Alterations in Causal Functional Brain Networks in Alzheimer's Disease: A resting-state fMRI study

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Abstract

Background: Alterations in functional connectivity (FC) of the brain is known to predate the onset of clinical symptoms of Alzheimer's disease (AD) by several decades. Identifying the altered functional brain networks in AD can help in its prognosis and diagnosis.

Objective: FC analysis is predominantly correlational. However, correlation does not necessarily imply causation. This study aims to infer causal functional connectivity (CFC) from functional magnetic resonance imaging (fMRI) data and obtain the sub-networks of CFC that are altered in AD compared to cognitively normal (CN) subjects.

Methods: We used the recently developed Time-aware PC algorithm to infer CFC between brain regions. The CFC outcome was compared with correlation-based functional connectivity obtained by sparse partial correlation. Then, Network-based Statistics (NBS) was used to obtain CFC sub-networks that altered in AD subjects compared to healthy controls while correcting for multiple comparisons at 5% level of significance.

Results: Our findings identified causal brain networks involving the inferior frontal gyrus, superior temporal gyrus (temporal pole), middle temporal gyrus (temporal pole), and different lobes of the cerebellum to be significantly reduced in strength in AD compared to CN group (p-value = 0.0299; NBS corrected). In the sample dataset that has been analysed, no brain networks were found to exhibit significant increase in strength in AD compared to CN group at 5% level of significance with NBS correction.

Conclusions: Our findings provide insights into disruptions in causal brain networks in AD. The corresponding brain regions are in agreement with published medical literature on brain regions impacted by AD. Our work establishes a methodology for finding causal brain networks that are affected by AD using TPC algorithm to compute subject-specific CFC and then using NBS for finding CFC subnetworks that alter between AD and CN groups. Larger datasets are expected to identify further subnetworks affected by AD.

Keywords: Causal inference, functional connectivity, brain mapping, directed graphical modeling, Alzheimer's disease, functional magnetic resonance imaging

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1. Introduction

Alzheimer's disease (AD), a leading cause of dementia worldwide, is characterized by a progressive decline in cognitive function such as impairments to working memory, episodic memory, attention/executive function (de LaCoste and White III, 1993, Christensen et al., 1997). Such cognitive function deficits in AD are believed to be due to abnormality in connectivity between different regions of the brain (Gomez-Isla and Hyman, 1997, Perry and Hodges, 1999, Delbeuck et al., 2003). Evidence from many anatomical and functional studies suggest that AD is a "disconnection syndrome" (Geschwind, 1965, Bozzali et al., 2011, Brier et al., 2014a). Anatomically, AD is characterized by progressive diffuse cortical atrophy, primarily located in the mesial-temporal regions (Yao et al., 2010, Amlien and Fjell, 2014). On the other hand, studies of functional connectivity (FC), which measures the degree of temporal synchrony between the activity of different brain regions, have found evidence of disrupted functional connectivity between brain regions (Wang et al., 2007, Zhao et al., 2020, Biswas and Sripada, 2023).

Several FC analyses measure the degree of statistical association between signals from different brain regions, in other words, associative FC (AFC) (Sheline and Raichle, 2013, Brier et al., 2014b, Briels et al., 2020). However, AFC does not account for the directionality and causality of information flow between regions (Biswas and Shlizerman, 2022a,b). In contrast to AFC, causal functional connectivity (CFC) extends beyond associations and captures the degree of causal interactions between brain regions. Thereby, CFC promises to be more informative than AFC. CFC represents functional connectivity between brain regions by a directed graph, with nodes as the brain regions, directed edges between nodes indicating causal relationships between the brain regions, and weights of the directed edges quantifying the strength of the corresponding causal relationship (Spirtes et al., 2000, Biswas and Shlizerman, 2022a). Yet, there are very few studies that explore causality in functional connectivity (Reid et al., 2019). In this paper, we aim to obtain CFC sub-networks that exhibit significant alteration in AD.

Functional magnetic resonance imaging (fMRI) is a widespread technique to record activity of different brain regions (Logothetis et al., 2001, Smith, 2004, Logothetis, 2008). It measures the Blood Oxygen Level Dependent (BOLD) signal from different brain regions. Studies in functional connectivity have popularly used fMRI data to obtain functional connectivity (Van Den Heuvel and Pol, 2010, Rogers et al., 2007, Kim and Ye, 2020). In an earlier work by Biswas and Sripada (2023), the authors obtained CFC in AD from fMRI data and performed an edge-wise analysis of differences in CFC edges between AD and CN groups. In contrast, in this present paper, we use network-based statistics (NBS) to perform a network-based comparison of the CFC resulting in altered CFC sub-networks in AD compared to CN subjects, while correcting for multiple comparisons in those sub-networks (Zalesky et al., 2010).

