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Tracking the visual system—from the optic chiasm to primary visual cortex

Introduction

While epilepsy is a complex and not thoroughly understood phenomenon of brain malfunction, the key to its understanding is in some cases actually hidden in the structure of the brain. This applies, for instance, to acquired causes of epilepsy, such as serious brain trauma, strokes, tumors, and other lesions [5], as well as to observed interactions between structure and function [24]. The comprehensive and accurate description of brain anatomy and connectivity is even more important for surgical interventions, particularly those with a high risk of causing damage to the integrity of the visual system (e.g., anterior temporal lobe resections and amygdalohippocampectomy, see **Fig. 1b**). These examples demonstrate that research on and treatment of epilepsy greatly benefits from the integration of brain imaging techniques, such as diffusion magnetic resonance imaging [23]. Diffusion MRI is a noninvasive imaging technique capable of capturing microstructural tissue properties and of mapping the fiber architecture, e.g., trajectories of neural pathways, which has been demonstrated to significantly contribute to the field of epilepsy. Its capability of capturing the microstructural properties has made it possible to link the structural compromise of fibers tracts in temporal lobe epilepsy (TLE) with memory and language impairments [24] and, as demonstrated in some cases of focal cortical dysplasias, to reveal thinning of white matter fibers and reduced connectivity between subcortical gray matter and the dysplastic cortex [10]. The latter study employed also the second feature of dMRI, i.e., its capability of reconstruction of trajectories of neural pathways. This process, referred to as "tractography" or "fiber or streamline tracking" [26], uses estimates of local fiber orientations (calculated from dMRI data) to produce a tractogram, i.e., the reconstruction of the given pathways in the brain.

The inherent risk of surgical interventions for visual function is well known in standard procedures, such as anterior temporal lobectomy and amygdalohippocampectomy. Although mostly nondisabling, a significant numbers of visual field defects were reported (respectively 78% and 73%; [14]), as for temporal lobe resections, where postoperative upper quadrant visual field deficits were reported for 29 of 38 patients [36]. Between the aforementioned resective approaches, there is no significant difference concerning the risk for visual field deficit [14]. The indicated need for more tailored surgical approaches can be addressed by incorporating tractography in the routine [32, 45], which is known to predict postoperative visual field deficits based on pre- and postoperative tractography [8, 20]. Apart from resective interventions, dMRI can also be integrated for the prevention of visual function deficits in minimally invasive approaches, such as laser interstitial thermal therapy (LiTT; also known as "MRI-guided laser ablation" or "stereotactic laser-thermoablation"; [7]), e.g., for amygdalohippocampectomies in mesial temporal lobe epilepsies ([11, 31]; see Fig. 1a) or parieto-occipital lesionectomies typically performed for periventricular heterotopias ([44]; see • Fig. 1b). At least for mesial temporal lobe epilepsies, more definitive information on risk can be awaited by larger prospective studies in the near future. However, initial results from consecutive case series suggest that the surgical risk for a visual field deficit after LiTT is less than that with the aforementioned resective procedures, particularly if an initial technical learning curve is acknowledged [17].

One risk-posing structure for all temporal surgical approaches is the individual anatomy of Meyer's loop. Nilsson et al. examined the intra- and interindividual variability of Meyer's loop in seven healthy volunteers and two patients undergoing anterior temporal lobe resection by measuring the distance between the anterior edge of the loop and the temporal lobe: the distance varied considerably between 34 and 51 mm (mean 44 mm; [27]) indicating the need for an individual approach during the presurgical planning (see section "Primary visual cortex").

