

REVIEW

A MACROSCOPIC VIEW OF MICROSTRUCTURE: USING DIFFUSION-WEIGHTED IMAGES TO INFER DAMAGE, REPAIR, AND PLASTICITY OF WHITE MATTER

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Abstract—Since its introduction in the early 1990s, diffusion-weighted magnetic resonance imaging (MRI) has played a crucial role in the non-invasive evaluation of tissue microstructure of brain parenchyma *in vivo*. Diffusion anisotropy, in particular, has been extensively used to infer histological changes due to brain maturation and pathology, as it shows a clear dependence on tissue architecture. Although the resolution used in most studies lies in the macroscopic range, the information provided originates at the microscopic level and, as such, diffusion MRI serves as a microscope that can reveal profound details of tissue with direct clinical and research applications. The interpretation of diffusion parameters of white matter rests on what is known to drive diffusion anisotropy, namely axonal membranes, density and coherence, as well as myelin sheaths. However, these factors interact to modulate anisotropy, making interpretations potentially difficult. While there are numerous publications that report diffusion changes in response to particular, histologically confirmed tissue abnormalities in animal models of disease, the microscopic correlates of altered diffusion parameters due to neurological disorders in humans have been difficult to characterize. Animal models may provide insight into the mechanisms involved, but do not necessarily provide accurate representations of the human condition, making human diffusion MRI studies with direct histological confirmation crucial for our understanding of tissue changes secondary to neurodevelopment and disease. This work provides a synopsis of tissue characteristics that give rise to highly informative, specific diffusion patterns, and also of how methodological and artifactual aspects can provide erroneous diffusion measurements that do not accurately reflect tissue and may lead to misinterpretation of results. Examples of diffusion changes due to human conditions are provided to illustrate the wealth of applications of diffusion MRI in clinical and research fields.

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Abbreviations: CHARMED, composite hindered and restricted model of diffusion; DTI, diffusion tensor imaging; FA, fractional anisotropy; FWHM, full width at half maximum; fODF, fiber orientation distribution function; MD, mean diffusivity; MRI, magnetic resonance imaging; PDF, probability density function.

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INTRODUCTION

The self-diffusion of water molecules is random, but barriers present in tissue hinder such free motion and, in effect, impart a particular signature to the displacement profile of water molecules. Far from being a drawback, this provides a mechanism with which to probe the micro-structural organization of tissue. In tissue with highly organized and coherent microstructure, water does not diffuse equally in all directions, but rather follows a preferred direction and is thus anisotropic. The brain's white matter and peripheral nerves are composed of bundles of packed axons with similar orientations and have been extensively analyzed in terms of their diffusion properties. Neural tissue is, however, not the only tissue capable of imparting anisotropy to the diffusion of water molecules, as skeletal and myocardial muscle fibers, and potentially any other tissue with a regular and directional organization, also show this phenomenon. The biological mechanisms that produce diffusion anisotropy in nervous tissue are well understood and have been reviewed before (Beaulieu, 2002). Briefly, the main contributor to diffusion anisotropy is the presence of ordered cellular membranes, such as the axolemma, with the spacing between them within such a range that a water molecule is likely to encounter them several times within a short time period. Water molecules

diffusing randomly at body temperature could potentially move as far as ten microns in 50 ms, providing ample opportunity for their interaction with axonal membranes. If the axons are spaced too far apart (or if the diffusion phenomenon is studied within a very small time window), water molecules will not have sufficient time to interact with the membranes, and diffusion anisotropy will appear to be reduced or not be evident at all. Semi-permeable barriers, such as the myelin sheaths that encapsulate certain axons, further modulate diffusion anisotropy. Thus, the diffusion profile is governed by the micro-architecture of the tissue (Pierpaoli et al., 1996), and if such architecture is coherent enough (and the diffusion profile is sampled appropriately), then the preferred direction of anisotropic diffusion will match the predominant orientation of the tissue.

Diffusion-weighted magnetic resonance imaging (MRI) allows the simultaneous assessment of water diffusion in a large number of samples (pixels) across an imaging plane (Lebihan and Breton, 1985; Merboldt et al., 1985; Taylor and Bushell, 1985; Le Bihan et al., 1986). Relating the signal in the presence of diffusion sensitization to that of an otherwise identical image, but lacking diffusion sensitization, provides a quantitative measure of the diffusion process in one dimension. If the experiment is repeated several times and the diffusion behavior is sampled in three dimensions, it is possible to reconstruct the diffusion profile on a pixel-by-pixel basis. As stated above, the preferred direction of diffusion on a given pixel (or voxel, in three dimensions) will relate to the tissue contained therein. White matter, being highly organized, will produce highly anisotropic diffusion profiles that follow the shape of major fiber bundles. Neighboring voxels belonging to the same structure will have similar preferred diffusion orientations: this is exploited by computational algorithms that enable the reconstruction of the three-dimensional trajectory of white matter bundles – a technique known as tractography (Mori and van Zijl, 2002). Neuroanatomical studies have benefited greatly from the use of tractography, as it is currently the only method available to delineate white matter pathways in the living human (Johansen-Berg and Behrens, 2006; Ciccarelli et al., 2008; Catani et al., 2012). Methodological aspects and applications of tractography are not covered here, as they warrant their own discussion.

Apart from providing the orientation of white matter fibers, the study of the diffusion of water molecules within each voxel provides a means to probe tissue microstructure in a non-invasive fashion in living tissue. It is of utmost importance, however, to acknowledge that MRI measurements of diffusion summarize an ensemble of water molecules that are simultaneously interacting with the tissue, and that a very large number of fibers may be present within the voxel. Thus, at each voxel (typically 2–3 mm in each dimension), diffusion MRI measurements provide information about the average characteristics of the tissue contained therein, bridging a spatial gap of three orders of magnitude. Fortunately, a leap of faith is not needed, as it has been

demonstrated repeatedly that diffusion measurements are capable of providing accurate representations of tissue properties (Beaulieu, 2002). What is needed, however, is a solid knowledge of the mechanisms involved, as many confounding factors hamper a clear and proper interpretation of the diffusion profile (Tournier et al., 2011; Jones et al., 2012; Le Bihan and Johansen-Berg, 2012). This article highlights particular scenarios in which histological characteristics have been inferred in humans by diffusion MRI, with a strong bias toward those studies in which direct histological confirmation was obtained, and with selected references to the animal research literature to provide context.

