to other fiber tracts (resulting in partial volume effects at the tract boundaries). Uncertainty visualization can provide helpful information in this case.

6 Conclusions and Future Work

In this paper, we have presented a framework for visualizing uncertainties in DTI fiber tracking due to noise and modelling errors. Our contribution consists of a processing pipeline for computing fiber confidence and an algorithm for visualizing fiber confidence intervals based on illustrative silhouettes and outlines. We also provide an interactive histogram widget to view the distribution of confidence values within a fiber set and adjust the visual parameters of each interval. Our uncertainty lens allows a Focus+Context view of both illustrative silhouettes and tumor (focus) and standard fiber visualization techniques with anatomical information (context). Finally, we have evaluated the added value of our uncertainty visualization for surgical risk assessment with three neurosurgeons. We concluded that our illustrative confidence intervals are a potentially useful approach to display uncertainty. Furthermore, we confirmed that uncertainty visualization is of great value for surgical risk assessment and may play an important role in patient counselling.

For future work, we would like to experiment with more visual styles and test these in a larger group of users. In this paper we have focused our user evaluation on neurosurgeons who are actually involved in diagnosis and surgical risk assessment. We would like continue and expand our collaboration with these medical experts to find ways to apply DTI uncertainty visualization in a clinical context for specific surgical procedures, such as tumor resection or epilepsy surgery. This requires dedicated visualization and specialized interaction paradigms that are fast, intuitive and easy to use without requiring a detailed understanding of the underlying algorithms.

Acknowledgements We would like to thank the following people for their valuable input, feedback and datasets: Geert-Jan Rutten MD PhD (St. Elisabeth Hospital Tilburg, Netherlands), Pieter Kubben MD and Olaf Schijns MD (University Hospital Maastricht, Netherlands), Albert Colon MD and Pauly Osehenblok PhD (Kempenhaeghe Epilepsy Center, Netherlands), and finally the organizers of the VisContest 2010.