In this paper, we use fMRI recordings to obtain CFC for each individual subject. We infer the CFC from fMRI data using the recently developed Time-aware PC algorithm (Biswas and Shlizerman, 2022b). The Time-aware PC algorithm infers causality in neural interactions instead of correlation, in a model-free non-parametric manner, computes

Characteristic	CN	MCI	AD	P
Number of subjects	41	54	34	-
Sex (M/F)	19/22	29/26	16/18	0.16
Age (years)	74.9 ± 6.4	74.2 ± 7.1	74.4 ± 7.4	0.86
Education (years)	16.5 ± 2.3	15.7 ± 2.6	15.4 ± 2.5	0.22
MMSE	29.1 ± 1.4	27.8 ± 1.9	21.9 ± 4.2	$< 10^{-14}$

Table 1: Summary of demographic information and Mini Mental State Examination (MMSE) for CN, MCI and AD subjects. The second to fourth columns present group characteristics, mean ± SD. The fifth column presents P values for the statistical significance of the inter-group differences. Differences in Sex was assessed using a Chi-Squared test and differences in Age, Education and MMSE using non-parametric analysis of variance by Kruskal-Wallis test.

whole brain-wide interactions, incorporates multi-regional interactions, and its utility has been demonstrated in fMRI data (Biswas and Sripada, 2023). We then use network-based statistics (NBS) to obtain CFC subnetworks that significantly alter between the subjects with AD and cognitively normal (CN) subjects, while correcting for multiple comparisons (Zalesky et al., 2010, Zhan et al., 2016, 2019). The resulting brain regions are supported by medical literature on brain regions impacted by Alzheimer's disease.

2. Materials and Methods

2.1. Participants

The resting fMRI and demographic data were downloaded from the Alzheimer's Disease Neuroimaging Initiative (ADNI; http://adni.loni.usc.edu/). A total of 129 subjects were included in the study: 41 subjects who are CN, 54 subjects with MCI, and 34 subjects with AD.

Table 1 includes a summary of the participants' demographic and medical information. In the experiments, the subjects with AD presented significantly lower scores in the screening assessment cognitive test Mini-Mental State Examination (MMSE) in comparison with the other groups. The subjects were age-matched (Kruskal-Wallis test: $p > 0.8$), gender-matched (Chi-Squared test: $p > 0.1$), and matching number of years of education (Kruskal-Wallis test: $p > 0.2$). As expected, MMSE scores had a significant difference between all pairs of groups (Kruskal-Wallis test: $p < 10^{-14}$).

2.2. Image Acquisition

The acquisition of fMRI images was performed using Philips Medical Systems scanner. The fMRI images were obtained using an echo planar imaging sequence at a field strength of 3.0 Tesla, with a repetition time (TR) of 3 seconds, an echo time (TE) of 30 milliseconds, and a flip angle of 80 degrees. The matrix size was 64×64 pixels, 140 volumes, 48 slices per volume, slice thickness of 3.3 mm, and voxel size of $3.3 \times 3.3 \times 3.3$ mm^3.

2.3. fMRI Preprocessing

The fMRI pre-processing steps were carried out using the CONN toolbox version 21a, which utilizes the Statistical Parametric Mapping (SPM12), both of which are MATLAB-based cross-platform software (Nieto-Castanon, 2021, Friston et al., 1994). We used the default pre-processing pipeline in CONN, consisting of the following steps in order: functional realignment and unwarp (subject motion estimation and correction), functional centering to (0,0,0) coordinates (translation), slice-time correction with interleaved slice order, outlier identification using Artifact Detection and Removal Tool, segmentation into gray matter, white matter and cerebrospinal fluid tissue, and direct normalization into standard Montreal Neurological Institute (MNI) brain space, and lastly, smoothing using spatial convolution with a Gaussian kernel of 8mm full-width half maximum. This pipeline was followed by detrending and bandpass filtering (0.001-0.1 Hz) to remove low-frequency scanner drift and physiological noise in the fMRI images. The first four time points have been filtered out to remove any artifacts.

For the extraction of Regions-Of-Interest (ROIs), the automated anatomical labeling (AAL) atlas was utilized on the pre-processed rs-fMRI dataset (Tzourio-Mazoyer et al., 2002). The list of all regions in the AAL atlas is provided in Appendix Appendix A along with their abbreviated, short, and full region names. This parcellation method has been demonstrated to be optimal for studying the FC between brain regions (Arslan et al., 2018). The voxels within each ROI were averaged, resulting in a time series for each ROI.

2.4. Functional Connectivity Analyses

2.4.1. Associative FC

We employed Sparse Partial Correlation to obtain AFC from fMRI time series data (Banerjee et al., 2008, Pervaiz et al., 2020, Schmittmann et al., 2015). The sparse partial correlation regularizes partial correlation with a graphical LASSO penalty, whose regularization parameters were selected by cross-validation (Meinshausen et al., 2006).