Tractography allows for the visualization and assessment of the connectivity between brain regions [10] and thus can also provide vital information for surgi-



Fig. 1 ▲ Relevance of visual system imaging in clinical routines. **a** Presurgical planning for LiTT on T1-weighted image in pseudo-sagittal view. Stereotactic trajectory (*red line*) and the aimed for lesion zone depicted by a brachytherapy planning tool (arbitrarily chosen: *red contour* resembles 50 Gy lsodose as desired limit of the irreversible damage zone, *green contour* resembles 100 Gy lsodose as core heat zone which needs to be monitored for peak temperatures). Neither trajectory nor lesion zone interfere with the visual tract (as defined by deterministic tractography; *white area* above trajectory), thus minimizing the risk of visual field loss. **b** Periventricular heterotopia (*pink contour*) adjacent to superimposed optic radiation (as defined by deterministic tractography; *white streak* adjacent to the *pink contour*) on T1-weighted image with gadolinium contrast. (Image courtesy of Dept. for Stereotactic Neurosurgery/Otto von Guericke University Magdeburg)

cal intervention planning and risk assessment in epilepsy. In the following, the different concepts of imaging the visual system using dMRI are introduced.

Methods

In order to generate the tractogram of the visual system (**■** Fig. 2), it is necessary to acquire data, select the analysis tools, preprocess the diffusion-weighted (DW) data, segment the structures of the visual system defining starting and ending points of reconstructed streamlines, fit the model to DW images, and perform the tractography. While each of these steps will be discussed separately, it should be noted that they should not be perceived as separate entities, but rather as interconnected and dependent.

Overview of required MRI data

The optimal MRI dataset acquired for the purpose of tracking of the visual system should comprise (a) T1-weighted (**©** Fig. 3a), and optionally a proton density (PD) map, to allow for the comprehensive delineation of brain tissue and structures, and (b) DW images necessary for the tractography (**©** Fig. 3b). In the dMRI analysis, the T1-weighted images



Fig. 2 A Tractography of the visual pathways. **a** A 3D graphic in sagittal view displaying orientation of right optic tract and optic radiation streamlines. **b** T1-weighted images including optic tract (*left*) and optic tract streamlines (*right*) in false color (color code describes streamline orientation: *green* anterior–posterior; *red* left–right; *blue* inferior–superior). Streamlines are not cropped to the shown slices. (Image courtesy of Otto von Guericke University Magdeburg)

and PD maps are used in order to identify and segment regions of interest (ROIs) and white matter mask, which enhances tractography. The DW images provide information about microstructural properties (**•** Fig. 3c) and what is critical for tractography, spatial organization of neural fibers (additionally, tractography may also provide its own contrast as shown in **•** Figure 3d). The basic underlying concept is the measurement of dephasing of excited protons along the given direction simultaneously in all of the brain voxels [37]. The contrast in a DW image depends on the properties of diffusion processes occurring in the tissue and is controlled by two parameters set for MR scanning: b-value (determines the sensitivity to diffusion; b0 is the lowest, where b-value equal to, e.g., 3000 s/mm² is considered high) and b-vector (describes the direction along which the diffusion is measured). The selection of DW acquisition parameters is a complex issue, as one needs to consider MRI scanner capabilities, the focus of study (imaging of the whole brain or a single structure), data quality requirements (signal-to-noise ratio, spatial resolution, number of b-values, and gradient directions) and external factors, e.g., scanning time (detailed in: [22, 40]). An efficient alternative solution is the adaptation of already validated and well-established protocols such as the Human Connectome Project (HCP) acquisition protocols [42]. Publicly available datasets should also be considered an option to tune the analysis procedures (see **Table 1**).

Processing

Apart from the data acquisition parameters, it is important to consider the choice of tools for data preprocessing and tractography (see **Table 2**). Diffusion-weighted data preprocessing is an important stage, as it determines the data quality and may affect the final outcome of the study. While the applicability and/or necessity of constituent preprocessing steps depends on parameters of the data and specifics of the study, the preprocessing pipelines applied in the HCP project [16] are a commonly recognized standard. **Table 3** provides a list of recommended preprocessing steps, created by extending the HCP preprocessing pipeline by more recent utilities, which result in a corrected DW dataset aligned to the T1-weighted image (Fig. 4). A recently emerging alternative to data preprocessing and analysis on local machines are online platforms, such as BrainLife [1]. BrainLife allows one to upload one's own data, or to use existing publicly available datasets (which are either stored directly on BrainLife or can be downloaded from other repositories) and perform neuroimaging analysis in the cloud using online services.

