

# Modeling cortical architectonic features by analyzing diffusion MRI data in the cortical reference frame

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## Synopsis

We quantified the alignment between the DTI reference frame (DRF) and the cortical reference frame (CRF) throughout the entire cerebral cortex in a macaque brain, and found relatively good correspondence, especially in regions with high curvature such as the gyral walls and the cortical sulci. Based on this correspondence, we analyze cortical diffusion signals in the CRF and construct a simple model of cortical diffusion with distinct radial (columnar) and tangential (sheet-like) diffusion processes in cortical layers. The variation of model parameters with cortical depth reflects architectonic features described in a histologically defined digital macaque brain atlas.

## INTRODUCTION

Diffusion MRI (dMRI) analysis in anisotropic tissues such as white matter (WM) can be simplified considerably by using the diffusion tensor imaging (DTI)<sup>1</sup> reference frame (DRF). The DRF is coincident with the dominant orientation of the underlying tissue microanatomy and provides a basis for making useful simplifying assumptions in the construction of many WM tissue models, such as CHARMED<sup>2</sup> or AxCaliber<sup>3</sup>. In tissues with low anisotropy, such as gray matter (GM), due to the similarity of the principal diffusivities, the DRF is poorly defined and prone to sorting bias<sup>4</sup> preventing the construction of a continuous and anatomically-consistent DRF tensor field approximation. Nevertheless, recent studies<sup>5-7</sup> suggest that at high spatial resolution diffusion anisotropy in the cortex varies with the folding geometry, i.e., the cortical reference frame (CRF), showing preferentially radial and tangential components<sup>8-10</sup> which evoke cortical columns and layers<sup>11,12</sup>, respectively, that can be observed with post-mortem histological staining.

We conduct a whole-cortex analysis of the alignment between the DRF and the CRF in the macaque brain and explore the possibility of employing the latter to construct eloquent, simplified models of water diffusion in the cortex. We analyze tissue model parameters in cortical regions-of-interest (ROIs) obtained from a histologically-defined macaque brain atlas<sup>13,14</sup>. dMRI models that reveal columnar (radial) and sheet-like (tangential) diffusion components in the cortex could automate *in vivo* cortical architectonic mapping, improve the clinical characterization of neuroinflammatory and neurodegenerative diseases, and advance our ability to study the developmental timelines of cortical cyto- and myelo-architecture<sup>15,16</sup> *inter alia*.

## METHODS

We acquired 101 diffusion-weighted images (DWIs) of a perfusion-fixed macaque brain<sup>17</sup> at 7T using a 250µm isotropic resolution, FOV=78x64x72cm, TE/TR=33.3/250ms. We used multiple b-values (100,600,1500,2800,4800,7200,10000s/mm<sup>2</sup>) with gradient orientations (3,4,8,12,18,24,32 respectively) uniformly sampling the unit sphere for each b-shell and across shells, and gradient pulse parameters δ=8ms and Δ=16.1ms. We also conducted a magnetization transfer (MT) prepared gradient-echo experiment, segmented<sup>18</sup> the WM and GM, reconstructed the GM/WM and pial cortical surfaces<sup>19</sup> and computed intermediate surfaces corresponding to cortical layers using the equivolumetric principle<sup>20,21</sup>. We registered<sup>22</sup> the histologically-defined D99 digital rhesus macaque brain atlas<sup>13,14</sup> to the EPI distortion-corrected DWIs<sup>23</sup> allowing for correlation analysis between dMRI parameters and histological stains in corresponding cortical areas.

We quantified the relative alignment between the CRF and the DRF throughout the cortex by measuring the radial and tangential deviations, angles  $\theta$  and  $\phi$ , respectively (Fig.1), and found relatively small deviations which justified dMRI analysis in the CRF. Consequently, we analyzed the DWIs (interpolated at each vertex of each layer surface) in the CRF using a simple two-component “stick-and-disc” tissue model that accounts for separate (non-exchanging) radial and tangential diffusion processes in each cortical layer (Fig.2):

$$E_c(\mathbf{g}) = f e^{-\mathbf{g}^T (\hat{\mathbf{n}} \hat{\mathbf{n}}^T) \mathbf{g} D_r} + (1-f) e^{-\mathbf{g}^T (\hat{\mathbf{c}}_1 \hat{\mathbf{c}}_1^T + \hat{\mathbf{c}}_2 \hat{\mathbf{c}}_2^T) \mathbf{g} D_t}$$

where  $E_c(\mathbf{g})$  is the diffusion signal attenuation as a function of the applied gradient  $\mathbf{g}$ ;  $f$  represents the signal fraction of the radial diffusion component; while the scalars  $D_r$  and  $D_t$  define the cylindrically symmetric degenerate rank-1 (radial) and rank-2 (tangential) diffusion tensors aligned with the cortical surface normal,  $\hat{\mathbf{n}}$ , and the tangent plane defined by the minimum and maximum Gaussian curvature orientations,  $\hat{\mathbf{c}}_1$  and  $\hat{\mathbf{c}}_2$ , respectively (Fig.1). We computed cortical depth profiles of  $f$ ,  $D_r$  and  $D_t$  and quantified their statistics in regions-of-interest (ROIs) obtained from the histologically-defined D99 macaque brain atlas.

## RESULTS

Matching the axes of the DRF and CRF (Fig.1) in regions with very low anisotropy, allows reordering of the diffusion tensor axes (i.e., sorting of the principal diffusivities) to produce a more continuously-varying tensor field approximation in the cortex. Moreover, the relatively small deviations between the DRF and CRF (Fig.2) suggest that the CRF may provide a well-defined anatomically-consistent and continuous reference frame for use in dMRI analysis, especially in regions with high curvature.

The largest misalignment is observed in regions with negative curvature (i.e., gyral crowns) and low diffusion anisotropy (e.g., the superior temporal gyrus) where the DRF is poorly-defined (Fig.2). Significantly better alignment can be observed in cortical areas with high curvature along the gyral walls and in the sulci.

The mid-cortical layer shows the largest values of  $f$  (Fig.4), consistent with the increased presence of radial projections in histological observations of cortical myeloarchitecture (Fig.3). The largest difference between  $D_r$  and  $D_t$  was also found in the mid-cortical layers, suggesting the presence of strong diffusion processes along cortical columns, in agreement with previous findings<sup>8-10</sup> (Fig.4).

## DISCUSSION AND CONCLUSIONS

Our results point to a remarkable correspondence between structural and functional reference frames across multiple length scales. The DRF describes a physiological process (water diffusion) at the microscopic scale ( $\sim 5\mu\text{m}$ ) while the CRF characterizes brain anatomy (cortical folding) at the macroscopic and mesoscopic scales ( $\sim 500\mu\text{m}$ ). The correspondence between these two reference frames (**Fig.2**) suggests that biological structures at the meso- and macroscopic scales may arise from processes at the microscopic scale, which would determine local transport properties, particularly the diffusion of various molecules (e.g., growth factors) during development<sup>15</sup>.

Concurrently, our results also imply that, from the cortical surface geometry, one could infer information about the microscopic tissue organization, which may simplify dMRI analysis within cortical areas, and among cortical layers, primarily by reducing the number of degrees of freedom in analyzing signals and constructing tissue models.

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