INFERRING MICROSTRUCTURE NON-INVASIVELY

Considering the difficulties of obtaining histological confirmation of human imaging studies, most of what is known regarding the changes of water diffusion due to white matter pathology comes from animal models of disease, which are assumed to correspond to the human condition. While this may be the case in some circumstances, great care must be exercised in order not to over-interpret diffusion parameters (Jones et al., 2012). Other important sources to support the interpretation of quantitative diffusion parameters include autopsy specimens (Schmierer et al., 2007, 2008), and prior knowledge regarding histological changes in specific neurological diseases.

The most widely used descriptor of water diffusion is the tensor model, with its imaging application known as diffusion tensor imaging (DTI) (Pierpaoli et al., 1996). In this mathematical model, water diffusion is assumed to be Gaussian and unidirectional, and it is represented as an ellipsoid shaped by the eigenvalues of the tensor. The eigenvector associated with the largest eigenvalue therefore represents the principal diffusion direction of water molecules of the tissue within a given voxel, and the remaining two eigenvectors are oriented orthogonal to it. Hence, the ellipsoid is oriented parallel to the predominant direction of white matter fibers. Anisotropy indices such as fractional anisotropy (FA), volume ratio (VR), relative anisotropy (Sands, 1995; Basser and Pierpaoli, 1996; Pierpaoli and Basser, 1996), or the linear component of the ellipsoid (CL) (Westin et al., 2002) relate longitudinal and radial diffusivities, while the mean apparent diffusion coefficient (ADC) or mean diffusivity (MD) provide information about the overall diffusivity, regardless of directionality.

In general, the diffusion tensor model is readily interpreted based on what is known to drive diffusion anisotropy (Beaulieu, 2002): longitudinal diffusivity (i.e., that associated with the largest eigenvalue, and thus, reflecting the diffusivity along the white matter bundle) is affected by axonal damage, particularly fragmentation (Song et al., 2003; Concha et al., 2006; Budde et al., 2007; Sun et al., 2008; Zhang et al., 2009; Liu et al., 2013), while radial (or perpendicular) diffusivity conveys information regarding axonal density, myelin integrity, and axonal diameter, as well as fiber coherence

(Takahashi et al., 2002; Song et al., 2003; Sun et al., 2006; Concha et al., 2010; Choe et al., 2012; Qin et al., 2012).¹ It is important to note that radial diffusivity is affected by several independent micro-architectural features, and it is impossible to discern their specific contributions by means of the tensor model alone.

Axons and myelin are not the only components within the white matter, yet they have been given a pivotal role in the biological interpretation of the diffusion tensor. Glial cells, for example, are abundant in the white matter, but are rarely taken into consideration when interpreting diffusion parameters (Ronen et al., 2013). Recently, highly organized formations of glial cells in the scar surrounding traumatic brain injury were found to be responsible for elevating diffusion anisotropy to a much larger degree than axonal or myelin contributions (Budde et al., 2011). Also, there is ample evidence that white matter contains a variable number of neurons, which show a descending density gradient from superficial to deep white matter (Larriva-Sahd et al., 2002; Suárez-Solá et al., 2009; DeFelipe et al., 2010). The contribution of these “ectopic” neurons to diffusion anisotropy of white matter is generally neglected, although it may be relevant in particular conditions, such as schizophrenia, in which their density is increased (Eastwood and Harrison, 2005).

Given its assumptions, the tensor model is only applicable where a single fiber population exists, as crossing, “kissing” or fanning fibers will result in oblate or isotropic tensors. More importantly, the determination of the vector associated with the largest eigenvalue in the presence of multiple fiber populations is an ill-posed problem, thus making the terms “axial” and “radial” diffusivities meaningless, along with their analysis (Wheeler-Kingshott and Cercignani, 2009). An added complication is that regions of crossing fibers may show a seemingly paradoxical increase of diffusion anisotropy when one of the two fiber populations undergoes degeneration (Pierpaoli et al., 2001; Douaud et al., 2011). It has been recently demonstrated (numerically and experimentally) that complex fiber architecture modulates all aspects of the diffusion tensor, and that such modulations are dependent on the geometrical properties of such crossings: Mean diffusivity decreases by as much as 5% as the angle between two fiber populations widens, and as a function of the volume fraction of the two fiber populations (Vos et al., 2012). Thus, the paradoxical increase of diffusion anisotropy following degeneration of a fiber population in a region of crossing fibers may be accompanied by an increase of MD that could be better interpreted by methodological (rather than physiological) considerations.

The described relationships of axial and radial diffusivities with tissue characteristics may be overly simplistic, yet they have served as the basis for

interpreting the tensor model in several conditions over the past two decades. In general and when properly analyzed (Jones and Cercignani, 2010), the tensor model is sufficient and highly informative in regions containing a single fiber orientation, such as the fornix or the corpus callosum. Despite the known limitations of the tensor model, its straightforward biological interpretation, as well as the fact that the required data are not very demanding (i.e., relatively few diffusion gradient directions are needed, with a b value on the order of 1000 s/mm²) (Lebel et al., 2012), has made its use widespread for the inference of tissue abnormalities due to injury or pathology in humans. On the other hand, the recognition that more than two-thirds of the voxels in a typical brain data set contain more than one fiber population (Behrens et al., 2007; Jeurissen et al., 2012) suggests that the tensor model is inappropriate for whole-brain analyses. The most straightforward way to deal with this problem is to estimate multiple tensors within each image voxel (Kreher et al., 2005; Peled et al., 2006; Ramirez-Manzanares et al., 2007), but the solution through this approach is non-unique, as different parameterizations of the underlying tensors may equally explain the observed data. Alternatives to the tensor model (discussed in Section 3) can cope with many of the shortcomings of the diffusion tensor and are rapidly gaining ground for the study of white matter microstructure.