2.4.2. Causal FC

In our study, CFC is inferred using the Time-aware PC (TPC) Algorithm, a recent method to infer CFC from brain signals (Biswas and Shlizerman, 2022b, Biswas and Mukherjee, 2024). By implementing the Directed Markov Property in a time series setting, the TPC algorithm models causal interactions as an unrolled Directed Acyclic Graph (DAG), representing the time series' spatial and temporal interactions. In this unrolled DAG, nodes represent regions of interest at specific times, and edges indicate directional causal influences between these nodes. The TPC algorithm transforms time series data into sequential variables with a maximum time delay ($\tau = 1$), and applies the Peter-Clark (PC) algorithm for DAG estimation, later rolling the DAG back to form a comprehensive CFC graph. This method allows for the analysis of complex, multivariate interactions and supports whole-brain CFC estimation, accommodating the BOLD signal's temporal delays and feedback loops (Biswas and Sripada, 2023). The Python package *Time-Aware PC* is used for implementing this approach. Our methodology is

particularly suited to fMRI data, accounting for the relatively slow temporal resolution of the BOLD signal and enabling the detection of both sequential and contemporaneous interactions within the brain network (Biswas and Sripada, 2023). The outcome of this approach offers a non-parametric, detailed mapping of causal functional connectivity across the brain, enhancing our understanding of the directional flow of information and its disruption in AD.

2.5. Alterations of CFC edges in Alzheimer's disease

NBS is a popular non-parametric statistical method for identifying inter-group differences in large brain networks while controlling for multiple comparisons (Zalesky et al., 2010, Zhan et al., 2016, Olivito et al., 2017, De Schipper et al., 2018, Zhan et al., 2019, Zhu et al., 2023). It is a method for controlling the FWER when a large number of univariate tests are conducted over the connections in the network. NBS method outputs the connected components in the brain network that exhibit inter-group difference and a corresponding FWER-corrected p-value for each such component. The FWER-corrected p-values are calculated for each component using permutation testing. We used 10,000 random permutations for the permutation tests. For implementation, we used the *NBS Matlab Toolbox* and its extension for directed networks (https://www.nitrc.org/projects/nbs/) (Zalesky et al., 2010). In the implementation of NBS, we used a primary threshold of 1.8 and FWER corrected significance level of 0.05. At that significance level, the critical threshold of cluster size was 6. For finding altered sub-networks between CN and AD groups, NBS considers the direction of each connection in the whole CFC among subjects in CN and AD groups. We outlined detailed steps of implementing NBS in Appendix B.

We applied NBS to the subject-specific CFCs estimated by TPC, to identify connected components of the CFC that are altered in strength for subjects with AD compared to the CN.

3. Results

3.1. Subject-specific CFC

Figure 1 shows the AFC obtained by sparse partial correlation and CFC obtained by TPC for an example subject who is CN. The AFC (top row) revealed an extensive associative network, as evidenced by dense symmetric adjacency matrix and dense undirected graphical representations. It is expected that the CFC will be a directed subgraph of the AFC and be consistent with the overall patterns present in the AFC. The overall patterns present in the CFC obtained by TPC indeed match with the AFC as visualized in their adjacency matrices (See Figure 1-left column). On a detailed level, there are differences between TPC-CFC and AFC: TPC results in a directed graph thereby having an asymmetric adjacency matrix while AFC is an undirected graph with symmetric adjacency matrix. Furthermore, in contrast to the dense adjacency matrix and graphical representations of the AFC, the CFC obtained by TPC results in a sparse adjacency matrix and sparse graphical representation devoid of noise since the connections are filtered by conditional dependence tests in the TPC algorithm.

