Fibre dispersion in the corpus callosum relates to interhemispheric functional connectivity

Presented During: Poster Session Thursday, June 29, 2017: 12:45 PM - 02:45 PM

Stand-By Time

Thursday, June 29, 2017: 12:45 PM - 2:45 PM

Submission No:

4019

Submission Type:

Abstract Submission

On Display:

Wednesday, June 28 & Thursday, June 29

Authors:

<u>Jeroen Mollink</u>^{1,2}, Saad Jbabdi², Stephen Smith², Fidel Alfaro-Almagro², Michiel Kleinnijenhuis², Anne-Marie van Cappellen van Walsum¹, Karla Miller²

Institutions:

¹Donders Institute for Brain, Cognition and Behaviour, Radboud University Medical Center, Nijmegen, Netherlands, ²FMRIB centre, University of Oxford, Oxford, United Kingdom

First Author:

Jeroen Mollink - Lecture Information | Contact Me

Donders Institute for Brain, Cognition and Behaviour, Radboud University Medical CenterlFMRIB centre, University of Oxford Nijmegen, NetherlandslOxford, United Kingdom

E-Poster

Introduction:

The corpus callosum (CC) plays a key role in cross-hemispheric information transfer. Although the CC is generally thought of as a well-organized bundle of parallel axons connecting homotopic areas, recent studies have demonstrated complex fibre architecture within the CC¹⁻³. In particular, fibre orientation dispersion at the midline is greater than in nearby lateral areas of the CC⁴. This might reflect interhemispheric connections of non-homotopic (heterotopic) areas. Here we hypothesize that dispersion in the CC is a signature of heterotopic connectivity. We evaluated this by looking for associations between diffusion MRI (dMRI) estimates of dispersion and resting-state functional MRI (rfMRI) estimates of interhemispheric connectivity.

Methods:

We analyzed dMRI and rfMRI data from 4903 subjects in the UK Biobank project (acquisition parameters and pre-processing detailed elsewhere)⁵.

Group-ICA yielded 55 resting-state networks. These were then split for each hemisphere and further split if a component contained non-contiguous brain areas. To estimate connectivity between the resulting brain areas (network 'nodes'), average time-series were generated for every node and correlated with each other using partial correlation. These connectivity values then form a node-by-node matrix of network connectivity. To quantify the balance of homo- to heterotopic interhemispheric connections for a given node, a functional asymmetry index (AI) was defined as the homotopic connection strength minus the mean of the 10% strongest heterotopic connections.

To calculate orientation dispersion (OD), the NODDI model⁶ was fit to dMRI data. OD values were extracted from the CC after transforming all subjects to the same space and mapping ODs onto a skeleton using TBSS⁷. To estimate which CC area connects

homotopic areas, structural connectivity probability maps for each node were obtained in the CC using a tractography template. Dispersion was calculated as the mean OD at the center minus the mean OD at the lateral aspects of the CC, referred to as OD hereafter (Figure 1.C).

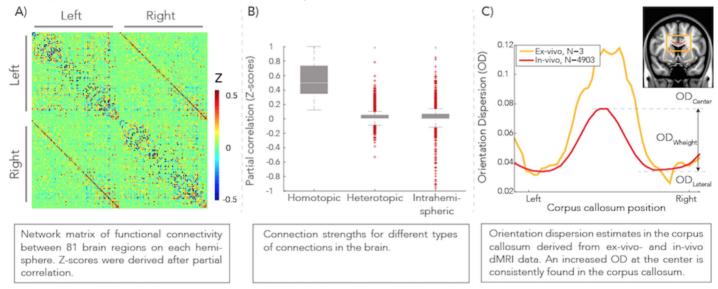
Associations between OD and AI were calculated, with statistical significance assessed using permutation testing. Reproducibility was demonstrated by dividing the subjects into two equally sized groups. For each AI vs OD comparison, the OD values were taken from a region of the CC that connects to the cortical node of interest. A negative control analysis was done by correlating AIs with OD derived from the CC region that is least likely to interconnect a node.

Results:

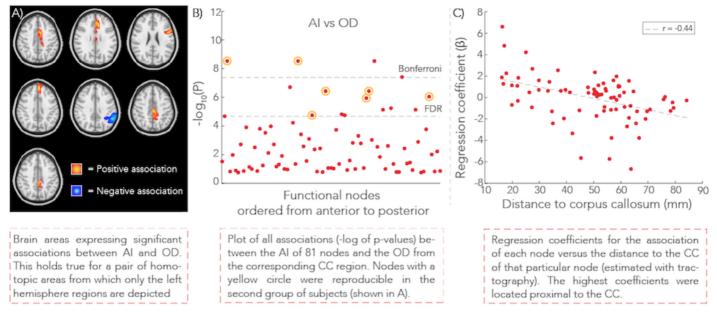
As reported previously⁴, high dispersion was found in the center of the CC (Figure 1.C). Figure 1.A illustrates the group-average functional network matrix.