Diffusion MRI, in general, is susceptible to imaging artifacts, such as image noise, geometric distortions due to eddy currents, motion artifacts (including pulsation), and ghosting. All these artifacts have a direct impact on the quantitative diffusion parameters and must be considered before attempting to provide biological interpretations of the data (Le Bihan et al., 2006; Jones and Cercignani, 2010). For example, an imaging voxel containing more than one tissue type will yield signal equal to the weighted average of its components. This artifact, known as partial volume, is most evident in voxels lying at the boundary between brain parenchyma and cerebrospinal fluid (CSF), which are prone to show fast and near-isotropic diffusion profiles due to their CSF signal contribution, obscuring the information from the tissue of interest (Concha et al., 2005; Metzler-Baddeley et al., 2012). This negative effect can be minimized by fitting the signal using a two-tensor model (Pierpaoli and Jones, 2004; Pasternak et al., 2009), or by nulling the CSF signal at the time of acquisition by means of an inversion pulse (Papadakis et al., 2002; Chou et al., 2005; Concha et al., 2005; Cheng et al., 2011). The smaller a brain structure, the greater the proportion of its voxels that may be affected due to partial volume (Vos et al., 2011), which highlights the importance of providing some form of control over this artifact in certain areas of the brain, such as the fornix or the pons.

Considering all the above, and regardless of the analytical method used, it must be stressed that diffusion MRI provides *indirect* information about tissue microstructure, and that several modeling assumptions, data acquisition limitations and image artifacts are likely to influence the resulting signal, and must be accounted for prior to making absolute claims about tissue integrity

¹ Too often, only MD and some sort of anisotropy index are reported, precluding the possibility to infer more details regarding tissue micro-architecture (Hasan, 2006). FA and MD are not enough to explain the shape of the tensor and are not orthogonal to each other (Ennis and Kindlmann, 2006): in general, the eigenvalues of the tensor should be reported for a more thorough interpretation of the results.

in the absence of direct histological confirmation (Jones et al., 2012).

White matter damage and repair

The evaluation of patients with acute ischemic stroke has been the hallmark application of diffusion MRI. This imaging technique has revolutionized the field, for it allows a prompt and accurate diagnosis followed by appropriate treatment. Its most important application is to identify patients with potentially salvageable parenchyma with the goal of providing treatment to those who could benefit from it, while safeguarding those patients in whom the treatment itself could produce further harm. Within minutes after an acute ischemic event, MD shows a marked decrease (Mintorovitch et al., 1991; Hjort et al., 2005), which is generally attributed to energy depletion, ionic imbalance, cellular shrinkage, release of excitatory aminoacids and a shift of water toward the intra-cellular space (cytotoxic edema) (Moseley et al., 1990; Benveniste et al., 1992; Liu et al., 2001; Sotak, 2004). It is still unclear how the increase of intracellular water results in hyper-intense signal on diffusion-weighted images (Duong et al., 2001; Ackerman and Neil, 2010; Budde and Frank, 2010; Badaut et al., 2011) as tortuosity is similar in the intra- and extracellular spaces (Duong et al., 1998), and reduced diffusivity following ischemia is not exclusive to either of the two compartments (Duong et al., 1998; Silva et al., 2002; Goodman et al., 2008). During cytotoxic edema, axons and dendrites adopt a beaded architecture (Murphy et al., 2008) that has been recently proposed to be a major contributor to the initial decrease of diffusivity following ischemia, due to the introduction of barriers (membranes) that hinder water mobility along these two neuronal structures (Budde and Frank, 2010). Further breakdown of the blood–brain barrier, damage to cell membranes, and vasogenic edema all have actions opposite to cytotoxic edema, pseudo-normalizing MD within the next 2 to 7 days (Knight et al., 1991; Duong et al., 1998; Gaudinski et al., 2008). Decreased MD is not associated with an established infarct (Liu et al., 2001), as tissue can potentially recover after reperfusion (Mintorovitch et al., 1991; Kidwell et al., 2000), although MD normalization following treatment does not guarantee tissue salvage (Li et al., 2002). Diffusion anisotropy is also time dependent with respect to stroke onset, with an initial slight increase in the hyper-acute phase (<5–7 h after onset), followed by a reduction within the next 24–36 h to values less than those found in non-affected tissue (Sorensen et al., 1999; Yang et al., 1999; Bhagat et al., 2006, 2008; Liu et al., 2007). While the mechanisms that increase diffusion anisotropy in the first few hours after stroke onset are still unclear, it has been suggested that cytotoxic edema could reduce the space between fibers and increase intra-axonal tortuosity, thus providing greater orientation-dependent barriers to diffusion (Sotak, 2002; Sen and Basser, 2005). Subsequent reduction of diffusion anisotropy is attributed to degradation of tissue microstructure in the

chronic phases of stroke (Liu et al., 2007) and can be seen in areas remote from the lesion due to secondary degeneration of the white matter fibers (Pierpaoli et al., 2001). Diffusion MRI can be used to monitor the evolution of stroke and to predict functional outcomes depending on the site of the lesion and the quantitative diffusion parameters contained therein and in remotely connected regions (Jiang et al., 2010). Although already established as an invaluable clinical tool in the evaluation of ischemic stroke, diffusion MRI is extensively used in animal models of the disorder, providing crucial information on the patho-physiology of ischemic injury, with the great advantage that this newly generated knowledge is rapidly being integrated into patient care.

Diffusion MRI is a useful tool in the field of oncology, as some brain tumors infiltrate white matter, while others merely displace it, a difference that may translate into very different treatments. For example, the infiltration of gliomas into organized tissue disrupts its architecture by virtue of disorganized cellular growth, neuronal death and axonal loss, all of which can be evaluated with DTI (Goebell et al., 2006; Stadlbauer et al., 2006). In a study of 20 patients with a histologically-confirmed diagnosis of glioma (grade II and III), the quantitative histological characteristics of 77 biopsies were correlated with pre-operative DTI indices, and the authors demonstrated a clear negative correlation between FA and both tumor cell number and degree of infiltration (Fig. 1) (Stadlbauer et al., 2006). In a different study, the same authors reported that the tract count per voxel (as derived from whole brain tractography) also correlated with quantitative histology (Stadlbauer et al., 2010), a finding that is explained in part by the dependence of the tracking algorithms on FA values. Edema and fluid accumulation in solid tumors lead to an enlarged extracellular space and, in turn, to increased MD, while highly cellular tumors show the reverse pattern (Humphries et al., 2007; Fatima et al., 2013; Koral et al., 2013). Brain abscesses generally display low values of MD, which shows a strong negative correlation with the number of viable inflammatory cells in the medium (Mishra et al., 2005), and thus diffusion MRI can be used to differentiate them from necrotic brain tumors (Reddy et al., 2006). Grading of brain tumors and their response to treatment, although debatable (Padhani et al., 2009), has also been attempted with diffusion MRI (Chenevert et al., 2000; Hamstra et al., 2005, 2008; Sijens et al., 2007). In a recent study, pre-operative super-resolution track density imaging (Calamante et al., 2010) was related to the histological characteristics of 43 image-guided biopsies from 18 patients with glioblastoma (Barajas et al., 2013). In particular, the necrotic core of the tumors corresponded to decreased track density, while the surrounding (non-Gd-enhancing) tissue showed complex characteristics, including edema, microscopic hemorrhage, necrosis and invading glioma cells within white matter that led to a paradoxical increase in track density. Further, tractography can be used to plan surgical interventions that spare displaced white matter