Segmentation of visual system structures

While the previous sections can be generalized for any study involving dMRI data, the selection of regions of interest (ROIs) is determined individually for each study. In the case of tractography of the visual system, it is required to delineate at least the optic chiasm, lateral geniculate nucleus (LGN), and the primary visual cortex (V1), as depicted in **Fig. 5**. The idea here is to localize the structures being connected by visual

Abstract · Zusammenfassung

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Tracking the visual system—from the optic chiasm to primary visual cortex

Abstract

Epilepsy surgery is a well-established method of treatment for pharmacoresistant focal epilepsies, but it carries an inherent risk of damaging eloquent brain structures. This holds true in particular for visual system pathways, where the damage to, for example, the optic radiation may result in postoperative visual field defects. Such risk can be minimized by the identification and localization of visual pathways using diffusion magnetic resonance imaging (dMRI). The aim of this article is to provide an overview of the step-by-step process of reconstructing the visual pathways applying dMRI analysis. This includes data acquisition, preprocessing, identification of key structures of the visual system necessary for reconstruction, as well as diffusion modeling and the ultimate reconstruction of neural pathways. As a result, the reader will become familiar both with the ideas and challenges of imaging the visual system using dMRI and their relevance for planning the intervention.

Keywords

DTI · Diffusion MRI · dMRI · Tractography · Epilepsy surgery

Traktographie des Sehsystems – vom Chiasma opticum zum primären visuellen Kortex

Zusammenfassung

Epilepsiechirurgische Interventionen sind etablierte Ansätze zur Behandlung pharmakoresistenter fokaler Epilepsien, bergen jedoch das Risiko der Schädigung eloquenter Hirnstrukturen. In Bezug auf die Sehbahn kann dies, beispielsweise bei Läsionen im Bereich der Sehstrahlung, zu postoperativen Gesichtsfelddefekten führen. Die Identifikation und Lokalisierung der Sehbahnstrukturen mittels Diffusionsbildgebung (diffusionsgewichtete Magnetresonanztomographie, dMRT) ermöglicht es, diese Risiken einzuschätzen und zu reduzieren. Der vorliegende Artikel vermittelt eine Schrittfür-Schritt-Darstellung der Prozesse, die der dMRT-basierten Sehbahnrekonstruktionen

zugrunde liegen. Dies umfasst die Datenakquise, Vorverarbeitung, Identifikation essenzieller Schlüsselstrukturen sowie die Diffusionsmodellierung und abschließend die Rekonstruktion der Sehbahn. So wird dem Leser ein Verständnis sowohl der Möglichkeiten und Herausforderungen der Bildgebung des Sehsystems mittels dMRT als auch ihrer Relevanz für die Interventionsplanung vermittelt.

Schlüsselwörter

Visuelles System · Diffusionsgewichtete Magnetresonanztomographie · dMRT · DTI · Epilepsiechirurgie

pathways and to perform the tractography in order to reconstruct these connections. It should be noted that while tractography directly between the optic chiasm and V1 is theoretically possible, in practice it is extremely challenging due to its length and complexity. Therefore, it is recommended to incorporate the LGN in the tractography.

In terms of segmentation there are two main approaches—manual, where the structures are identified and marked by hand by a trained user, or automated, which is performed by designated software. For the latter, the FreeSurfer segmentation software (**Table 2**) is widely used and functions as a semi-standard of automated segmentation. It should also be noted that recent rapid developments in deep learning (DL) methods, although not yet widely established, resulted in new emerging tools that are of great promise.

