

The interplay between insomnia and Alzheimer's disease on the triple brain network

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INTRODUCTION

Insomnia is a common sleep disorder characterized by perceived difficulties in falling asleep, maintaining sleep, and early waking times [3]. A majority of insomnia symptoms are related to aberrant connectivity within and between the default mode network (DMN), salience network (SN) and central executive network (CEN) [3,6]. These three brain networks, primarily involved in working memory, self-referential processing, and switching between mind-wandering and directed thought, comprise the triple brain network [7].

Insomnia is a significant risk factor for Alzheimer's disease (AD) in the prodromal stage [2], while also being a common comorbidity in the clinical stage [10]. Moreover, much like insomnia, AD is characterized by triple network deficiencies [8]. As such, we hypothesize that there may be an interaction between the effects of insomnia and Alzheimer's disease pathology within the triple brain network system.

A1

S

A3

AD

0.9

METHODS

Participants. 320 ADNI [1] subjects: N=178 cognitively normal (CN), N=132 Mild Cognitive Impairment (MCI), N=40 Alzheimer's disease (AD).

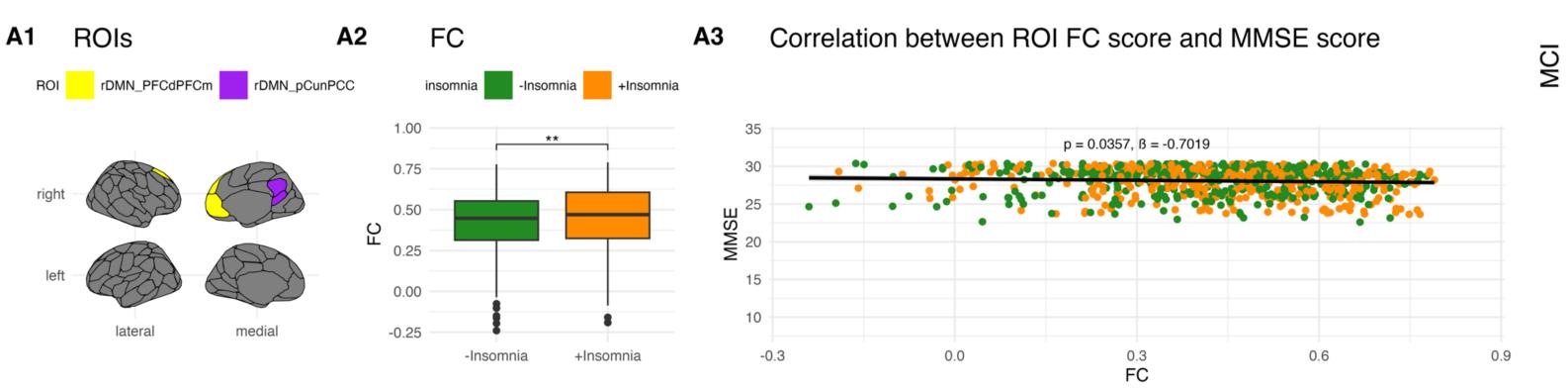
Clinical Variables. 50% of all subjects reported insomnia symptoms on the NPI [4]. All subjects had performed a MMSE. N=58 subjects had CSF biomarker (beta-amyloid, ptau) data available.