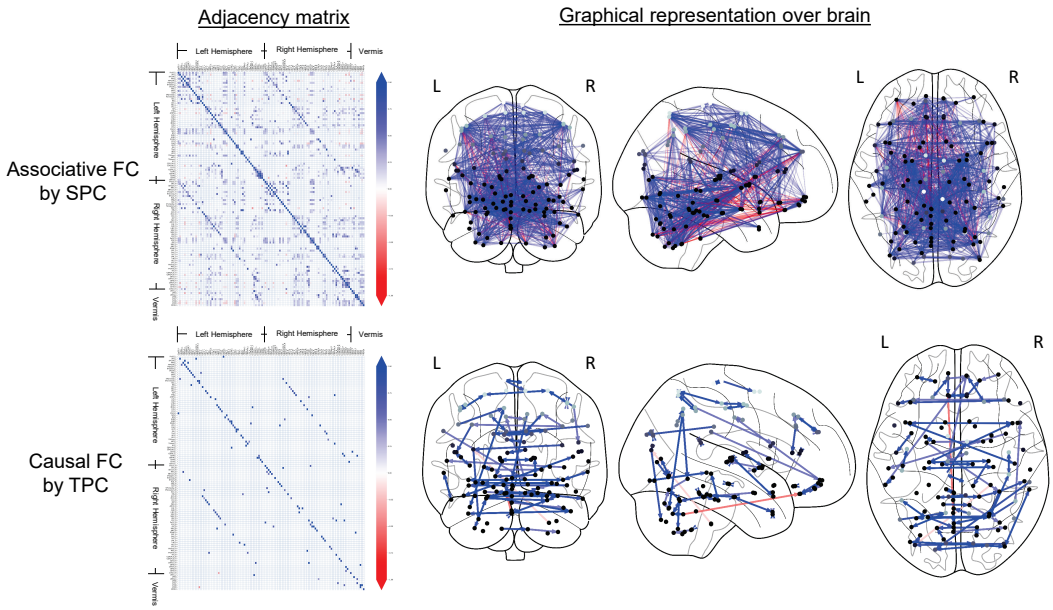

Figure 1: CFC for an example subject who is CN, estimated by TPC algorithm. The estimated CFC is represented by its adjacency matrix (left column), whose non-zero entry (i, j) represents the connection of region $i \to j$. The CFC is also visualized with directed graph edges on the Frontal, Axial and Lateral brain maps (2nd to 4th columns from left). The nodes correspond to brain region centers, ranging from superficial (light gray) to deeper (darker gray) regions, in the AAL brain atlas. The brain regions are annotated by Left (L) and Right (R) hemispheres of the brain and Vermis (V).

3.2. Altered CFC sub-networks in AD

Figure 2 shows the significantly altered ($P < 0.05$; NBS corrected) CFC sub-networks in AD compared to CN. In the implementation of NBS, t-statistic thresholds of 0.1 to 3.1 in steps of 0.1 were tried out. Among those, thresholds of 1.8, 2.2 and 2.3 yielded altered sub-networks at 5% level of significance. At each of these thresholds, a sparse thresholded adjacency matrix is obtained (Figure 2-left row). A subset of the thresholded edges is outputted by NBS as an altered sub-network at 5% level of significance. Figure 2 shows the altered sub-network's adjacency matrix representation (2nd column from left) as well as graphical representation (3rd to 5th columns from left) on brain schematics.

3.3. Brain regions in the altered CFC sub-networks

In Table 2, we report the CFC edges that are present in the altered CFC sub-networks with greater average strength in CN subjects compared to AD at NBS corrected 5% level of significance, and their source brain regions. The corresponding brain regions are in agreement with published medical literature cited in Table 2. No significantly altered sub-networks were obtained by NBS for lower average strength in CN subjects compared to AD at 5% level of significance.