Optic nerves

If required, the currently recommended strategy would be a manual delineation based on T1-weighted images. This may change with emerging DL-based meth-



Fig. 3 ▲ T1-weighted and dMRI-derived sample images. **a** T1-weighted images. **b** Single post-processed diffusion-weighted volume acquired with b-value = 1600 s/mm². **c** Map offractional anisotropy (*FA*) describing anisotropy of local diffusion. **d** Track-density imaging (*TDI*), tractography-derived contrast obtained by mapping density of streamlines. (Image courtesy of Otto von Guericke University Magdeburg)

ods, which demonstrate good performance [25], but are not yet widely available.

Optic chiasm

A FreeSurfer software, regarded as a standard tool in automated brain segmentation, is capable of segmenting the optic chiasm, yet its results can be suboptimal (**Fig. 5b**, top). As such it is recommended to segment the optic chiasm manually in order to achieve optimal accuracy (**Fig. 5b**, bottom)—an approach widely used in optic chiasm imaging [30]. Similar to optic nerves, current DL developments allow for the segmentation of the optic chiasm, but they are still not widely tested and available—especially in the case of optic chiasm malformations.

Lateral geniculate nucleus

The segmentation of the LGN is particularly challenging, as this structure does not stand out in T1-weighted images. While it is possible to use automated segmentation tools (e.g., FreeSurfer, SPM, FSL, GIF, MALP-EM, Mipav) to this end, this approach has limitations, e.g., LGN segmentation can even vary between versions (as shown for FreeSurfer in **Fig. 5c**). Furthermore, the lack of validation undermines the study's reliability. Therefore, it is recommended to obtain PD maps during the data acquisition, as their contrast allows for an LGN identification. In the absence of this information, an alternative strategy is the tractography-based identification of the LGN [28]. We propose that this strategy would be even further improved by simultaneous tractography from both the optic chiasm and V1 with the target in the thalamus. The intersection of the streamline endpoints might serve to identify the LGN, which can be combined with the previously introduced FreeSurfer segmentation.

Primary visual cortex

For the identification of the primary visual cortex (V1), three approaches

are available: (1) the use of automated segmentation tools, (2) individualized retinotopic mapping, and (3) the estimation of retinotopic maps from anatomical priors. (1) Automated segmentation tools in principle allow for an estimation of V1; however, individual differences from the general template introduce errors in the results of the V1 ROI definition. Further, this approach does not provide an estimate of the retinotopy of V1. (2) The alternative is to obtain retinotopic maps of the visual cortex specifically for the respective individuals with dedicated functional magnetic resonance imaging (fMRI) measurements [32]. Due to the retinotopic organization of the visual cortex, visual areas can be identified via fMRI-based retinotopic mapping (reviewed in [18, 19, 43]). T2*weighted BOLD gradient-EPI sequences are acquired in fMRI scans during visual stimulation, typically with a contrast-inverting or moving checkerboard pattern section that travels through the visual field in a systematic manner to create a specific spatiotemporal response pattern on the visual cortex. Conventional phase-encoded retinotopic mapping [33] or the neuro-computationally more demanding population receptive field mapping [13] can be applied to obtain cortical maps of the eccentricity and polar angle representations of the visual field. These are typically visualized on the computationally inflated or flattened surface of the visual cortex as derived from high-resolution T1 images. Visual area boundaries can be delineated from these maps [33] as depicted in **Fig. 5d** (top row) for the primary visual cortex (V1). (3) Based on the evidence of qualitatively consistent organization of primary (V1) and extra-striate (V2 and V3) visual area topography [12], Benson et al. [4] demonstrated the ability of an anatomical template to predict the retinotopic organization of the visual cortex with high accuracy using only a participant's brain anatomy, i.e., the patterns of the gyral and sulcal curvatures. In the absence of retinotopic mapping data, a viable alternative is therefore the use of Benson's atlas, which utilizes known anatomical priors in order to estimate V1 location, as well as polar

