## Singular behavior of time-dependent diffusion in a fiber bundle geometry due to a disordered packing

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Introduction: How important is disorder in the packing geometry of an axonal fiber bundle from the standpoint of a diffusion measurement? Here we show that the randomness in fiber arrangement in a bundle crucially affects diffusion in the extra-axonal space, which can have implications for both pulse-gradient<sup>1</sup> and oscillating gradient methods<sup>2</sup> of characterizing white matter fiber integrity. Specifically, we focus on the time-dependence of the diffusion coefficient, D(t), in the extra-axonal space by diffusion measurements in a phantom made of randomly packed parallel aligned impermeable fibers and Monte-Carlo simulations. We show that D(t) transverse to a fiber bundle has a logarithmic singularity at long diffusion



**Figure 1:** a) Diffusion tensor eigenvalues  $\lambda_1 > \lambda_2 > \lambda_3$  measured in the phantom, as function of diffusion time t. b) Taking  $D(t)=\lambda_2$ , we fit it to Eq (1), and show the difference D(t)- $D_{\infty}$  in log-log scale. The systematic slope change relative to the 1/t fit is a hallmark of logarithmic singularity.



Figure 2: a) This singularity is also manifest in the "bend" in D(t) vs. 1/t using  $\lambda_2(t)$  from Fig.1. **b**) The logarithmic "bend" straightens up when D(t) is replotted with respect to  $\ln(t/t_0)/t$ . c) and d) The same as for a and b, but with MC simulated data.



axons are ~1µm in diameter, almost any measurement is in long-time limit, and hence the above singularity may significantly affect the interpretation of timedependent diffusion-based methods of evaluating fiber integrity, as described below. **Methods:** Theory. As recently shown<sup>3</sup>, the power law exponent  $\vartheta$  describing the approach of the tortuosity limit  $D_{\infty}$  can yield information about the type of disorder in a system. This exponent appears both in the instantaneous diffusion coefficient  $D_{\text{inst}}(t)$ , and in OGSE  $\mathcal{D}(\omega)$ 

$$D_{inst}(t) \equiv \frac{1}{2} \frac{\partial}{\partial t} \langle x^2 \rangle = D_{\infty} + at^{-\vartheta} + \dots \quad \Leftrightarrow \quad \mathcal{D}(\omega) = D_{\infty} + b|\omega|^{\vartheta} + \dots \quad (2)$$
  
In two dimensions,  $\vartheta = 1$  for the most commonplace random packing characterized

by short-ranged disorder in fiber placement<sup>3</sup>. The integration  $\hat{J}_{inst}^{4,5}$  of  $D_{inst}(t)$  up to t and dividing by t yields the  $\ln t / t$  behavior of D(t) in Eq. (1) above. Conversely, for any more ordered arrangement (e.g. periodic),  $\vartheta > 1$ , yielding  $D(t) \sim D_{\infty} + c/t$ . Hence,

in a D(t) measurement, the effect of disorder is in the extra ln t factor. Phantom Construction. The diffusion phantom for this study was constructed with approximately

195,000 Dyneema® fibers tightly held together with a shrinking tube measuring 8 cm long. The fibers are  $17\pm2.6 \,\mu\text{m}$  in diameter, ultrahydrophobic and impermeable to water. The fiber bundle was suspended in a 1.5 L plastic bottle filled with a distilled water solution of 0.09% w/v NaCl to reduce B<sub>1</sub> field inhomogeneities.

MRI Measurements. Imaging was performed at 15°C on a 7T Siemens clinical MRI scanner using a 28 channel knee coil. DTI was carried out using a STEAM sequence which allows for long diffusion times while minimizing echo attenuation caused by T2 relaxation. Twenty five measurements were performed at b values of 0 and 500 in 20 directions, each with TE of 57 ms and TM ranging from 10ms to 1000 ms, corresponding to diffusion times, t, of 38.5 ms to 1028.5 ms. Three slices of resolution  $3 \text{ mm} \times 3 \text{mm} \times 10 \text{ mm}$  were used. The fiber bundle was placed parallel to the B<sub>0</sub> field to eliminate the possibility of internal field inhomogeneties.

**Results:** The DTI eigenvalues are plotted vs t in Fig. 1a. We focus on the second eigenvalue, D(t) $= \lambda_2$ . Fig. 1b shows a plot of  $D(t) - D_{\infty}$  vs t on a log-log scale, along with a fit of  $D(t) - D_{\infty}$  to c  $\ln(t/t_0)/t$  (black dashed line) and a fit to c/t (red dashed line), showing clearly that the c/t fit is insufficient to properly describe the data. Fig. 2a and b show the same D(t) data plotted with respect to 1/t and  $\ln(t/t_0)/(t/t_0)$ , respectively. In Fig. 2a, a slight curve can be seen in the data indicating the logarithmic singularity. Fig. 2b shows that the bend is removed when plotted with respect to  $\ln(t/t_0)/(t/t_0)$ ,  $t_0 = 11$  ms. Fig. 2c and d show Monte Carlo simulation data using a free diffusion coefficient of 2  $\mu$ m<sup>2</sup>/s, and fiber size distribution centered around 17  $\mu$ m, with  $t_0 = 7.3$ ms, in agreement with experiment.

Discussion: Implication for PGSE: The diffusion restricted inside axons, giving the 1/t contribution, is used<sup>1</sup> to probe internal diameter distribution. However, as we have shown, the t-dependence in the extra-axonal space is more relevant, as  $\ln(t)/t$  eventually exceeds 1/t in long-t limit. Hence, modeling the disorder in extra-axonal space is essential for interpreting such measurements.

Implication for OGSE: The non-analytic  $\ln(t)/t$  in D(t) translates into linear behavior in  $|\omega|$  (a sharp kink in  $D(\omega)$  for near  $\omega = 0$ ). Indeed, such sharp non-parabolic behavior is clearly seen is recent OGSE measurements in brain [Fig 6 of ref 6], which may indicate 2-dimensional disorder in extracellular space. For ordered arrangements, or for confined diffusion (e.g. inside axons), the 1/t behavior in D(t) translates into  $\omega^2$  in  $D(\omega)$ .<sup>2,4,7</sup> This parabolic behavior will be less relevant than  $|\omega|$  at small  $\omega$  and the effect of packing disorder will again dominate over that of confined water.

**Conclusions:** The logarithmic singularity in two-dimensional diffusion has been demonstrated for a first time as a hallmark of disordered packing geometries. This singularity dominates the time-dependence of diffusion across axonal fiber bundles and should be included in any quantification scheme for adequate fiber characterization.

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