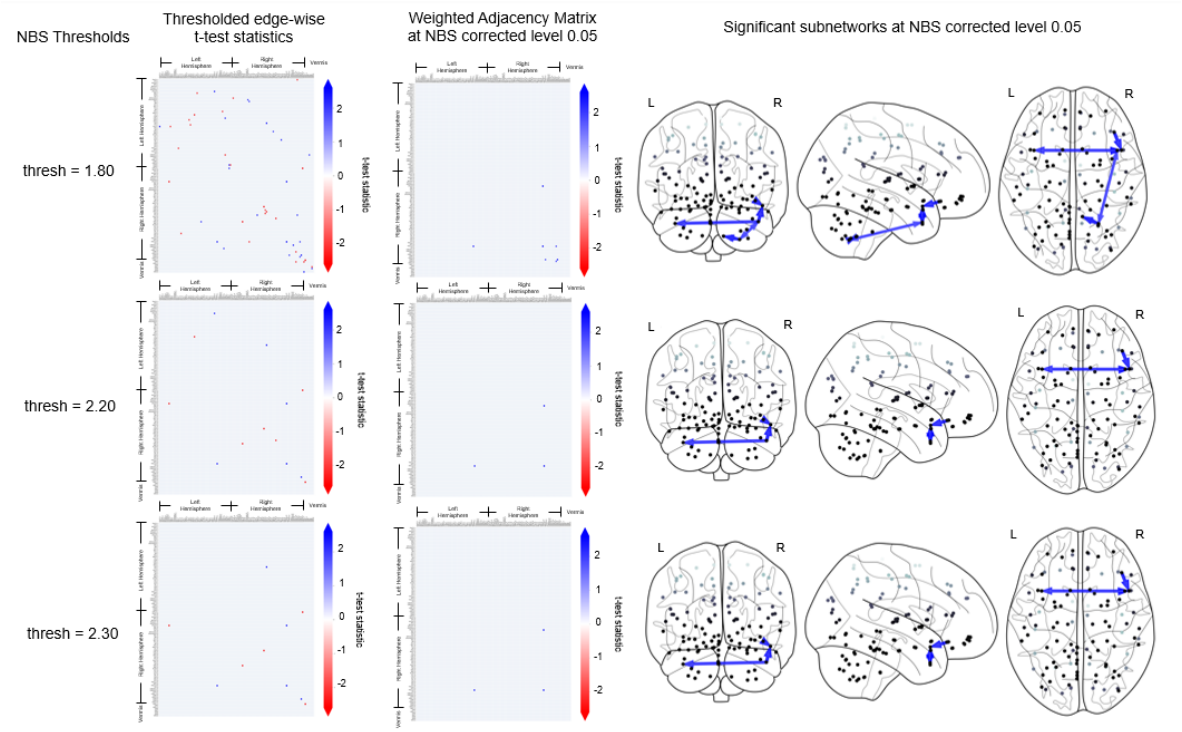

Figure 2: This figures shows the significantly altered sub-networks obtained by NBS implemented on the CFC outcome of TPC. For NBS test-statistics thresholds of 1.8, 2.2 and 2.3 (top to bottom), the thresholded edge-wise t-test statistics (left column) and the result of the NBS corrected sub-network are shown: weighted adjacency matrix representation (2nd column from left), graphical representation on brain schematics on the Frontal, Axial and Lateral brain maps (3rd to 5th columns from left).

4. Discussion

In this study, we obtained CFC sub-networks of the whole brain that are altered in AD compared to CN from resting state fMRI time series. We used the recently developed TPC algorithm based on directed graphical modeling in time series, to compute the CFC. We first computed the subject-specific CFC using TPC and compared it with AFC obtained by sparse partial correlation. We then used the CFC outcomes of TPC for further investigation into altered CFC sub-networks in AD. In this regard, we used NBS to obtain CFC sub-networks that alter in AD compared to CN subjects at 5% level of significance, while correcting for multiple comparisons in those sub-networks. The brain regions in those altered sub-networks were found to be in agreement with medical literature for regions impacted by AD.

In Figure 2 and Table 2, the presence of CFC sub-network with weight of edges in AD greater than that in CN is consistent with published studies in the literature. The absence of CFC sub-networks with weight of edges in AD lower than that in CN at 5% level of significance, supports the hypothesis that AD is a "disconnection syndrome" (Geschwind, 1965, Bozzali et al., 2011, Brier et al., 2014a).

In Table 2, the brain regions of Inferior frontal gyrus (Inferior frontal gyrus, pars

(a) CFC sub-network for CN > AD at thresh = 1.80 having p-value = 0.0299

Edge	**Region Name**	**Reported by**
IFGorb_R → TPOsup_R	Inferior frontal gyrus	Eliasova et al. (2014), Cajanus et al. (2019)
TPOmid_R → TPOsup_R	Temporal pole: middle temporal gyrus	Harasty et al. (1999), Convit et al. (2000)
TPOmid_R → TPOmid_L	Temporal pole: middle temporal gyrus	Harasty et al. (1999), Convit et al. (2000)
CER8_R → TPOmid_R	Cerebellum	Joachim et al. (1989), Jacobs et al. (2018)
TPOmid_R → CER8_R	Temporal pole: middle temporal gyrus	Harasty et al. (1999), Convit et al. (2000)
CER9_R → CER8_R	Cerebellum	Joachim et al. (1989), Jacobs et al. (2018)
CER8_R → CER9_R	Cerebellum	Joachim et al. (1989), Jacobs et al. (2018)

(b) CFC sub-network for CN > AD at thresh = 2.20 having p-value = 0.0465

Edge	**Region Name**	**Reported by**
IFGorb_R → TPOsup_R	Inferior frontal gyrus	Eliasova et al. (2014), Cajanus et al. (2019)
TPOmid_R → TPOsup_R	Temporal pole: middle temporal gyrus	Harasty et al. (1999), Convit et al. (2000)
TPOmid_R → TPOmid_L	Temporal pole: middle temporal gyrus	Harasty et al. (1999), Convit et al. (2000)

(c) CFC sub-network for CN > AD at thresh = 2.30 having p-value = 0.0233

Edge	**Region Name**	**Reported by**
IFGorb_R → TPOsup_R	Inferior frontal gyrus	Eliasova et al. (2014), Cajanus et al. (2019)
TPOmid_R → TPOsup_R	Temporal pole: middle temporal gyrus	Harasty et al. (1999), Convit et al. (2000)
TPOmid_R → TPOmid_L	Temporal pole: middle temporal gyrus	Harasty et al. (1999), Convit et al. (2000)