Significant associations between AI and OD were primarily found in medial nodes, with correlation coefficients ranging between r=0.05-0.11 (Figure 3). Of 16 significant associations in the "discovery" group of subjects (that survive FDR multiple comparison correction), seven were found in the replication group. The highest regression coefficients (β) were found closer to the CC, and lower regression coefficients distal to the CC (Figure 3.D).

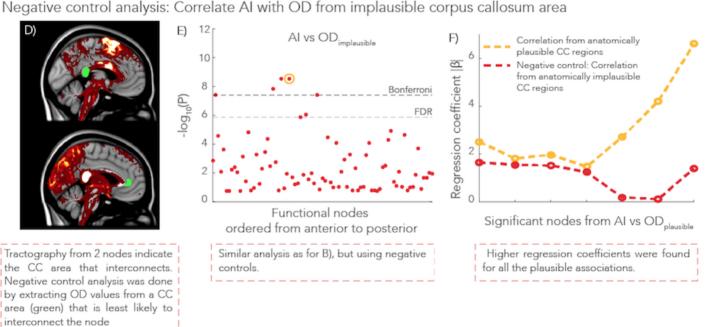
Finally, the negative control analysis yielded one significant association between AI and OD in an implausible CC region (Figure 4), with a lower regression coefficient (β =1.55) than the anatomically plausible CC area (β =1.82).



·Figure 1



Negative central analysis Carrelate Al with OD from implemental across collegem area



·Figure 2

Conclusions:

Significant associations were found between AI vs OD in regions close the CC, such as the cingulate cortex. In addition, the AI of a node appeared to be less associated with OD if the node was more laterally located. We hypothesize that dispersion in the CC is more crucial for midline nodes to project to nearby heterotopic regions, while more distant nodes can form a path to heterotopic areas in other tracts. The negative control analysis yielded one significant node that may be explained by correlation of ODs among different areas of the CC.

Modeling and Analysis Methods:

Diffusion MRI Modeling and Analysis ² fMRI Connectivity and Network Modeling ¹

Poster Session:

Poster Session - Thursday

Keywords:

FUNCTIONAL MRI Tractography White Matter

WHITE MATTER IMAGING - DTI, HARDI, DSI, ETC Other - Dispersion

^{1|2}Indicates the priority used for review

Would you accept an oral presentation if your abstract is selected for an oral session?

Yes

I would be willing to discuss my abstract with members of the press should my abstract be marked newsworthy:

Yes

Please indicate below if your study was a "resting state" or "task-activation" study.

Resting state

By submitting your proposal, you grant permission for the Organization for Human Brain Mapping (OHBM) to distribute the presentation in any format, including video, audio print and electronic text through OHBM OnDemand, social media channels or other electronic media and on the OHBM website.

I accept

Healthy subjects only or patients (note that patient studies may also involve healthy subjects):

Healthy subjects

Internal Review Board (IRB) or Animal Use and Care Committee (AUCC) Approval. Please indicate approval below. Please note: Failure to have IRB or AUCC approval, if applicable will lead to automatic rejection of abstract.

Not applicable

Please indicate which methods were used in your research:

Functional MRI Diffusion MRI

For human MRI, what field strength scanner do you use?

3.0T

Which processing packages did you use for your study?

FSL

Provide references in author date format

- 1. Mikula S. et al (2012). Staining and embedding the whole mouse brain for electron microscopy. Nature Methods.
- 2. Ronen I. et al (2014). Microstructural organization of axons in the human corpus callosum quantified by diffusion-weighted magnetic resonance spectroscopy of N-acetylaspartate and post-mortem histology. Brain Structure and Function.
- 3. Budde M.D. et al (2013). Quantification of anisotropy and fiber orientation in human brain histological sections. Frontiers in integrative neuroscience.
- 4. Mollink J. et al (2016). Exploring fibre orientation dispersion in the corpus callosum: Comparison of Diffusion MRI, Polarized Light Imaging and Histology. Proceedings ISMRM.
- 5. Miller K.L. et al (2016). Multimodal population brain imaging in the UK Biobank prospective epidemiological study. Nature Neuroscience.
- 6. Zhang H. et al (2012). NODDI: Practical in vivo neurite orientation dispersion and density imaging of the human brain. Neuroimage.
- 7. Smith S.M et al (2006). Tract-based spatial statistics: Voxelwise analysis of multi-subject diffusion data. Neuroimage.