Table 1 Sample of online, publicly available repositories offering dMRI datasets									
Source	Modality	Sample size	Link						
Human Connectome Project (HCP; [42])	dMRI, T1w	1206 controls	http://www.humanconnec tomeproject.org/						
Beijing Normal Univer- sity data	dMRI, T1w	180 controls	http://fcon_1000.projects. nitrc.org						
Image & Data Archive (IDA; 138 studies)	Various	Various	https://ida.loni.usc.edu						
Information eXtraction from Images (IXI)	dMRI, T1w, PD	500 controls	http://brain-development.org						
dMRI diffusion magnetic resonance imaging, PD proton density, T1w T1-weighted									

Table 2 Sample list of neuroimaging tools used in DW and T1w data processing								
Name	Format conversion	T1w pro- cessing	DW pro- cessing	Analysis of mi- crostructural properties	Tracto- graphy	Function and remarks		
dcm2niix	X	-	-	-	-	Well-established converter from Digital Imaging and Communications in Medicine format (DICOM; scanner output) to Neuroimaging Informatics Technology Initiative format (NIfTI; neuroimaging standard)		
FSL [21]	-	Х	Х	Х	Х	Well-established, high-quality distortion correc- tion tools		
FreeSurfer [15]	-	Х	-	-	-	Well-established standard in T1-weighted image segmentation		
mrDiffusion	Х	Х	Х	Х	Х	MATLAB-dependent, part of VISTASOFT package		
MRtrix [41]	Х	Х	Х	Х	Х	Notable software for processing, analysis and visualization of dMRI data		
Trekker [2]	-	-	-	-	Х	Uses state-of-the-art parallel transport tractogra- phy algorithm		
FiberNavigator	-	-	-	-	Х	Supports real-time tractography from the data		
DIPY	Х	Х	Х	Х	Х	Python library for analysis of dMRI data		
DW diffusion weighted, dMRI diffusion magnetic resonance imaging, PD proton density, T1w T1-weighted								

Table 3 Suggested DW preprocessing pipeline Name Software Purpose Comments Conversion from Converting the scanner's DICOM to format Diffusion image in NIfTI format is accompanied by files stordcm2niix DICOM standardly handled by neuroimaging softing information on b-values (.bvals) and gradient directions ware (NIfTI) (.bvecs) MRtrix Denoising Removal of noise Requires unmodified data, therefore must be first in the preprocessing Gibbs rings MRtrix Removal of Gibbs ringing artifacts from im-Must be run before any interpolation is performed on the age data MRtrix/FSL Estimation of motion in the images Benefits from low b-value volumes interspersing the DW Topup series (Fig. 4a) Acquisition of at least portion volumes with opposite Eddy MRtrix/FSL Correction of motion, geometry-induced and phase-encoding direction is recommended (**Fig. 4b**) eddy current distortions Gradient non-linear-FSL Correction of gradient non-linearity of diffu-Recommended if not performed automatically during data ity correction sion gradients acquisition Removes bias field in order to normalize **Bias field correction** MRtrix/ANTS MRtrix function interacting with ANTs software voxels intensities Any transformations of DW image must also be applied to Alignment MRtrix/FSL Aligns the DW and T1w and PD maps in order to ensure match between images on the gradient directions (.bvecs file) (Fig. 4c) voxel scale Upsampling MRtrix Optional step allowing data to be interpo-Optional, in some cases recommended lated to increase resolution DW diffusion weighted, dMRI diffusion magnetic resonance imaging, PD proton density, T1w T1-weighted



Fig. 4 A Exemplary distortions affecting the quality of the diffusion-weighted (*DW*) b0 images. **a** Example of motion-induced distortions. First (*left*) and last (*middle*) b0 image in the DW series. Calculation of voxel-wise difference between volumes reveals displacement introduced by motion. **b** Example of Echo-Planar Imaging (EPI) distortions. Discontinuity of the medium causes the inhomogeneity of the magnetic field, which warps the image. These distortions are particularly pronounced in the region of the optic chiasm (*white arrows*) and near the ear canals, and depending on the phase-encoding direction (*PED*) are manifested either by squeezing (*left*) or stretching (*middle*). Images acquired with opposite PEDs can be combined in order to unwarp the distortion (*right*). **c** Alignment of the T1-weighted (*T1w; middle*) and DW image (*left*) ensures the correspondence between voxels (*right*). (Image courtesy of Otto von Guericke University Magdeburg)

angle and eccentricity map (see **Fig. 5d**, middle row). This information can be used to specify ROIs for the purpose of tractography.