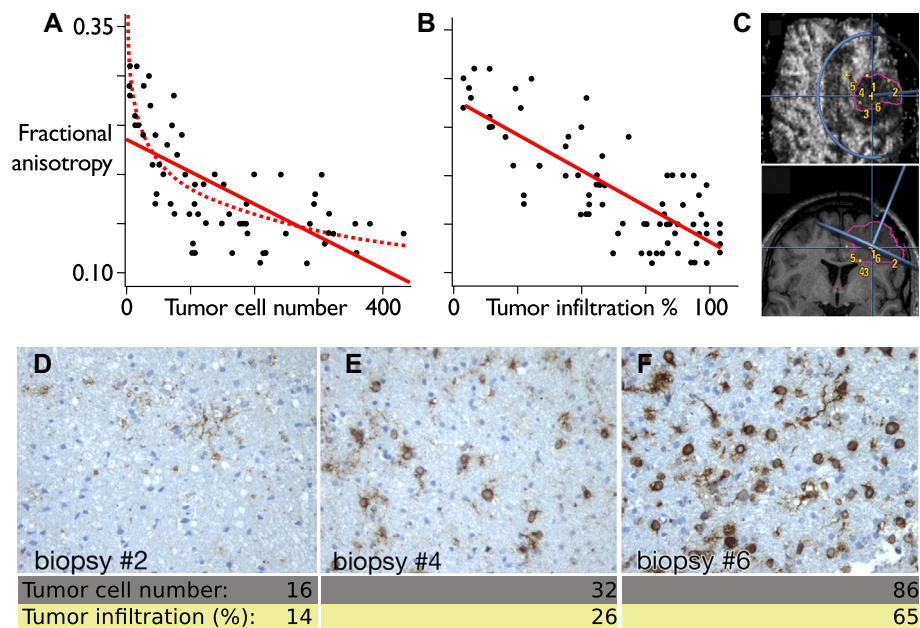


Fig. 1. Correlations between fractional anisotropy (FA) and quantitative histology of glioma biopsies. Pre-operative FA values show a negative correlation with tumor cell number (A) and tumor infiltration (B) defined as the ratio between tumor and non-tumor cells ($n = 77$ biopsies, 20 patients). (C) Biopsy sites are shown for a representative patient overlaid on an axial FA map (top) and a coronal T1-weighted image (bottom). The corresponding photomicrographs of three biopsies taken from the sites specified in C are shown in D–F. Tumoral cells are stained in brown over a hematoxylin–eosin stain, and the corresponding values (averaged over five fields) are shown below each field. (Modified from [Stadlbauer et al. \(2006\)](#), with permission from the author and the Radiological Society of North America).

fibers, thereby reducing secondary deficits brought about by their ablation [see, for example, ([Leclercq et al., 2011](#))].

While computing correlations of diffusion parameters with histology is relatively straightforward in the case of biopsy specimens, the same is not true in the case of human fiber bundles affected by disease, for there are few scenarios where pre-operative imaging and subsequent procurement of white matter tissue is feasible. For example, the fimbria-fornix of 11 patients with temporal lobe epilepsy undergoing surgery was analyzed using DTI prior to surgery, which demonstrated reduced FA and increased radial diffusivity. Quantitative electron microscopic evaluation of the surgically resected tissue revealed a positive correlation between the surface area of axonal membranes and FA and a similar positive trend between FA and axonal density ([Concha et al., 2010](#)). Although myelin has been shown to modulate diffusion anisotropy ([Gulani et al., 2001](#); [Beaulieu, 2002](#)), it did not play a role in the case of the fornix specimens, although other studies of ex-vivo human brains have proven its participation ([Schmierer et al., 2007, 2008](#); [Saksena et al., 2008](#)). The direct confirmation of microstructural changes that were predicted by pre-surgical diffusion further validates the use of DTI to detect white matter pathology, particularly in the case of fiber bundles with a single fiber population.

DTI is also a very sensitive tool for the detection of Wallerian degeneration, and the time course of the diffusion parameters reflects the underlying microstructural changes that occur during the

degeneration process ([Fig. 2](#)). During the acute phases of degeneration (i.e., the first week after the primary lesion), axons within the fiber bundles fragment and form beaded structures ([Kerschensteiner et al., 2005](#); [Zhang et al., 2009](#); [Qin et al., 2012](#)) that hinder diffusivity along the axons, which translates into reduced anisotropy secondary to a decrease of the major eigenvalue ([Thomalla et al., 2004, 2005](#); [Concha et al., 2006](#); [Liu et al., 2013](#)). Next, over a period of several months, there is further axonal membrane breakdown, progressive degradation and phagocytosis of the myelin sheaths, glial proliferation and macrophage infiltration (particularly in the vicinity of the primary lesion) ([Vaughn et al., 1970](#); [Cook and Wisniewski, 1987](#); [George and Griffin, 1994](#)), and all these events lead to a minimization or disorganization of the barriers that normally restrict radial diffusivity, thus increasing the second and third eigenvalues (reducing anisotropy) –this has been extensively demonstrated in humans ([Pierpaoli et al., 2001](#); [Thomalla et al., 2004, 2005](#); [Concha et al., 2006](#); [Yu et al., 2009](#); [Liu et al., 2013](#)) and is in line with results in animal models ([Zhang et al., 2009](#); [Qin et al., 2012](#)). The ability to stage Wallerian degeneration by characterizing the diffusion parameters of white matter enables DTI to be used as a tool for the prognosis of patients following stroke ([Yu et al., 2009](#); [Puig et al., 2010](#)) and traumatic ([Betz et al., 2012](#)) or other ([Naismith et al., 2010](#)) acute brain injuries.