Table 2: CFC edges in significantly altered CFC sub-networks with greater weight in CN compared to AD group at NBS corrected 5% level of significance. The sub-networks are obtained at different NBS's t-statistic thresholds: (a) 1.80 , (b) 2.20 and (c) 2.30 yielding altered sub-networks with p-values of 0.0299, 0.0465 and 0.0233 respectively. The corresponding source brain regions are in agreement with regions reported in literature (right column) as impacted by AD.

orbitalis Right → Temporal pole: superior temporal gyrus Right) and Temporal pole: middle temporal gyrus (Temporal pole: middle temporal gyrus Right → Temporal pole: superior temporal gyrus Right, Temporal pole: middle temporal gyrus Right → Temporal pole: middle temporal gyrus Left) are prominent in sub-networks with lower CFC weight in AD compared to CN subjects. Among these regions, the Inferior frontal gyrus plays a role in attentional control (Hampshire et al., 2010), and language processing (Rota et al., 2009), and is known to be impacted by AD (Eliasova et al., 2014, Cajanus et al., 2019). The temporal pole: middle temporal gyrus, another region in Table 2, is considered to be important in word memory and meaning (Luria, 1976, Benson, 1980, Ojemann and Mateer, 1979, Damasio and Damasio, 1980, Frackowiak et al., 1981, Mesulam, 1990, Demonet et al., 1993, Gazzaniga, 1993). It has been reported to be impacted by AD (Harasty et al., 1999, Convit et al., 2000). Also present in Table 2(a) is the Cerebellum (Lobule VIII of cerebellar hemisphere Right → Temporal pole: middle temporal gyrus Right, Lobule IX of cerebellar hemisphere Right → Lobule VIII of cerebellar hemisphere Right, Lobule VIII of cerebellar hemisphere Right → Lobule IX of cerebellar hemisphere Right), which plays an essential role in motor control and perception (Paulin, 1993, Akshoomoff and Courchesne, 1992, Botez et al., 1989), and has been reported to be impacted by AD (Joachim et al., 1989, Jacobs et al., 2018).

4.1. Conclusion

In this paper, we have demonstrated the following: (a) Application of the TPC algorithm to compute whole-brain CFC for each subject (b) Comparison of CFC with AFC computed using sparse partial correlation (c) Identified CFC sub-networks with altered connectivity in subjects with AD compared to CN using NBS at 5% level of significance. (d) Interpretation of altered CFC sub-networks in the context of AD using domain (neuropathological) knowledge. The findings are consistent with published medical literature. The results demonstrate the potential of computing the whole-brain CFC from fMRI data using the TPC algorithm to gain prognostic and diagnostic insights.

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Appendix A. Automated Anatomical Labeling (AAL) Atlas

The regions in the AAL atlas along with their abbreviated, short and full names are listed in Table A.3.

Table A.3: Names of regions in the AAL Atlas

No	Abbr. Name	Short Name	Full Region Name
1	PreCG_L	Precentral_L	Precentral gyrus Left
2	PreCG_R	Precentral_R	Precentral gyrus Right
3	SFG_L	Frontal_Sup_L	Superior frontal gyrus, dorsolateral Left
4	SFG_R	Frontal_Sup_R	Superior frontal gyrus, dorsolateral Right
5	SFGorb_L	Frontal_Sup_Orb_L	Superior frontal gyrus, pars orbitalis Left
6	SFGorb_R	Frontal_Sup_Orb_R	Superior frontal gyrus, pars orbitalis Right
7	MFG_L	Frontal_Mid_L	Middle frontal gyrus Left
8	MFG_R	Frontal_Mid_R	Middle frontal gyrus Right
9	MFGorb_L	Frontal_Mid_Orb_L	Middle frontal gyrus, pars orbitalis Left
10	MFGorb_R	Frontal_Mid_Orb_R	Middle frontal gyrus, pars orbitalis Right
11	IFGoperc_L	Frontal_Inf_Oper_L	Inferior frontal gyrus, opercular part Left
12	IFGoperc_R	Frontal_Inf_Oper_R	Inferior frontal gyrus, opercular part Right
13	IFGtriang_L	Frontal_Inf_Tri_L	Inferior frontal gyrus, triangular part Left
14	IFGtriang_R	Frontal_Inf_Tri_R	Inferior frontal gyrus, triangular part Right
15	IFGorb_L	Frontal_Inf_Orb_L	Inferior frontal gyrus, pars orbitalis, Left
16	IFGorb_R	Frontal_Inf_Orb_R	Inferior frontal gyrus, pars orbitalis, Right
17	ROL_L	Rolandic_Oper_L	Rolandic operculum Left
18	ROL_R	Rolandic_Oper_R	Rolandic operculum Right
19	SMA_L	Supp_Motor_Area_L	Supplementary motor area Left
20	SMA_R	Supp_Motor_Area_R	Supplementary motor area Right
21	OLF_L	Olfactory_L	Olfactory cortex Left
22	OLF_R	Olfactory_R	Olfactory cortex Right
23	SFGmedial_L	Frontal_Sup_Medial_L	Superior frontal gyrus, medial Left
24	SFGmedial_R	Frontal_Sup_Medial_R	Superior frontal gyrus, medial Right
25	SFGmedorb_L	Frontal_Med_Orb_L	Superior frontal gyrus, medial orbital Left
26	SFGmedorb_R	Frontal_Med_Orb_R	Superior frontal gyrus, medial orbital Right
27	REC_L	Rectus_L	Gyrus rectus Left
28	REC_R	Rectus_R	Gyrus rectus Right
29	INS_L	Insula_L	Insula Left
30	INS_R	Insula_R	Insula Right
31	ACC_L	Cingulum_Ant_L	Anterior cingulate & paracingulate gyri Left
32	ACC_R	Cingulum_Ant_R	Anterior cingulate & paracingulate gyri Right
33	MCC_L	Cingulum_Mid_L	Middle cingulate & paracingulate gyri Left
34	MCC_R	Cingulum_Mid_R	Middle cingulate & paracingulate gyri Right
35	PCC_L	Cingulum_Post_L	Posterior cingulate gyrus Left