White matter

In order to enhance the accuracy of the tractography it is recommended to use the known anatomical priors, such as the limitation of the reconstructed pathways to white matter only. Such white matter masks can be extracted using, e.g., FSL or FreeSurfer software. This approach can be further extended by taking into account other types of tissues, such as in five-tissue-type segmentation (implemented in MRtrix), which is a critical component in their proposed anatomically constrained tractography (ACT; [34]).

DW data modeling

Subsequent to DW data preprocessing, it is necessary to fit the diffusion model to the data (**Fig. 6b**). The available modeling approaches depend on the quality and properties of DW data, such as the number of gradient directions, number of shells etc. As the in-depth discussion of available models is beyond the scope of this paper, the following section will cover a sample of two well-established models.

Diffusion tensor

Basic, yet successful and well-established, is the DT [3] model (**\Box Fig. 6b**, *right*), which's popularity caused confusion of Diffusion Tensor Imaging (DTI; application of DT model) with the general dMRI term (covering all models). The DT model represents diffusion as a 3×3 matrix with six independent terms (i.e., from at least six volumes with unique gradient directions), which can be graphically represented as an ellipsoid. This model has limitations, as it models only one diffusion direction per voxel, and as such fails to represent more complex structures, such as the optic chiasm (due to the presence of crossing fibers), Meyer's loop (due to its curvature), or posterior parts of optic radiation (due to fanning).

Constrained spherical deconvolution

Among other alternatives (such as Q-ball imaging, diffusion spectrum imaging), we would like to discuss in detail CSD [39], which treats signal in each voxel as a convolution of the response from a single fiber population and distribution of the local fiber's orientations. As such, CSD is capable of resolving multiple fiber bundles crossing a single voxel, which are described by orientation distribution functions (ODFs). The fitting of the CSD model requires more than six gradient directions (typically 30-60), which grants noise reduction and higher angular resolution at the cost of longer scanning time.

Tractography

In addition to the chosen diffusion model, the outcome tractogram depends on multiple parameters governing the tracking process (**2** Fig. 6). These options are introduced and briefly discussed here.

Probabilistic vs. deterministic algorithms

Generally, there are two classes of tracking algorithms—deterministic and probabilistic. Deterministic algorithms assume that fibers in each voxel are oriented in only one direction as determined by the given model (see **Fig. 6c**)—as such they offer robust results, but fail to grasp complex architectures. Alternative probabilistic algorithms at each step of tracking sample the final direction from the distribution of all possible directions. This approach makes it possible to uncover connections missed by the deterministic algorithms (**Fig. 6c**) and has been proven to be superior to deter-



Fig. 5 ▲ Segmentation of visual system structures. a A sagittal view of the 3D graphic displaying the optic chiasm (*orange*), *left* and *right* lateral geniculate nucleus (*LGN*; *purple*) and primary visual cortex (*dark gray*) and their location in the T1-weighted image. b Tilted axial slices displaying comparison of the optic chiasm masks obtained via FreeSurfer software (*top*) and manual segmentation (*bottom*). c Axial slices comparing LGN masks from FreeSurfer v6.0 (*purple*) or FreeSurfer v7.1 (*cyan*; *overlap in blue*). d Correspondence of the functional magnetic resonance imaging (fMRI)-derived retinotopic maps, their estimates derived from anatomy using Benson's atlas and the volumetric representation of the latter. *Left* and *right* columns depict eccentricity and polar angle maps, respectively. *Top row* displays surface representation of retinotopic maps of the right hemisphere derived from fMRI data using population Receptive Field (pRF) models. *Middle row* displays analogical representation derived directly from T1-weighted image using Benson's atlas. The *bottom row* displays Benson's prediction in a volumetric representation. It should be noted that the representation is limited to the central visual field (14° radius). (Image courtesy of Otto von Guericke University Magdeburg)

ministic tracking. Consequently, modern tracking algorithms relying on advanced models, such as CSD, employ probabilistic tracking algorithms.