Multiple sclerosis (MS) is particularly amenable to study with diffusion-weighted MRI, as degradation of myelin, one of the modulators of diffusion anisotropy, is the hallmark of the disease ([Fox, 2008](#); [Rovira and](#)

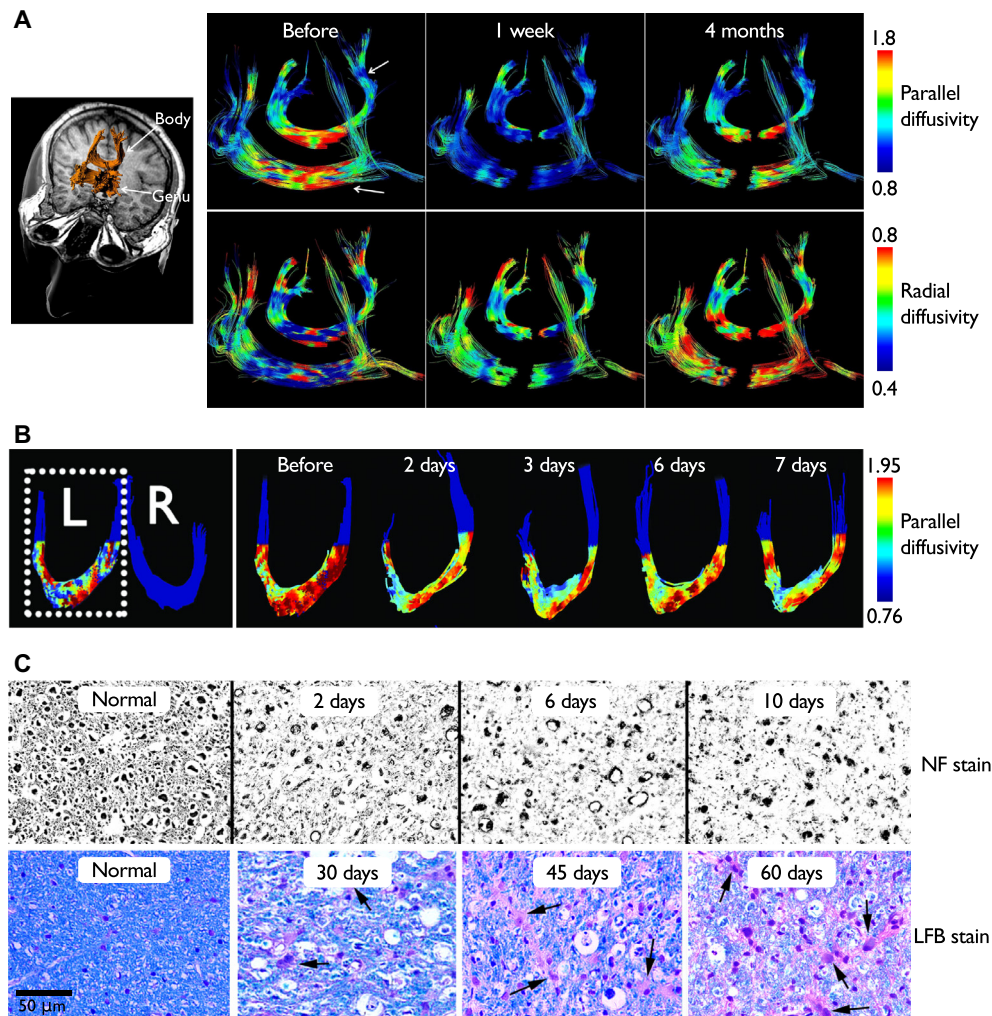


Fig. 2. Acute and chronic phases of Wallerian degeneration. (A) Diffusion parameters of the genu and body of the corpus callosum in a patient before and after surgical transection of the fibers. There is a marked decrease of parallel diffusivity at one week, with pseudo-normalization at 4 months, while perpendicular diffusivity shows a gradual increase over time. (B) The first week following transection of the fornix in a patient with temporal lobe epilepsy shows that the decrease of parallel diffusivity occurs immediately after the lesion. (C) Immunohistochemical analysis of the cortico-spinal tract in a feline model of Wallerian degeneration confirms the acute degeneration of axons, as revealed by the neurofilament protein stain (NF), and the chronic and progressive degradation of myelin evidenced by Luxol-fast blue stain (LFB). Arrows in bottom row of panel C indicate axonal debris (purple) that has been phagocytized by activated microglia. Adapted, with permission from the authors, from Concha et al. (2006) (A), Liu et al. (2013) (B) and Qin et al. (2012) (C).

León, 2008; Hygino da Cruz et al., 2011). There is, in addition, direct axonal damage (Trapp et al., 1998), which can further decrease diffusion anisotropy. Tissue inflammation is present in active lesions, and the degree to which the histological changes occur is extremely variable between lesions and among patients. Therefore, there are several imaging profiles of MS lesions with any given technique. The most severe and irreversible white matter lesions are clearly visible with conventional T1- and T2-weighted images, and show increased overall diffusivity and decreased anisotropy; these changes are more pronounced in the case of lesions that show post-Gd-enhancement or T1 hypointensity (Filippi et al., 2001). Diffusion MRI, however, has the advantage of being able to demonstrate MS lesions before conventional MRI (Rovaris et al., 2005). The diffusion abnormalities of these so-called “normal-appearing white matter” regions