36	PCC_R	Cingulum_Post_R	Posterior cingulate gyrus Right
37	HIP_L	Hippocampus_L	Hippocampus Left
38	HIP_R	Hippocampus_R	Hippocampus Right
39	PHG_L	ParaHippocampal_L	Parahippocampal gyrus Left
40	PHG_R	ParaHippocampal_R	Parahippocampal gyrus Right
41	AMYG_L	Amygdala_L	Amygdala Left
42	AMYG_R	Amygdala_R	Amygdala Right
43	CAL_L	Calcarine_L	Calcarine fissure and surrounding cortex Left
44	CAL_R	Calcarine_R	Calcarine fissure and surrounding cortex Right
45	CUN_L	Cuneus_L	Cuneus Left
46	CUN_R	Cuneus_R	Cuneus Right
47	LING_L	Lingual_L	Lingual gyrus Left
48	LING_R	Lingual_R	Lingual gyrus Right
49	SOG_L	Occipital_Sup_L	Superior occipital gyrus Left
50	SOG_R	Occipital_Sup_R	Superior occipital gyrus Right
51	MOG_L	Occipital_Mid_L	Middle occipital gyrus Left
52	MOG_R	Occipital_Mid_R	Middle occipital gyrus Right
53	IOG_L	Occipital_Inf_L	Inferior occipital gyrus Left
54	IOG_R	Occipital_Inf_R	Inferior occipital gyrus Right
55	FFG_L	Fusiform_L	Fusiform gyrus Left
56	FFG_R	Fusiform_R	Fusiform gyrus Right
57	PoCG_L	Postcentral_L	Postcentral gyrus Left
58	PoCG_R	Postcentral_R	Postcentral gyrus Right
59	SPG_L	Parietal_Sup_L	Superior parietal gyrus Left
60	SPG_R	Parietal_Sup_R	Superior parietal gyrus Right
61	IPG_L	Parietal_Inf_L	Inferior parietal gyrus, excluding supramargina...
62	IPG_R	Parietal_Inf_R	Inferior parietal gyrus, excluding supramargina...
63	SMG_L	SupraMarginal_L	SupraMarginal gyrus Left
64	SMG_R	SupraMarginal_R	SupraMarginal gyrus Right
65	ANG_L	Angular_L	Angular gyrus Left
66	ANG_R	Angular_R	Angular gyrus Right
67	PCUN_L	Precuneus_L	Precuneus Left
68	PCUN_R	Precuneus_R	Precuneus Right
69	PCL_L	Paracentral_Lobule_L	Paracentral lobule Left
70	PCL_R	Paracentral_Lobule_R	Paracentral lobule Right
71	CAU_L	Caudate_L	Caudate nucleus Left
72	CAU_R	Caudate_R	Caudate nucleus Right
73	PUT_L	Putamen_L	Lenticular nucleus, Putamen Left
74	PUT_R	Putamen_R	Lenticular nucleus, Putamen Right
75	PAL_L	Pallidum_L	Lenticular nucleus, Pallidum Left
76	PAL_R	Pallidum_R	Lenticular nucleus, Pallidum Right
77	THA_L	Thalamus_L	Thalamus Left
78	THA_R	Thalamus_R	Thalamus Right