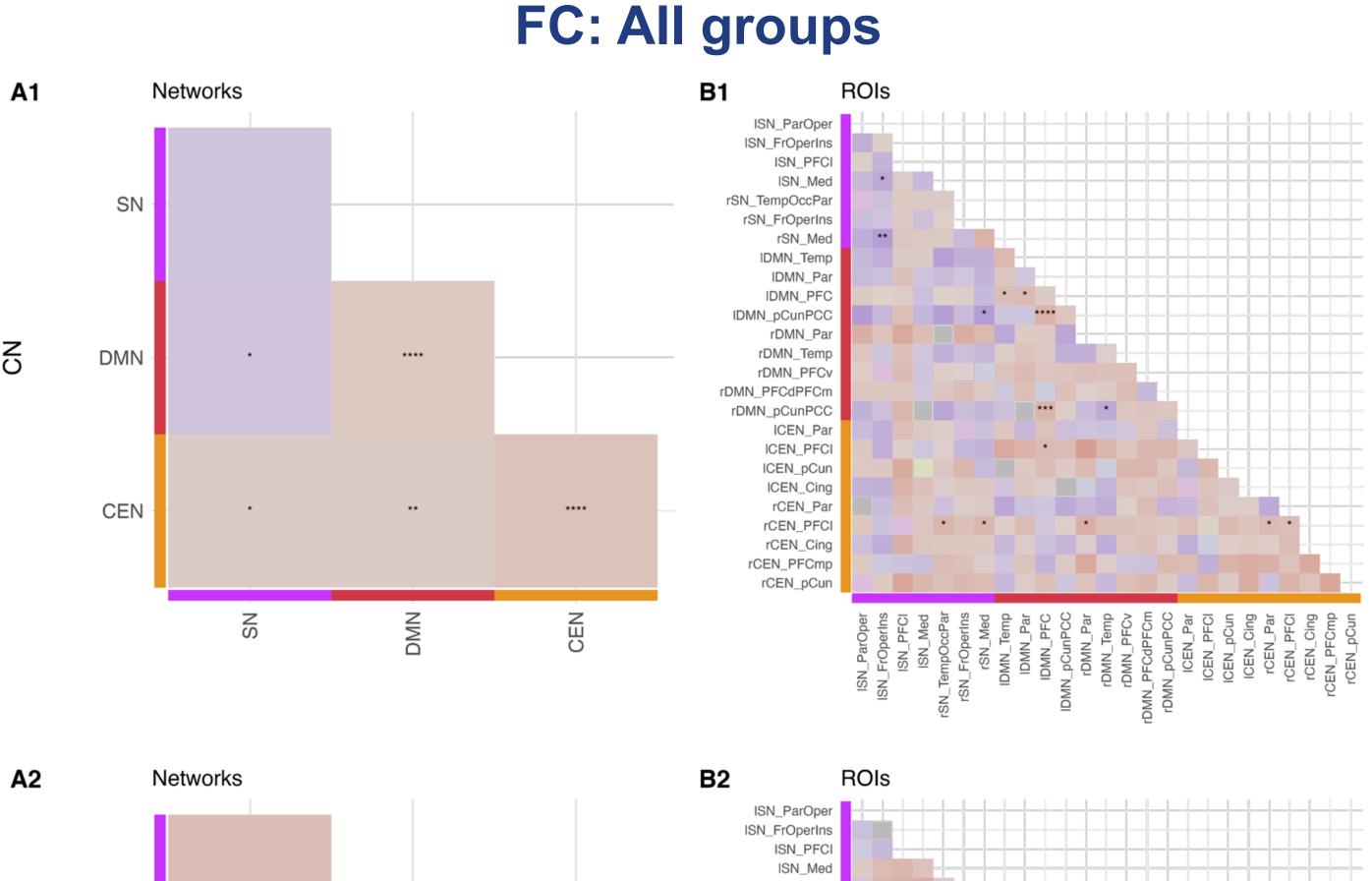
Preprocessing. T1 MRI and resting state (rs-)fMRI were preprocessed using fMRIprep [5]. Triple network nodes were identified with the Schaefer atlas [9]

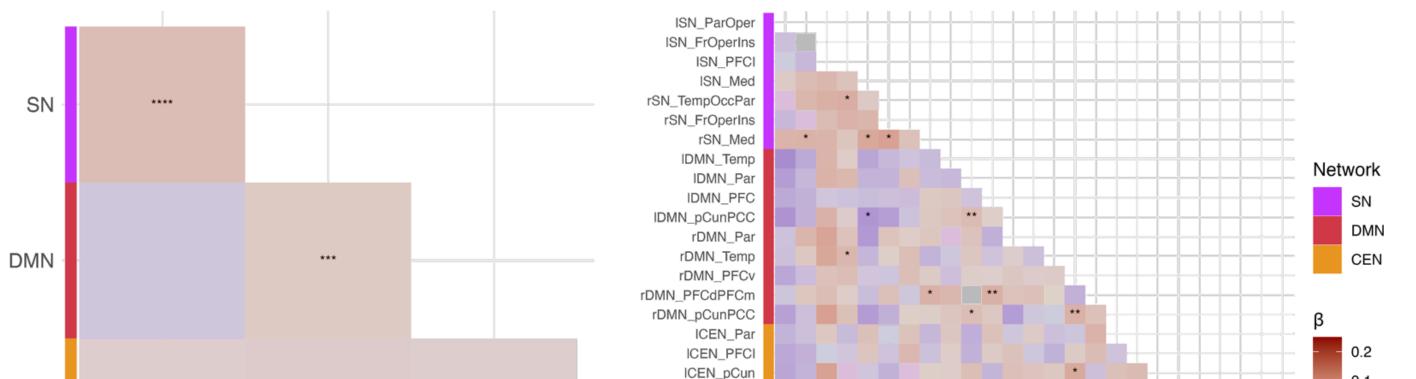
MRI indices. The triple network gray matter volume (GMV) and structural covariance were obtained to assess (inter-)nodal morphological changes. The degrees (SC) centrality (DC) and functional connectivity (FC) were obtained to assess (inter-)nodal functional changes.