become more marked as the disease progresses and neurological disability increases (Rovaris et al., 2003; Garaci et al., 2007). Furthermore, the diffusion abnormalities cannot be solely attributed to secondary Wallerian degeneration of the axons affected within visible lesions (Rovaris et al., 2005; Saindane et al., 2007). Post-mortem examination of the brains of patients with MS demonstrated that reduced diffusion anisotropy and increased MD within white matter lesions correspond to reduced myelin staining and axonal count (Mottershead et al., 2003; Schmierer et al., 2007). Given that axonal loss is a key predictor of disease progression (Petzold, 2013), diffusion MRI should be able to provide clinically relevant prognostic information regarding progression and neuroprotection, but so far it has been insufficiently accurate to become clinically viable (Barkhof et al., 2009). This may be due to the fact that axonal loss does not occur in isolation, but in

combination with several other histological changes that may modulate diffusion in contrasting ways, such as myelin degradation, edema, inflammation, astrocyte hyperplasia, activation of the microglia, increased tissue water, and alterations in neurofilament phosphorylation (Werring et al., 2000). The contribution of inflammation and increased cellularity has been recently disentangled from axonal damage using a modified multi-tensor approach known as diffusion basis spectrum imaging (Wang et al., 2011). Moreover, animal models have shown that re-myelination can be measured with DTI, particularly through radial diffusivity (Sun et al., 2006), a feat that has been replicated in human spinal cord in MS patients who recover from a spinal cord relapse (Freund et al., 2010), reaffirming the utility of diffusion MRI in longitudinal assessments of subtle white matter changes in MS patients.

Evaluating recovery through diffusion-weighted MRI has received considerable attention recently with good reason, as it provides a non-invasive way to longitudinally monitor the effect of treatment. A longitudinal study of 22 patients with traumatic brain injury (Sidaros et al., 2008) showed that DTI parameters could be used as prognostic factors regarding the patients' recovery. Shortly after injury, the patients showed reduced FA and increased radial diffusivity in several white matter regions, particularly the cerebral peduncles, which correlated with the neurological state 1 year later. Moreover, the DTI parameters of the internal capsule and centrum semiovale showed normalization in patients with good outcomes, while elevated MD and a lack of FA normalization in the cerebral peduncle and corpus callosum were associated with limited recovery. Other studies have shown that quantitative diffusion parameters of the white matter, particularly of the corticospinal tract, are indicative of motor function several months following stroke (Watanabe et al., 2001; Cho et al., 2007; Schaechter et al., 2009; Qiu et al., 2011). The tissue at the periphery of post-ischemic necrosis undergoes drastic changes that can be modulated or accentuated by neuro-restorative therapy (Jiang et al., 2010). Following stroke induced in rodents, the border zone of the ischemic lesion shows an initial reduction of FA, followed by a normalization and even elevation that is commonly attributed to axonal and myelin regeneration. The reorganization of white matter can be modulated by certain treatments such as neural progenitor cell treatment (Jiang et al., 2006), sildenafil (Ding et al., 2008) or erythropoietin (Li et al., 2009), all of which result in increased FA (Dijkhuizen et al., 2012). However, it is important to recognize that reactive gliosis shows a high degree of directionality, which can also increase diffusion anisotropy (Li et al., 2005). Indeed, this phenomenon has been clearly demonstrated to also occur in the scar that surrounds experimentally injured brain tissue, where the contribution of gliosis to the variance of diffusion anisotropy was greater than that of axons (Budde et al., 2011), meaning that not all increases of diffusion anisotropy are signs of recovery.

Plasticity of white matter

Structural plasticity of the adult brain was classically considered small, if it existed at all. However, there is at present little doubt that the adult brain is capable of undergoing molecular, structural and functional changes, many of which form the bases of memory and learning processes, or certain neurological disorders (Taubert et al., 2012). Understanding these mechanisms, as well as their time course, is pivotal for the development of restorative and neuroprotective interventions to treat neurodegenerative disorders such as Parkinson's disease, or acute brain lesions such as stroke or traumatic brain injury. Longitudinal and non-invasive investigations are necessary, particularly to study high-order processes that are mostly evident in the human brain. Neuroimaging techniques, MRI and others, can fulfill this role, but diffusion MRI in particular, with its ability to confer information of the microscopic domain, seems specially suited to probe tissue changes that are not necessarily translated into gross morphological or metabolic modifications.

Learning and training induce many types of structural and functional changes in the gray and white matter in a time-dependent manner (Johansen-Berg et al., 2012; Zatorre et al., 2012); these may include neurogenesis, synaptogenesis, dendritic arborisation, axonal and synaptic sprouting, gliogenesis, angiogenesis and myelin formation (Blumenfeld-Katzir et al., 2011; Zatorre et al., 2012). Long-term effects of cortical plasticity or training result in macroscopic morphological changes that relate to remodeling of neural processes (Lerch et al., 2011), which can be detected through MRI in humans (Maguire et al., 2000; Draganski et al., 2004; Boyke et al., 2008; Driemeyer et al., 2008; Hyde et al., 2009). The white matter of highly trained individuals, such as musicians, shows differences in diffusion parameters from non-trained subjects in key areas of the brain (Bengtsson et al., 2005; Imfeld et al., 2009; Steele et al., 2013). Diffusion MRI of human adults who learned a complex visuo-spatial task (juggling) revealed increased diffusion anisotropy of the white matter underneath the intraparietal sulcus within a 6-week period (Scholz et al., 2009). Similarly, diffusion anisotropy of the white matter of the prefrontal cortex showed a positive correlation with learning performance of a balancing task in subjects within a two-month period (Taubert et al., 2010). These white matter changes are not restricted to motor-related brain regions, as the amount of training in a working memory task correlates positively with diffusion anisotropy of the corpus callosum and the white matter underneath the intraparietal sulcus (Takeuchi et al., 2010). In an animal model of spatial learning and memory which involved learning to navigate a water maze, there was evidence of decreased diffusivity in several brain structures, including the dentate gyrus, and the piriform and cingulate cortices, as compared to their baseline assessed five days earlier (Blumenfeld-Katzir et al., 2011). These changes were related to an increased immunoreactivity to synaptophysin and glial fibrillary acidic protein, as well as a substantial increase in the