Algorithm types

Even within the class of probabilistic algorithms, there exists a wide range of possible choices, which impacts the outcome tractogram. The difference between two probabilistic algorithm—iFOD2 (implemented in MRtrix) and parallel transport tractography (implemented in Trekker)—is demonstrated in **Fig. 6c.**

Seeding region

The term "seed" refers to locations chosen as starting points for reconstructed streamlines. Global seeding describes allowance to track from any brain voxel (usually white matter voxel), while ROI seeding refers to limiting seeds to defined ROIs. Global seeding allows for the reconstruction of all possible pathways, which is important in studies on the connectivity within the whole brain, but at the same time is much more computationally demanding. It also does not guarantee that the pathways of interest will indeed be reconstructed. The ROI seeding limits the tractography only to the selected structures, which is faster, but may, at the same time, limit the application of streamline post-processing options (see next section).

Target ROIs

For the recommended probabilistic algorithms, the generated streamlines will not only be limited to the "true" pathways between seeds, but will also cover a wide range of positive, but anatomically implausible, connections (**©** Fig. 6d). In order to limit the tractography outcome to only valid streamlines, it is advised to employ information about the destination of the streamlines, known as a "target." A combination of information about start (seed) and end (target) ROIs greatly improves the accuracy of tractography (**©** Fig. 6d).



Fig. 6 ▲ Exemplary diffusion models, tractography algorithms, and tracking strategies. **a** The axial slice of the T1-weighted image with highlighted LGN(*cyan*) and V1 (*red*). **b** Two exemplary models fit to the data: constrained spherical deconvolution (*CSD; left*) and diffusion tensor (*DT; right*). **c** Sample streamlines connecting LGN and V1 generated using different tracking algorithms (from *left* to *right*): second-order Integration over Fiber Orientation Distrbutions (iFOD2) based on CSD model, Parallel Transport Tractography (PTT) based on CSD model, deterministic algorithm based on DT, and probabilistic algorithm based on DT. **d** Impact of incorporating regions of interest (*ROIs*) in the iFOD2-based tractography. *From left to right*: streamlines connecting LGN and V1, streamlines connecting LGN and V1 and limited to white matter but not requested to reach V1; streamlines connecting LGN and V1 and limited to white matter; streamlines connecting LGN and V1, limited to white matter; streamlines connecting LGN and V1, limited to white matter; streamlines connecting LGN and V1, limited to white matter; streamlines connecting LGN and V1, limited to white matter; streamlines connecting LGN and V1, limited to white matter; streamlines connecting LGN and V1, limited to white matter; streamlines connecting LGN and V1, limited to white matter; streamlines connecting LGN and V1, limited to white matter; streamlines connecting LGN and V1, limited to white matter; streamlines connecting LGN and V1, limited to white matter; streamlines connecting LGN and V1, limited to white matter; streamlines connecting LGN and V1, limited to white matter; streamlines connecting LGN and V1, limited to white matter; streamlines connecting LGN and V1, limited to white matter; streamlines connecting LGN and V1, limited to white matter; streamlines connecting LGN and V1, limited to white matter and requested to pass through user-defined ROI (enforcing anatomical plausibility of streamlines). (Image courtesy of Otto von Guerick

Additional inclusion/exclusion ROIs

Still, the definition of seed and target ROIs does not necessarily unambiguously determine the reconstructed streamlines, especially in the case of complex neural connections, such as the optic radiation. Therefore, it is necessary to always visually inspect the tractography results. For uncertain results, it is recommended to introduce further inclusion or exclusion ROIs (based on anatomical knowledge) in order to impose other restrictions on generated streamlines and remove implausible connections (Fig. 6d). A good and widely used example of the incorporation of anatomical priors in tractography is using the white matter mask as a limitation for the tractography. This option is supported in all tracking software, as well as

its extensions to tissue types other than WM (such as in ACT; [34]).