Statistical analysis. For each index, pertaining to each (combination of) node(s), we fit a linear regression model, corrected for age and sex, assessing the effect of insomnia symptoms in different diagnostic groups. P<0.05, FDR-corrected.

MMSE: MCI







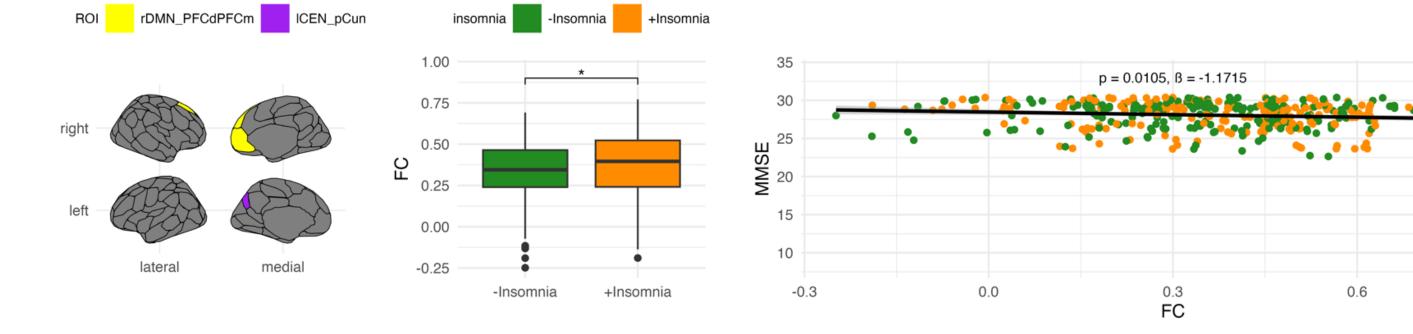
CEN SN CEN





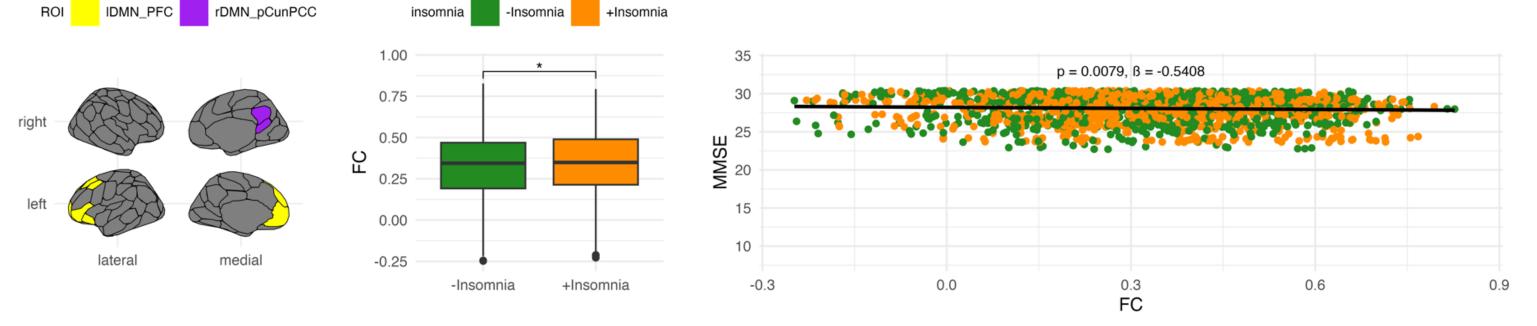
B1 ROIs **B**2 FC

Correlation between ROI FC score and MMSE score **B**3

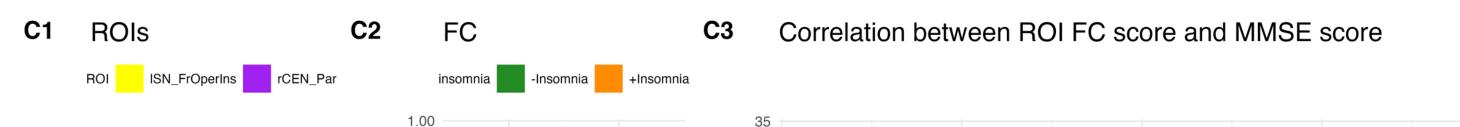


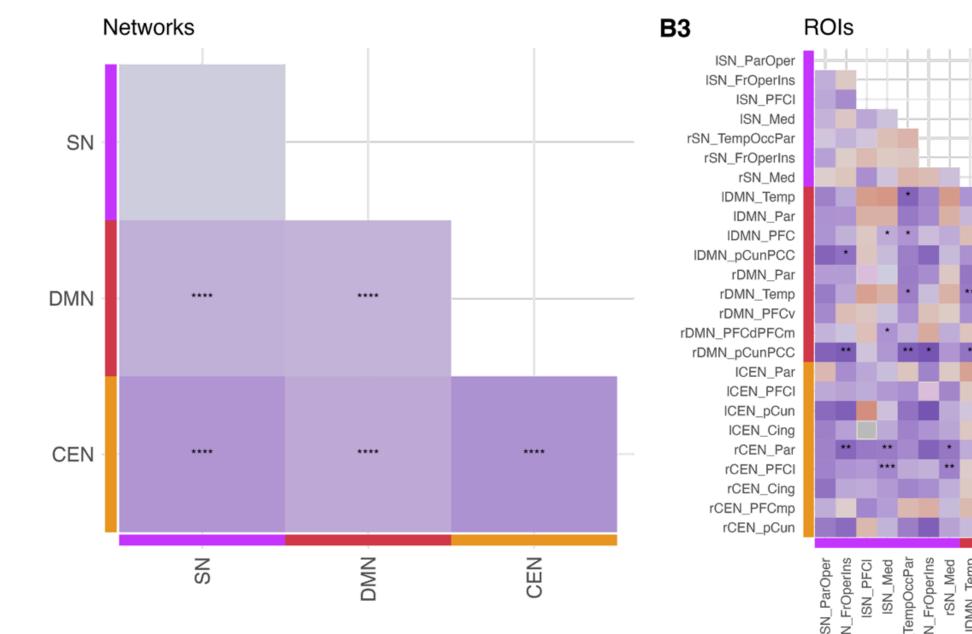
C1 ROIs

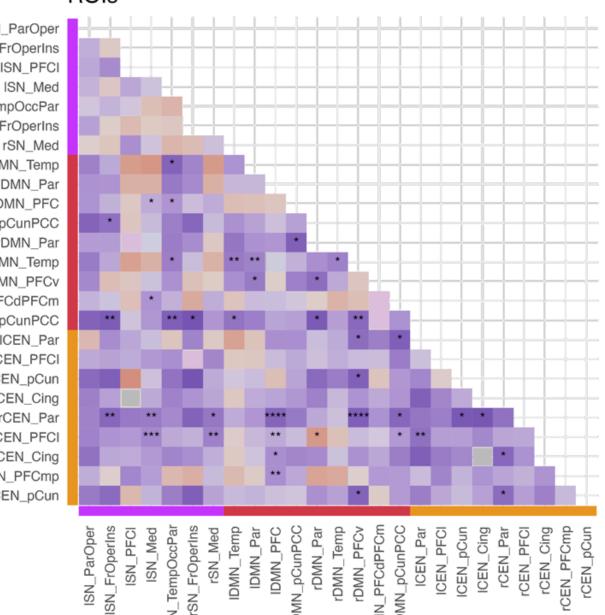




MMSE: AD

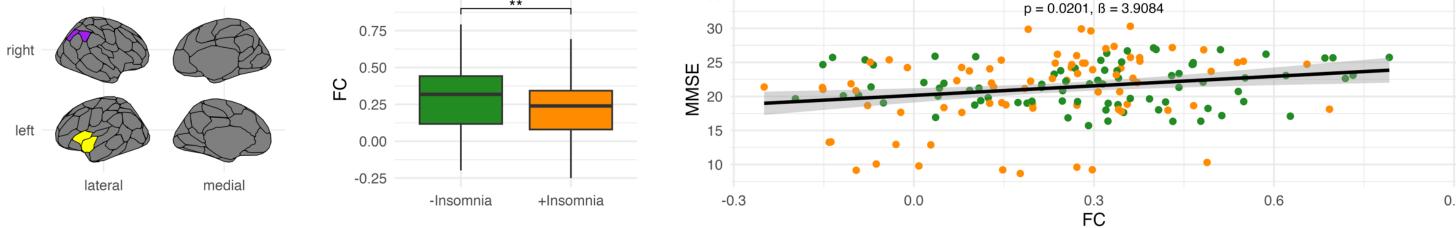






RESULTS

There were no significant changes in GMV, SC, and DC across all groups. CN and MCI individuals displayed patterns of hyperconnectivity in the DMN, SN and CEN. In



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FC

insomnia

C2

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contrast, AD individuals displayed patterns of hypoconnectivity across the DMN and CEN. Intra-DMN hyperconnectivity in CN and MCI and CEN hypoconnectivity in AD primarily associated with decreased a MMSE score. These connectivity were alterations were not associated with significant change in CSF biomarker burden.

CONCLUSION

- Insomnia symptoms modify the effect of AD on triple network FC
- Prodromal stages (CN, MCI) are associated with classical markers of hyperarousal, including intra-DMN hyperconnectivity.
- Clinical stages (AD) are associated with markers of emotional dysregulation and post-insomnia anxiety
- Comorbidities with anxiety and depression may underlie a heightened sensitivity to the affective symptoms of insomnia in AD.



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