number of astrocytic processes. In the same study, the corpus callosum showed a significant increase of diffusion anisotropy following training, which was related to an increase of myelin basic protein observed by immunohistochemistry. In follow-up human studies, it was shown that the MD of the hippocampus and parahippocampus decreased after being involved in only 2 h of a spatial learning task (playing a car-racing video game) (Sagi et al., 2012), and that these changes are echoed in the fornix (Hofstetter et al., 2013). These findings were supported by a similar paradigm in rodents, in which the same diffusion decreases were found, and were related to an increased number of synaptic vesicles, astrocytic processes, and expression of brain-derived neurotrophic factor evaluated through immunohistochemistry (Sagi et al., 2012) (Fig. 3). Other plastic changes such as synaptic reorganization and axonal sprouting can also occur in response to brain injury, as is the case of the hippocampus following status epilepticus (Sutula, 2002). In that case, the coherent organization of the newly formed axonal processes is capable of increasing diffusion anisotropy, as seen through high-resolution diffusion MRI, both ex-vivo and in the living animal (Kuo et al., 2008; Laitinen et al., 2010). Moreover, diffusion MRI is sensitive enough to demonstrate tissue abnormalities that lie beyond the classically recognized sites of alterations following status epilepticus in rats (Sierra et al., 2011; van Eijsden et al., 2011). Although the

resolution needed for its application in humans is still not sufficient, these findings could be of great relevance for the diagnosis and follow-up of patients with temporal lobe epilepsy (Sutula et al., 1989) and other forms of brain injury. For a review of plastic changes of gray and white matter, evidenced with diffusion MRI and other neuroimaging methods, the reader is referred to (Zatorre et al., 2012).

WHEN THE TENSOR MODEL IS NOT ENOUGH

With the recognition that using typical resolution nearly 90% of the white matter voxels in the human brain contain crossing fibers (Jeurissen et al., 2012) and that the tensor model is inadequate under these circumstances, there is a definite need to find alternative ways to analyze the diffusion MRI signal that conveys information about complex tissue microstructure. There are several methods to deal with this problem (Tournier et al., 2011), but all have one thing in common, and that is their high demands in terms of data acquisition compared to DTI. Some *q*-space methods require several dozen diffusion gradient directions with one or more high *b* values (~2000–50,000 s/mm²), which potentially translate into prohibitively long acquisition times (> 25 min). Fortunately, recent developments in hardware, pulse sequence design and analytical methods (Tuch et al., 2002; Prčková et al., 2013) are progressively reducing

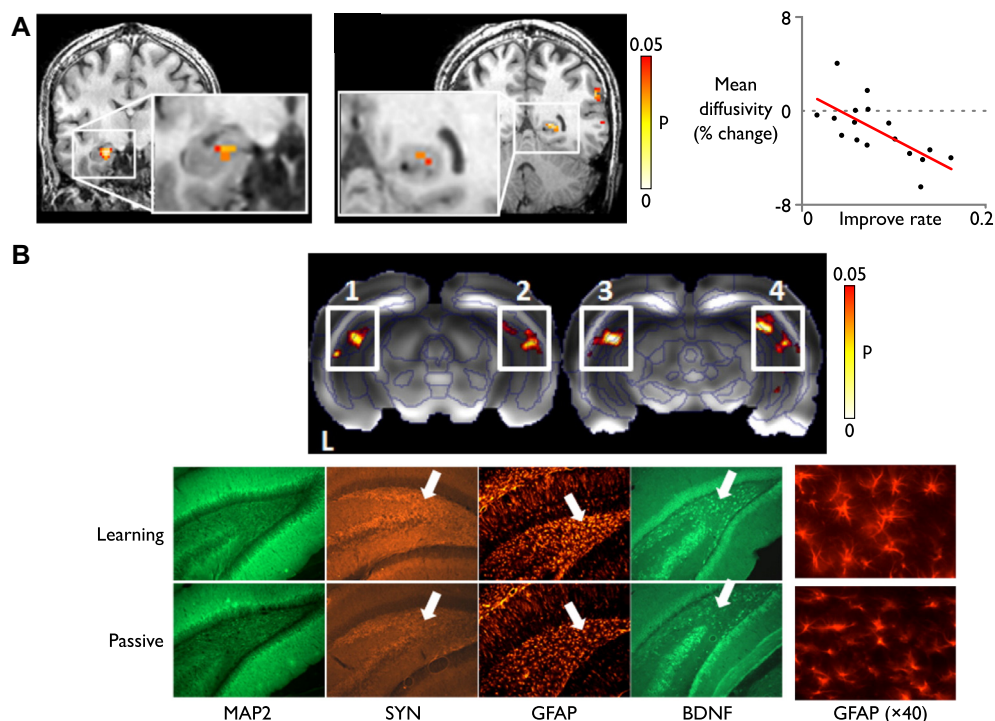


Fig. 3. Changes in mean diffusivity due to training. (A) Decreased mean diffusivity of the left hippocampus and right parahippocampus following 2 h of training on a spatial learning and memory task. There was a negative correlation between the subject's behavioral improvement rate and the change in mean diffusivity of the right parahippocampus (scatter plot). (B) Using a water maze task in rodents, there was also evidence of decreased diffusivity in homologous regions in trained animals, as compared to the passive group (top row; four of five clusters found are shown, 1–4). Immunohistochemical evaluation of the hippocampus (mid- and bottom rows) demonstrated regional differences (arrows) in immunoreactivity of synaptophysin (SYN), glial fibrillary acid protein (GFAP) and brain-derived neurotrophic factor (BDNF), but not microtubule-associated protein 2 (MAP2, a marker of dendrites) between the learning and passive group. Adapted, with permission from the author, from Sagi et al. (2012).

