



## Basal forebrain septal nuclei are enlarged in healthy subjects prior to the development of Alzheimer's disease



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### ABSTRACT

Alzheimer's disease (AD) is known to be associated with loss of cholinergic neurons in the nucleus basalis of Meynert, located in the posterior basal forebrain. Structural changes of septal nuclei, located in the anterior basal forebrain, have not been well studied in AD. Using a validated algorithm, we manually traced septal nuclei on high-resolution coronal magnetic resonance imaging (MRI) in 40 subjects with mild cognitive impairment (MCI) or AD, 89 healthy controls, and 18 subjects who were cognitively normal at the time of MRI but went on to develop AD an average of 2.8 years later. We found that cognitively normal subjects destined to develop AD in the future had enlarged septal nuclei as compared to both healthy controls and patients with current MCI or AD. To our knowledge, this is the first time a brain structure has been found to be enlarged in association with risk of AD. Further research is needed to determine if septal enlargement reflects neuroplastic compensation, amyloid deposition, inflammation, or another process and to determine whether it can serve as an early MRI biomarker of AD.

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### 1. Introduction

Based on autopsy and neuroimaging studies, Alzheimer's disease (AD) is associated with loss of cholinergic neurons in the nucleus basalis of Meynert (NBM), located in the posterior basal forebrain (BF) (Teipel et al., 2005; Whitehouse et al., 1981). NBM atrophy occurs early in the course of AD (Grothe et al., 2012) resulting in a cortical cholinergic deficit that can be partially ameliorated with cholinergic medication; this is the rationale for the use of acetylcholinesterase inhibitors to treat symptoms of AD.

The role of anterior BF structures in AD, in particular the septal region, has been less studied. The human septal region is located under and contiguous with the septum pellucidum and contains well-developed nuclei including the ventrolateral, dorsolateral, intermediolateral, septofimbrial, and medially, the vertical limb of the diagonal band of Broca (Andy and Stephan, 1968; Mai et al., 2004). Septal nuclei are strongly interconnected with hippocampi via the fimbria/fornix and are critical for generating the

hippocampal theta rhythm needed for learning and memory (Buzsaki, 2002; Gu and Yakel, 2011; Hangya et al., 2009; Huerta and Lisman, 1993; Stewart and Fox, 1990; Winson, 1978). Septal lesions impair memory in animals (Baxter et al., 2013; Winson, 1978) and humans (Alexander and Freedman, 1984; Fujii et al., 2002). Larger septal volume has been associated with better contextual memory in young healthy subjects (Butler et al., 2012). One might expect that septal atrophy, like NBM atrophy, would be an early finding in the development of AD. However, there are hints that this is not the case: autopsy studies demonstrate preserved septal cholinergic cell bodies (Mufson et al., 1989; Vogels et al., 1990) with increased cholinergic innervation of the hippocampus in early AD (DeKosky et al., 2002; Geddes et al., 1985; Hyman et al., 1987). Neuroimaging studies that divide the BF into anterior and posterior divisions show atrophy in early AD *only* in the posterior BF and NBM, with relative preservation of the anterior BF structure until the late stages of AD (Grothe et al., 2012; Kerbler et al., 2015; Kilimann et al., 2014).

Complicating assessment of the human septal region using neuroimaging is the fact that they are small ( $\sim 290 \text{ mm}^3$ ) (Butler et al., 2014) and in a location at the base of the brain, in between the ventricles that can be prone to artifacts in magnetic resonance

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imaging (MRI) studies, especially those that use automated processing. Septal nuclei are not included in any of the standard neuroanatomical parcellation schemes typically used to interpret human neuroimaging studies (Fischl et al., 2002; Tzourio-Mazoyer et al., 2002). To address this issue, we developed a manual tracing algorithm based on histology to accurately measure septal nuclei volume in vivo using MRI (Butler et al., 2014). Here, we used this tracing algorithm to measure septal volume in healthy controls, patients with current MCI or AD, and participants who were clinically normal at the time of MRI but later went on to develop AD.

## 2. Materials and methods

### 2.1. Participant selection

All subjects were recruited by the Center for Brain Health (CBH) at the NYU School of Medicine for institutional review board–approved longitudinal studies of aging, cognitive decline, and AD risk factors. A convenience sample was selected from the CBH data set to include all participants scanned on the same 1.5-T MRI, aged older than 55 years and diagnosed as MCI or AD at any time point, as well as a random sample of normal participants. The purpose of this selection strategy was to “enrich” the sample with participants with current or future MCI/AD, which constitutes a low proportion of the CBH data set. Later, we excluded MCI subjects who did not eventually decline to AD to avoid diagnostic uncertainty and heterogeneity inherent to the diagnosis of MCI. Each subject had an MRI (see acquisition protocol below).