Other parameters and ensemble tractography

Apart from the aforementioned points, it is also necessary to choose the parameters governing tracking, such as the number of steps, maximal angle between consecutive steps, and possible thresholds preventing tractography from entering false regions etc. A detailed discussion of this topic requires a separate study, as the selection of fixed parameters introduces tracking bias. A solution to this problem is the repetition of tractography multiple times (using different algorithms, sets of parameters, and possibly even models) and combining all these into one outcome [38]. This will, however, naturally extend the complexity of the analysis and its duration.

Streamline post-processing editing and filtering

Once the tractogram is generated, it can be subjected to further editing and adjustments. All approaches can generally be grouped into two classes: (a) Userinformed tractograms can be merged, divided with respect to the number of streamlines or their properties (such as length), split using newly defined ROIs, transformed to different templates etc. The common denominator here is the user-made decision about the action. (b) Signal-informed tractograms are automated methods that refine the selection of streamlines using initially measured signal as a reference, a validation process referred to as filtering. As an example, linear fascicle evaluation (LiFE; [29]) calculates the predicted signal based on generated streamlines and compares it with the currently measured signal. Redundant streamlines with zero contribution to the currently measured signal are subsequently discarded. Other filtering methods include, e.g., Spherical-deconvolution Informed Filtering of Tractograms (SIFT; [35]) or Convex Optimization Modeling for Microstructure Informed Tractography (COMMIT; [**9**]).

Outlook

The previous sections presented the procedure involved in the tractography of the visual system and its relevance for presurgical planning. From a more general viewpoint, dMRI with its unparalleled capability of capturing the architecture and microstructural properties is a versatile tool also in research on epilepsy, or any kind of neuroscientific research involving brain anatomy. Neurosurgery is continuously seeking to be less invasive, yet attempts in this direction are often hindered by limitations in knowledge, e.g., of the function of higher-order visual cortices. This is expected to eventually change with the scientific progress regarding structure-function relationships in the visual system and thus foster its integration with clinical applications and presurgical planning. A fine example of such developments are research initiatives aiming to design visual prosthesis by microelectrode stimulation of the V1 [6]. Here, it is anticipated that dMRI will be of assistance in deciphering retinal projection fields in V1 a priori to the application of the high-resolution stimulation grids. As such, this would be a meaningful step in the development of visual prosthesis, even though it is still a long way from creating phosphenes capable of inducing meaningful visual impressions by neuronal stimulation.

Practical conclusion

- Surgical planning and risk assessment in epilepsy benefit greatly from an individualized reconstruction of the visual pathways.
- Integrated diffusion magnetic resonance imaging (dMRI)-based tractography allows for the individualized identification of the visual pathways, including optic tracts and optic radiation.
- To acquire the information essential for successful tractography and to cope with imaging artifacts, dMRI requires careful consideration of data acquisition settings and preprocessing tools.
- Tractography requires the segmentation of seed structures, i.e., optic chiasm, lateral geniculated nucleus, primary visual cortex, and white matter masks.
- The choice of the correct dMRI data modeling framework is critical for the successful tractographybased reconstruction of the visual pathways.

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Compliance with ethical guidelines

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All procedures performed in studies involving human participants or on human tissue were in accordance with the ethical standards of the institutional and/or national research committee and with the 1975 Helsinki declaration and its later amendments or comparable ethical standards. Informed consent was obtained from all individual participants included in the study.

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