scan time, and the alternatives to DTI are becoming feasible in the clinical realm (Reijmer et al., 2012, 2013a,b; Liégeois et al., 2013). The advantages of non-tensorial approaches to tractography are evident and indisputable (Farquharson et al., 2013), and it is often absolutely necessary to be able to resolve crossing fibers in order to identify certain white matter bundles, such as the acoustic radiation, or the lateral aspects of the cortico-spinal tracts (Behrens et al., 2007). At the single voxel level it is possible to obtain diffusion anisotropy metrics similar to FA, although their biological interpretation has not been as thoroughly studied as their tensor counterpart (Fritzsche et al., 2010). For example, the generalized fractional anisotropy (GFA) index (Tuch, 2004), defined as the standard deviation divided by the root mean square of the diffusion orientation density function, provides a straightforward replacement for FA in non-tensorial data. By design it captures the anisotropy of all the fiber populations at once and, therefore, is unable to extricate tissue abnormalities in separate fiber populations. The construction of the entire diffusion probability density function (PDF) through one of the q -space methods [e.g., diffusion spectrum imaging (Wedeen et al., 2005) or q -ball imaging (Tuch, 2004)] allows the extraction of parameters that can be altered by pathology or development, such as the zero displacement probability, full width at half maximum (FWHM), root mean square displacement and kurtosis excess. For example, during the natural process of myelination of the spinal cords in rats, the PDF of the white matter becomes increasingly narrow, yielding increasing values of zero displacement probability and decreasing values of FWHM (Assaf et al., 2000). In contrast, an increased FWHM is associated with axonal damage and myelin degradation (Assaf et al., 2003; Biton et al., 2005, 2007). Excess kurtosis, particularly parallel to white matter fibers, is a highly sensitive indicator of degeneration following axotomy in a rodent model, both in the acute and chronic phases of degeneration (Farrell et al., 2010). In patients with multiple sclerosis, levels of N-acetylaspartate (a neuronal marker assessed by MR spectroscopy) correlates with zero displacement probability and FWHM not only in regions of white matter with frank abnormalities, but also in so-called normal-appearing white matter. Further, q -space diffusion parameters are more sensitive than DTI to detect these abnormalities (Assaf et al., 2005).

Using multiple b value images, the tensor model can be extended into diffusion kurtosis imaging (Jensen et al., 2005; Jensen and Helpert, 2010; Wu and Cheung, 2010), which provides similar information to the excess kurtosis metric derived from the diffusion PDF. More complex models have also been proposed that take into consideration the geometrical properties of tissue and assign different compartments in which diffusion behavior varies (Stanisz et al., 1997; Peled et al., 1999; Assaf et al., 2004; Jespersen et al., 2007; Alexander, 2008). In the composite hindered and restricted model of diffusion (CHARMED) (Assaf et al., 2004), the diffusion signal is separated into a

fast-diffusing component, assumed to be extra-axonal, and a slow-diffusing component, assumed to be intra-cellular. The extra-axonal space poses some barriers to free diffusion and can indeed produce anisotropy, but with longer diffusion times, the mean water displacement increases linearly. In contrast, the slow-diffusing, intra-axonal water encounters barriers perpendicular to the fibers that cannot be overcome (i.e., the axolemma and myelin sheaths) and as diffusion times are increased, the water displacement gradually increases toward a plateau. This separation of intra- and extra-axonal diffusivities provides more detailed and accurate estimates of fiber microarchitecture. Further, an extension of the CHARMED model, AxCaliber, exploits the restricted diffusivity to infer the distribution of axon diameters within a given voxel, which correspond closely to those derived from electron microscopy (Assaf et al., 2008; Barazany et al., 2009). While the orientation of the fibers must be known a priori to derive axon diameters from AxCaliber, recently introduced restriction spectrum imaging (White et al., 2013) uses high b -value data to study the behavior and orientation of water diffusion within and outside neurites (axons and dendrites), thus providing a more thorough description of tissue organization. In an ex vivo preparation of rodent brain, restriction spectrum imaging was able to capture important structural details of cortical layers that corresponded to histology but were not evident through other diffusion MRI methods. Data requirements must be less strict when imaging the living human brain, for which adaptations of these analytical methods have been made. For instance, the average axonal diameter and density of large white matter bundles can be determined within acceptable acquisition times (Alexander et al., 2010) – while simulations showed the method performed well, and the results in humans are in line with known histological features, there is no direct evidence that the derived axonal metrics correspond to histology. Also, neurite dispersion and density (Jespersen et al., 2007) have been estimated in humans in under 30 min using high angular resolution and two non-zero b values (Zhang et al., 2012). As fractional anisotropy can be affected similarly by increased fiber dispersion or reduced fiber density (Tournier et al., 2011), each with very different origins and meanings, the ability to disambiguate the two conditions can provide important biological information, and the relatively short scan times required to acquire all the necessary data make this technique appealing for clinical research.

Using a single non-zero b value and multiple gradient directions, the fiber orientation distribution function (fODF) can be obtained through spherical deconvolution (Tournier et al., 2004). The amplitude of the fODF is proportional to the intra-axonal fraction, as well as to axonal diameter and spacing (Dell'Acqua et al., 2012; Raffelt et al., 2012). Convincing results have been obtained by simulation of fibers, and the applicability of this method has been shown in human data, proving it to be more sensitive than DTI parameters to identify

pathological changes in motor neuron disease (Raffelt et al., 2012). Nonetheless, no study has yet confirmed that the suspected mechanisms driving the changes in fODF amplitude truly correspond to histological changes.

CONCLUSIONS

Diffusion MRI is a powerful tool for the non-invasive assessment of tissue microstructure; it cleverly exploits the information regarding the behavior of water diffusion, using it to probe the microscopic environment while reporting at a macroscopic scale. It provides a window into the microscopic realm at each image voxel, while at the same time enabling the identification of macroscopic, long-range white matter pathways through tractography. The relative ease with which it can be applied to study the human brain has made it very popular among scientists and clinicians, and the technique is now available on virtually all modern scanners. As with any powerful tool, however, one must be aware of its reach and its limitations, keeping in mind that any inferences made about tissue microstructure arise from an indirect measurement, and must consider all the factors (biological and artifactual) that may influence the diffusion signal prior to making bold claims about white matter integrity.

The diffusion tensor model has been largely responsible for the widespread adoption of diffusion MRI since its introduction nearly two decades ago, likely due to its simplicity and clear biological interpretation. While DTI remains a valid tool for the analysis of particular brain regions and for testing carefully considered hypotheses, its limitations force us to look for extensions or alternatives for the analysis of diffusion MRI. As such, more complete models or even model-free approaches (some of them as seasoned as the tensor model) are gradually gaining a central role in the study of brain connectivity and tissue characteristic. As data acquisition times are shortened, imaging hardware is improved (or diffusion gradient requirements are relaxed), and histological correlates of complex water diffusivity are understood, these advanced methods are likely to dominate the diffusion MRI literature, and will surely provide us with more accurate and complete descriptions of human white matter microstructure *in vivo* during normal development and in the presence of pathology, recovery and plasticity.

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