References

1. Basser, P., Pajevic, S., Pierpaoli, C., Duda, J., Aldroubi, A.: In Vivo Fiber Tractography using DT-MRI Data. Magnetic Resonance in Medicine 44, 625–632 (2000)
2. Behrens, T., Woolrich, M., Jenkinson, M., Johansen-Berg, H., Nunes, R., Clare, S., Matthews, P., Brady, J., Smith, S.: Characterization and Propagation of Uncertainty in Diffusion-Weighted MR Imaging. Magnetic Resonance Medicine 50, 1077–1088 (2003)
3. Berman, J., Chung, S.W., Mukherjee, P., Hess, C., Han, E., Henry, R.: Probabilistic Streamline Q-Ball Tractography Using the Residual Bootstrap. NeuroImage 39, 215–222 (2008)
4. Born, S., Jainek, W., Hlawitschka, M., Trantakis, C., Meixensberger, J., Bartz, D.: Multimodal Visualization of DTI and fMRI Data Using Illustrative Methods. In: Bildverarbeitung fuer die Medizin, pp. 6–10 (2009)
5. Brechelsen, R., Vilanova, A., Platel, B., ter Haar Romenij, B.: Parameter Sensitivity Visualization for DTI Fiber Tracking. IEEE Transactions on Visualization and Computer Graphics 15, 1441–1448 (2009)
6. Chen, W., Zhang, S., Correia, S., Ebert, D.: Abstractive Representation and Exploration of Hierarchically Clustered Diffusion Tensor Fiber Tracts. Computer Graphics Forum 27, 1071–1078 (2008)
7. Delmarcelle, T., Hesselink, L.: Visualizing Second-Order Tensor Fields with Hyperstreamlines. IEEE Computer Graphics and Applications 13, 25–33 (2003)
8. Enders, F., Saubers, N., Merhof, D., Hasteit, P., NimsKY, C., Stamminger, M.: Visualization of White Matter Tracts with Wrapped Streamlines. In: Proceedings of IEEE Visualization ’05, pp. 51–58 (2005)
9. Everts, M., Bekker, H., Roerdink, J., Isenberg, T.: Depth-Dependent Halos: Illustrative Rendering of Dense Line Data. IEEE Transactions on Visualization and Computer Graphics 15, 1299–1306 (2009)
10. Friman, O., Farneback, G., Westin, C.F.: A Bayesian Approach for Stochastic White Matter Tractography. IEEE Transactions on Medical Imaging 25, 965–978 (2006)
11. Jainek, W., Born, S., Bartz, D., Strasser, W., Fischer, J.: Illustrative Hybrid Visualization and Exploration of Anatomical and Functional Brain Data. Computer Graphics Forum 27, 855–862 (2008)
12. Jiao, F., Philips, J., Stinstra, J., Krüger, J., Varma, R., Hsu, E., Kornberg, J., Johhson, C.: Metrics for Uncertainty Analysis and Visualization of Diffusion Tensor Images. In: Proceedings of Medical Imaging and Augmented Reality, pp. 179–190 (2010)
13. Jones, D.: Tractography Gone Wild - Probabilistic Fibre Tracking Using the Wild Bootstrap with Diffusion Tensor MRI. IEEE Transactions on Medical Imaging 27, 1268–1274 (2008)
14. Klein, J., Hahn, H., Rexilious, J., Erhard, P., Althaus, M., Leibfritz, H.O.: Efficient Visualization of Fiber Tracking Uncertainty Based on Complex Gaussian Noise. In: Proceedings of International Society for Magnetic Resonance Medicine (ISMRM '06) (2006)
15. Lazar, M., Weinstein, D., Tsuruda, J., Hasan, K., Arfanakis, K., Meyerand, M., Badie, B., Rowley, H., Haughton, V., Field, A., Alexander, A.: White Matter Tractography Using Diffusion Tensor Deflection. Human Brain Mapping 30, 306–321 (2003)
16. Liu, R.: Bootstrap Procedures Under Some Non-IDD Models. The Annals of Statistics 16, 1696–1708 (1988)
17. Lodha, S., Pang, A., Sheehan, R., Wittenbrink, C.: UFLOW: Visualizing Uncertainty in Fluid Flow. In: Proceedings of IEEE Visualization '96, pp. 249–254 (1996)
18. Merhof, D., Meister, M., BingöI, E., Nimksy, C., Greiner, G.: Isosurface-Based Generation of Hulls Encircling Neuronal Pathways. Stereotactic and Functional Neurosurgery 87, 50–60 (2009)
19. Moberts, B., Vilanova, A., Wijk, J.v.: Evaluation of Fiber Clustering Methods for Diffusion Tensor Imaging. In: Proceedings of IEEE Visualization ’05, pp. 65–72 (2005)
20. Mori, S., Crain, B., Chacko, V., Zijl, P.v.: Toward a Quantitative Assessment of Diffusion Anisotropy. Magnetic Resonance in Medicine 36, 893–906 (1996)
21. O’Donnell, L., Westin, C.F., Golby, A.: Tract-Based Morphometry for White Matter Group Analysis. NeuroImage 45, 832–844 (2009)
22. Otten, R., Vilanova, A., Wetering, H.v.d.: Illustrative White Matter Fiber Bundles. Computer Graphics Forum 29, 1013–1022 (2010)
23. Pajevic, S., Basser, P.: Parametric and Non-Parametric Statistical Analysis of DT-MRI Data. Journal of Magnetic Resonance 161, 1–14 (2003)
24. Pang, A.: Visualizing Uncertainty in Geo-spatial Data. In: Proceedings of Workshop on the Intersections Between Geospatial Information and Information Technology ’01, pp. 1–14 (2001)
25. Pang, A., Wittenbrink, C., Lodha, S.: Approaches to Uncertainty Visualization. The Visual Computer 13, 370–390 (1997)
26. Pierpaoli, C., Basser, P.: Toward a Quantitative Assessment of Diffusion Anisotropy. Magnetic Resonance in Medicine 36, 893–906 (1996)
27. Schultz, T., Sauber, N., Arwander, A., Theisel, H., Seidel, H.P.: Virtual Klingler Dissection: Putting Fibers into Context. Computer Graphics Forum 27, 1063–1070 (2008)
28. Sherbondy, A., Dougherty, R., Ben-Shachar, M., Napel, S., Wandell, B.: ConTrack: Finding the Most Likely Pathways Between Brain Regions Using Diffusion Tractography. Journal of Vision 8, 1–16 (2008)
29. Svetachov, P., Everts, M., Isenberg, T.: DTI in Context: Illustrating Brain Fiber Tracts in Situ. Computer Graphics Forum 29, 1024–1032 (2010)
30. Vilanova, A., Berenschot, G., Pul, C.v.: DTI Visualization with Streamsurfaces and Evenly-Spaced Volume Seeding. In: Proceedings of IEEE TVCG Symposium on Visualization ’04, pp. 173–182 (2004)
31. Vila Nova, A., Zhang, S., Kindllmann, G., Laidlaw, D.: Visualization and Image Processing of Tensor Fields, chap. An Introduction to Visualization of Diffusion Tensor Imaging and Its Applications, pp. 121–153. Mathematics and Visualization. Springer Verlag (2004)
32. Weiler, F., Hahn, H., Koehn, A., Friman, O., Klein, J., Peitgen, H.O.: Dealing with Inaccuracies in Multimodal Neurosurgical Planning - A Preliminary Concept. In: Proceedings of the 22nd Internal Congress and Exhibition of Computer Assisted Radiology and Surgery (CARS) (2008)
33. Whitcher, B., Tuch, D., Wisco, J., Sorensen, A., Wang, L.: Using the Wild Bootstrap to Quantify Uncertainty in Diffusion Tensor Imaging. Human Brain Mapping 29, 346–362 (2008)
34. Wittenbrink, C., Pang, A., Lodha, S.: Glyphs for Visualizing Uncertainty in Vector Fields. IEEE Transactions on Visualization and Computer Graphics 2, 266–279 (1996)