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Non central chi estimation of multi-compartment models improves model selection by reducing overfitting

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PURPOSE – Diffusion images are known to be corrupted with a non-central chi (NCC)-distributed noise [1]. There has been a number of proposed image denoising methods that account for this particular noise distribution [1,2,3]. However, to the best of our knowledge, no study was performed to assess the influence of the noise model in the context of diffusion model estimation as was suggested in [4]. In particular, multi-compartment models [5] are an appealing class of models to describe the white matter microstructure but require the optimal number of compartments to be known a priori. Its estimation is no easy task since more complex models will always better fit the data, which is known as over-fitting. However, MCM estimation in the literature is performed assuming a Gaussian-distributed noise [5,6]. In this preliminary study, we aim at showing that using the appropriate NCC distribution for modelling the noise model reduces significantly the over-fitting, which could be helpful for unravelling model selection issues and obtaining better model parameter estimates.

METHODS – Noise in diffusion MRI is NCC-distributed as 

\[ f(S_j|A_j, S_0, \sigma^2) = \frac{2}{\sqrt{2\pi}} \frac{1}{\sigma^2} e^{-\frac{\sigma^2}{2\sigma^2} (S_j - A_j)^2} I_{L-1} \left( \frac{2 \sigma^2}{\sigma^2} \right), \]

where \( L \) and \( \sigma^2 \) are respectively the effective number of coils and effective variance of each coil that depend on the coils’ correlations, the acceleration factor of the acquisition and the signal reconstruction method [3,4]. \( S_j \) is the reconstructed measured signal, \( A_j \) is the signal attenuation predicted by the model and \( I_{L-1} \) is the \((L-1)\)-order modified Bessel function of the first kind. We set up an NCC estimation framework that can be used to fit any diffusion model that provides an analytic expression of the predicted signal attenuation \( A_j \). Starting from \( N \) acquired diffusion images, we compute the NCC log-likelihood by replacing \( A_j \) with the appropriate expression and then we maximize it using a derivative-free optimizer. Researchers usually rely on the assumption of high SNR to resort to the Gaussian approximation of the noise. However, at high SNRs, the NCC log-likelihood reads 

\[ \log L = \log \left( 2\pi e^{-1/2} \exp \left( -\sum_{i=1}^{N} (S_i - A_i)^2 / (2\sigma^2) \right) \right) - (2L - 1)/2 \sum_{i=1}^{N} \log(S_0 A_i / S_i), \]

which does not boil down to the Gaussian log-likelihood (blue term). Indeed, the red term penalizes any model that predicts a signal greater than the observed one, i.e., any model that predict physically implausible signals. For smaller SNRs, the penalty can even dominate the likelihood. Gaussian noise does not worry for these physical incoherences.

RESULTS – The objective of the experimental design was to demonstrate that accounting for the real nature of the noise in diffusion images helps in unravelling the model selection problem by reducing over-fitting. We used two cases provided by the Human Connectome Project [7]. For each case, the data consisted in 270 DW images with three different b-values at 1000, 2000 and 3000 s/mm² and 18 non-DW images, at a spatial resolution of 1.25 × 1.25 × 1.25 mm³. We estimated the same ball-and-stick models with increased number of fascicle compartments using both Gaussian and NCC (with \( L=32 \) degrees of freedom) noise models and performed a model selection based on Akaike information criterion for each noise model to output maps of the optimal number of fascicles voxelwise. We proposed a qualitative assessment of the resulting maps and we reported, for each noise model, the proportion of 0, 1, 2 and 3 fascicle voxels across subjects.

CONCLUSION – We set up an NCC framework for the estimation of multi-compartment models and performed a preliminary study to understand whether it is important to account for the real NCC nature of the noise or if the usual Gaussian approximation yields equivalent results. In this work, we specifically compared the maps of optimal number of fascicles obtained assuming Gaussian noise instead of NCC noise yields a lot of over-fitting that favors the 3-fascicle model. As a result, 1-fascicle regions are much better delineated using NCC noise. Specifically, using the exact same fitting procedure with only an improved noise model, the proportion of 0,1,2-fascicle voxels (models with low complexity) is increased and the proportion of 3-fascicle voxels is decreased as summarized in Table 1.

![Figure 1: Coronal view of the maps of optimal number of fascicles for subject #1 (a,b) and #2 (c,d); assuming Gaussian noise (a,c) and NCC noise (b,d).](image)

<table>
<thead>
<tr>
<th># Fascicles</th>
<th>0</th>
<th>1</th>
<th>2</th>
<th>3</th>
</tr>
</thead>
<tbody>
<tr>
<td>Gaussian (%)</td>
<td>9</td>
<td>4</td>
<td>27.5</td>
<td>59.5</td>
</tr>
<tr>
<td>NCC (%)</td>
<td>18.5</td>
<td>7.5</td>
<td>32</td>
<td>42</td>
</tr>
</tbody>
</table>

Table 1: Averaged proportion of 0,1,2,3-fascicle voxels across subjects