### 2.2. Participant assessment

All participants underwent medical, psychiatric, and neurological assessments, as well as routine blood tests, electrocardiogram, and detailed neuropsychological testing that included the Mini–Mental Status Examination (Folstein et al., 1975) and Clinical Dementia Rating scale (Morris, 1993) as well as other elements of the National Alzheimer’s Coordinating Center (NACC) Uniform Data Set Neuropsychological Test Battery. A diagnosis of normal, MCI, or AD dementia was made in accord with standard criteria (Albert et al., 2011; McKhann et al., 2011) by an experienced clinician based on the diagnostic interview with the participant and his/her study partner and review of all available information including cognitive test scores and neuroimaging.

### 2.3. MRI acquisition

MRI was performed on the same quality-controlled high-resolution 1.5-T GE scanner between 1994 and 2013. The scanner did not undergo hardware or software upgrades during this time period. The examination included a T1-weighted spoiled gradient echo sequence with repetition time = 35 ms, time to echo = 2 ms, flip angle = 60°, number of excitations = 1, voxel size = 0.8 × 0.8 × 1.6 mm, field of view = 200 mm, and matrix = 256 × 192 × 124, reconstructed as 256 × 256. Although the examination also included Fluid-attenuated inversion recovery (FLAIR), T2-weighted, and diffusion-weighted sequences, only T1-weighted data were used for septal nuclei volumetric analysis.

### 2.4. MRI analysis

Tracing was performed on de-identified T1 MRIs by 5 independent tracers. After a training period during which tracers reached consensus on septal boundaries in a set of training scans, each tracer traced between 30 and 60 scans. All tracers traced a random subset of 11 scans (different than the training scans) to

evaluate interrater reliability. After tracing was completed, it was found that 1 tracer had not accurately followed the tracing rules, and another had traced scans from only 1 of the 3 diagnostic categories, so their scans were excluded from further analysis.

### 2.5. Diagnostic classification of scans

Scans were divided into 3 categories as follows: For subjects with a diagnosis of AD at any time point, a scan obtained at least 1 year before the diagnosis of MCI/AD, when the participant was cognitively normal, was selected for analysis if available. This group was deemed “future AD” (n = 18). Scans from AD patients obtained after a diagnosis of MCI or AD had been made were deemed “current MCI/AD” (n = 40). This group included 5 participants scanned at the time they were diagnosed with MCI who later declined to AD and 35 participants scanned after they had been diagnosed with AD. Normal participants had at least 1 year of follow-up and were diagnosed as normal at all time points (n = 89). For participants with more than 1 MRI scan that qualified for the aforementioned categories, the earliest scan was selected.

Tracing was performed using a validated and published semi-automated tracing algorithm implemented in FireVoxel (Mikheev et al., 2008). As described in more detail in the study by Butler et al., 2014, all scans were first reformatted to coronal planes perpendicular to a line connecting the anterior and posterior commissures. Coronal slices were traced from anterior to posterior. The *anterior septal boundary* was defined as the most anterior coronal slice in which the bilateral globus pallidus was visible and gray matter at the base of the septum pellucidum was present. The *superior extent of septal nuclei* was defined by the plane where the membranous septum pellucidum widened into septal nuclei. *Lateral boundaries* were defined by parallel sagittally oriented planes through the most inferior and medial aspect of each lateral ventricle. When a slice was anterior to the crossing fibers of the anterior commissure, the *inferior boundary* was the base of the brain. When the crossing fibers of the anterior commissure were fully visible, they served as the inferior boundary. The *posterior boundary* was defined by the following rule—A slice was considered to contain septal gray matter if it met at least 2 of the following 3 criteria: (1) the T1-weighted signal intensity indicates the presence of some gray matter (rather than pure white matter intensity); (2) presence of the crossing fibers of the anterior commissure; and (3) lack of cerebrospinal fluid (CSF) space in the center of the septal region (corresponding to CSF space between columns of the fornix). We used these boundaries and landmarks to create an overinclusive 3D region of interest (ROI). In the next step, we identified CSF voxels as having intensity less than half the average intensity of the white matter. The final step consists of automatic removal of CSF voxels from the overinclusive 3D ROI, yielding the septal ROI, whose volume in cubic millimeter was calculated and recorded.

In addition, we measured the width of the lateral ventricles, which could potentially confound septal measurement (marked ventriculomegaly may distort/stretch the septal region; Butler et al., 2013, 2014) and which also provides a measure reflecting overall brain atrophy. The ventricular width was defined as the widest distance in the left-right direction at the superior extent of the ventricles measured on a coronal slice also containing the septal gray matter.

### 2.6. Statistical analysis

General linear models were used to compare septal volume and ventricular width between groups (normal, future AD, and current MCI/AD.) Tracer was included as a covariate. The dependent variable was residualized against age and height (a proxy for head size).

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