79	HES_L	Heschl_L	Heschl's gyrus Left
80	HES_R	Heschl_R	Heschl's gyrus Right
81	STG_L	Temporal_Sup_L	Superior temporal gyrus Left
82	STG_R	Temporal_Sup_R	Superior temporal gyrus Right
83	TPOsup_L	Temporal_Pole_Sup_L	Temporal pole: superior temporal gyrus Left
84	TPOsup_R	Temporal_Pole_Sup_R	Temporal pole: superior temporal gyrus Right
85	MTG_L	Temporal_Mid_L	Middle temporal gyrus Left
86	MTG_R	Temporal_Mid_R	Middle temporal gyrus Right
87	TPOmid_L	Temporal_Pole_Mid_L	Temporal pole: middle temporal gyrus Left
88	TPOmid_R	Temporal_Pole_Mid_R	Temporal pole: middle temporal gyrus Right
89	ITG_L	Temporal_Inf_L	Inferior temporal gyrus Left
90	ITG_R	Temporal_Inf_R	Inferior temporal gyrus Right
91	CERCRU1_L	Cerebellum_Crus1_L	Crus I of cerebellar hemisphere Left
92	CERCRU1_R	Cerebellum_Crus1_R	Crus I of cerebellar hemisphere Right
93	CERCRU2_L	Cerebellum_Crus2_L	Crus II of cerebellar hemisphere Left
94	CERCRU2_R	Cerebellum_Crus2_R	Crus II of cerebellar hemisphere Right
95	CER3_L	Cerebellum_3_L	Lobule III of cerebellar hemisphere Left
96	CER3_R	Cerebellum_3_R	Lobule III of cerebellar hemisphere Right
97	CER4_5_L	Cerebellum_4_5_L	Lobule IV, V of cerebellar hemisphere Left
98	CER4_5_R	Cerebellum_4_5_R	Lobule IV, V of cerebellar hemisphere Right
99	CER6_L	Cerebellum_6_L	Lobule VI of cerebellar hemisphere Left
100	CER6_R	Cerebellum_6_R	Lobule VI of cerebellar hemisphere Right
101	CER7b_L	Cerebellum_7b_L	Lobule VIIB of cerebellar hemisphere Left
102	CER7b_R	Cerebellum_7b_R	Lobule VIIB of cerebellar hemisphere Right
103	CER8_L	Cerebellum_8_L	Lobule VIII of cerebellar hemisphere Left
104	CER8_R	Cerebellum_8_R	Lobule VIII of cerebellar hemisphere Right
105	CER9_L	Cerebellum_9_L	Lobule IX of cerebellar hemisphere Left
106	CER9_R	Cerebellum_9_R	Lobule IX of cerebellar hemisphere Right
107	CER10_L	Cerebellum_10_L	Lobule X of cerebellar hemisphere Left
108	CER10_R	Cerebellum_10_R	Lobule X of cerebellar hemisphere Right
109	VER1_2	Vermis_1_2	Lobule I, II of vermis
110	VER3	Vermis_3	Lobule III of vermis
111	VER4_5	Vermis_4_5	Lobule IV, V of vermis
112	VER6	Vermis_6	Lobule VI of vermis
113	VER7	Vermis_7	Lobule VII of vermis
114	VER8	Vermis_8	Lobule VIII of vermis
115	VER9	Vermis_9	Lobule IX of vermis
116	VER10	Vermis_10	Lobule X of vermis

Appendix B. The Network-Based Statistics (NBS) method

For comparing the CFCs between two groups, NBS initially calculates a Generalized Linear Model based statistic to score each connection. These scores are then subjected to a primary threshold, leading to the selection of suprathreshold links with scores surpassing this threshold. The thresholding is done separately for positive and negative scores, identifying connected components where individuals with CN exhibit notably higher or lower connectivity strength than those with AD. Various primary thresholds were evaluated in the range of 0.1 to 3 in steps of 0.1, and fixed at 1.8 which is the lowest in the range yielding significance at family wise error rate (FWER) level 0.05. Although the primary threshold impacts method sensitivity, yet FWER control is ensured regardless of the chosen threshold (Zalesky et al., 2010). Subsequently, components within the suprathreshold link set are identified, and their sizes are recorded. To establish component significance, a non-parametric permutation test is conducted. In each permutation, subjects are randomly reassigned to CN and AD groups. Independent scores are computed for each link, and the size of the largest connected component among suprathreshold links is noted. This process is reiterated 10,000 times to generate a null distribution of the pseudo size of the connected component. Ultimately, the corrected p-value for a connected component of size M is determined based on the proportion of permutations where the largest connected component size exceeds M. The NBS method is implemented in the *NBS Matlab Toolbox* and its extension for directed networks (https://www.nitrc.org/projects/nbs/) (Zalesky et al., 2010).