
Unified representation of tractography and diffusion-weighted MRI data using sparse multidimensional arrays

SUPPLEMENTARY MATERIAL

Cesar F. Caiafa*

Department of Psychological and Brain Sciences
Indiana University (47405) Bloomington, IN, USA
IAR - CCT La Plata, CONICET / CIC-PBA
(1894) V. Elisa, ARGENTINA
ccaiafa@gmail.com

Olaf Sporns

Department of Psychological and Brain Sciences
Indiana University (47405) Bloomington, IN, USA
osporns@indiana.edu

Andrew J. Saykin

Department of Radiology - Indiana University
School of Medicine. (46202) Indianapolis, IN, USA
asaykin@iupui.edu

Franco Pestilli†

Department of Psychological and Brain Sciences
Indiana University (47405) Bloomington, IN, USA
franpest@indiana.edu

1 Proofs of the theoretical bound for accuracy and compression factor

In this section, we derive a theoretical bound on the accuracy of LiFE_{SD} compared to the original LiFE model (Proposition 3.1) and we theoretically analyze the compression factor associated to the factorized tensor approximation (Proposition 3.2). Hereafter, we assume a given connectome having N_f fascicles, each fascicle having a fixed number of N_n nodes, and where diffusion weighted measurements were taken on N_θ gradient directions with a gradient strength b .

Proof of Proposition 3.1: The error in modeling the diffusion signal for a particular voxel v , fascicle f and gradient direction θ is given by:

$$\Delta_{\mathbf{O}}(\theta) = |\mathbf{O}_f(\theta) - \mathbf{D}(\theta, a)|, \quad (\text{S1})$$

where $\mathbf{O}_f(\theta)$ is the orientation distribution function as defined in equation (2.3) (we avoided making reference to the voxel v for clarity) and $\mathbf{D}(\theta, a)$ is the diffusion signal of atom a

*<http://web.fi.uba.ar/~ccaiafa/Cesar.html>

†<http://www.brain-life.org/plab/>

at gradient direction $\boldsymbol{\theta} = [\theta_x, \theta_y, \theta_z]^T$. By defining $\mathbf{v} = [v_x, v_y, v_z]^T$ and $\mathbf{v}_a = \mathbf{v} + \boldsymbol{\Delta}_v = [v_x + \Delta_{v_x}, v_y + \Delta_{v_y}, v_z + \Delta_{v_z}]^T$ as the vectors pointing out at the directions of the fascicle f and its closest dictionary atom a , respectively (see Fig. 3c), we arrive at:

$$\Delta_{\mathbf{O}}(\boldsymbol{\theta}) = \left| \Delta_g(\boldsymbol{\theta}) - \frac{1}{N_{\boldsymbol{\theta}}} \sum_{\boldsymbol{\theta}'} \Delta_g(\boldsymbol{\theta}') \right|, \quad (\text{S2})$$

where $\Delta_g = |g(\mathbf{v}_1 + \boldsymbol{\Delta}_v, \boldsymbol{\theta}) - g(\mathbf{v}, \boldsymbol{\theta})|$ with $g(\mathbf{v}, \boldsymbol{\theta}) = e^{-b(\boldsymbol{\theta}^T \mathbf{v})^2}$. For a sufficiently small error vector $\boldsymbol{\Delta}_v = [\Delta_{v_x}, \Delta_{v_y}, \Delta_{v_z}]^T$, we can approximate $\Delta_g(\boldsymbol{\theta})$ as follows:

$$\Delta_g(\boldsymbol{\theta}) \approx \left| \frac{\partial g(\mathbf{v}, \boldsymbol{\theta})}{\partial v_x} \right| \Delta_{v_x} + \left| \frac{\partial g(\mathbf{v}, \boldsymbol{\theta})}{\partial v_y} \right| \Delta_{v_y} + \left| \frac{\partial g(\mathbf{v}, \boldsymbol{\theta})}{\partial v_z} \right| \Delta_{v_z}, \quad (\text{S3})$$

and, by using the fact that $|\boldsymbol{\theta}^T \mathbf{v}| \leq 1$, $e^{-b(\boldsymbol{\theta}^T \mathbf{v})^2} \leq 1$, $\Delta_{v_x}, \Delta_{v_y}, \Delta_{v_z} \leq \|\boldsymbol{\Delta}_v\| \leq \frac{\pi}{\sqrt{2}L}$, and $\|\boldsymbol{\theta}\|_1 \leq \sqrt{3}\|\boldsymbol{\theta}\|$ in equation (S3), we obtain: $\Delta_g(\mathbf{v}, \boldsymbol{\theta}) \leq \frac{b\pi\sqrt{6}}{L}$. Thus, by using this result in equation (S2), we obtain an upper bound for the error of modeling the diffusion signal of one fascicle and at one gradient direction in a voxel: $\Delta_{\mathbf{O}}(\boldsymbol{\theta}) \leq \frac{2b\pi\sqrt{6}}{L}$. Finally, by summing up all over the nodes in the connectome, it implies that

$$\|\underline{\mathbf{M}} - \hat{\underline{\mathbf{M}}}\|_F^2 \leq N_f N_n N_{\boldsymbol{\theta}} \left(\frac{2b\pi\sqrt{6}}{L} \right)^2. \quad (\text{S4})$$

□

Proof of Proposition 3.2: The memory load necessary to store each fascicle in a sparse matrix \mathbf{M} is $3N_{\boldsymbol{\theta}}N_n$, because using a sparse matrix structure, three numbers are required for each node, i.e., the row-column indices plus the entry value. Thus the storage cost of \mathbf{M} is:

$$C(\mathbf{M}) = \mathcal{O}(3N_n N_{\boldsymbol{\theta}} N_f). \quad (\text{S5})$$

Conversely, storing fascicles in the LiFE_{SD} model requires $4N_n$ values per fascicle plus the dictionary matrix (i.e., the set of the non-zero entries and their locations within the tensor $\underline{\Phi}$ plus matrix \mathbf{D}). Thus, the amount of memory required in the LiFE_{SD} model is:

$$C(\hat{\underline{\mathbf{M}}}) = \mathcal{O}(4N_n N_f + N_{\boldsymbol{\theta}} N_a), \quad (\text{S6})$$

where $N_{\boldsymbol{\theta}} N_a$ is the storage associated with the dictionary matrix $\mathbf{D} \in \mathbb{R}^{N_{\boldsymbol{\theta}} \times N_a}$. Finally, by taking the ratio of equations (S5) and (S6), we arrive at the expression of the compression factor as shown in equation (3.7). □