

The effects of sleep-disordered breathing on neuroimaging biomarkers of Alzheimer's disease

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INTRODUCTION

- Unclear neurobiological mechanism of the association of SDB with AD progression [1, 2]: Higher cortical Aβ accumulation in middle-aged patients with severe SDB [3], decreased Cerebrospinal Fluid (CSF) CSF Αβ, and increased phosphorylated tau (p-tau) in healthy and MCI groups, particularly in subjects with SDB [4].
- SDB may interact with AD pathophysiology in preclinical increase the stages risk of to AD finally exacerbate developing and the development of AD-related cognitive impairments.
- biomarkers, amyloid-beta Positron Developing including Emission Tomography (A β -PET), 18F-Fluorodeoxyglucose PET (FDG-PET), and structural Magnetic Resonance Imaging (sMRI) [5], as well as A β and tau assessment in CSF, pathophysiological advanced understanding of our mechanisms, early detection, and monitoring of the progress and treatment of AD [6].

How Aβ-PET, FDG-PET, and sMRI biomarkers may help to understand the underlying neurobiological mechanisms of the association between SDB and AD progression.





Fig. 1 Cognitive status-SDB interaction medium effect sizes on AD biomarkers. The brain regions that have medium effect sizes of the interaction between cognitive status (Alzheimer's disease, mild cognitive impairment and cognitively unimpaired) and SDB condition (with and without sleep-disordered breathing) for different biomarkers: A) amyloid-plaque burden (Aβ), B) regional fluorodeoxyglucose (rFDG), C) grey matter volume (GMV).

Fig. 2 AD biomarkers alteration in different groups. Differences in each biomarker value in different states of SDB condition (with and without SDB) through different disease groups (Alzheimer's disease, mild cognitive impairment, and cognitively unimpaired).



Fig. 3 Association between cognitive status-SDB interaction and cognitive score and pattern of **biomarker's changes.** A) Interaction between cognitive status and SDB condition has a medium effect size on highlighted regions. Furthermore, the highlighted regions are significantly associated with cognitive status-SDB interaction effect size on MMSE. B) changes pattern (A β , rFDG, and GMV) through different cognitive status in the highlighted regions from A. C) MMSE score changes through different cognitive status. D) CSF Aβ42 changes through different cognitive status. E) CSF p-tau changes through different cognitive status.



in AD+SDB

Fig. 4 The neurobiological interaction between SDB and AD. SDB induces hypoxia and sleep fragmentation, which leads to neuroinflammation and glymphatic stasis. These changes, together with genetic, environmental, and lifestyle factors, exacerbate aging effects on the accumulation of amyloid-beta, and insoluble tau protein, which results in brain metabolic and structural alterations and precedes AD pathology. * Glymphatic system illustration obtained from [8].



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CONCLUSION



