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Brainstem volumetric integrity in preclinical and prodromal Alzheimer's disease

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Abstract

Background: Neuropathological studies have suggested the tau pathology observed in Alzheimer's disease (AD) originates in brainstem nuclei, but no studies to date have quantified brainstem volumes in clinical populations with biomarker-confirmed mild cognitive impairment (MCI) or dementia due to AD or determined the value of brainstem volumetrics in predicting dementia.

Objective: The present study examined whether MRI-based brainstem volumes differ among cognitively normal older adults and those with MCI or dementia due to AD and whether preclinical brainstem volumes predict future progression to dementia.

Methods: Alzheimer's Disease Neuroimaging Initiative participants (N = 1,629) underwent baseline MRI scanning with variable clinical follow-up (6–120 months). Region of interest and voxel-based morphometric methods assessed brainstem volume differences among cognitively normal (n = 814), MCI (n = 542), and AD (n = 273) participants, as well as subsets of CSF biomarker-confirmed MCI (n = 203) and AD (n = 160) participants.

Results: MCI and AD cases showed smaller midbrain volumes relative to cognitively normal participants when normalizing to whole brainstem volume, and showed smaller midbrain, locus coeruleus, pons, and whole brainstem volumes when normalizing to total intracranial volume. Cognitively normal individuals who later progressed to AD dementia diagnosis exhibited smaller

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DISCLOSURE STATEMENT

The authors have no conflicts of interest to report.

SUPPLEMENTARY MATERIAL

Supplementary material is available in the electronic version of this article.

baseline midbrain volumes than individuals who did not develop dementia, and voxel-wise analyses revealed specific volumetric reduction of the locus coeruleus.

Conclusion: Findings are consistent with neuropathological observations of early AD-related pathology in brainstem nuclei and further suggest the clinical relevance of brainstem substructural volumes in preclinical and prodromal AD.

Keywords

Alzheimer's disease; Biomarkers; Brainstem; Cognitive aging; Locus coeruleus; Mild cognitive impairment; Neuroimaging; Magnetic resonance imaging

INTRODUCTION

Neuropathological studies have suggested tau protein-related Alzheimer's disease (AD) pathophysiological processes begin in midbrain and pontine nuclei and precede any observable cortical changes [1,2]. The classic Braak staging of AD pathology was subsequently updated to include precortical stages whereby neurofibrillary tangles first appear in brainstem nuclei and later spread to transentorhinal, hippocampal, and neocortical regions in a stereotypical fashion [3,4]. However, there has been recent debate regarding whether brainstem nuclei represent the actual origin sites of tau seeding activity or simply the earliest regions showing phospho-tau signal [5–8]. Thus, identification of an origin site for tau seeding in AD remains controversial. Despite strong evidence from postmortem autopsy studies, it remains unclear whether corresponding pathological abnormalities may be detected with *in vivo* brain MRI and whether observable brainstem pathology is clinically relevant for cognitive impairment and dementia.

A growing number of studies have identified progressive accumulation of neurofibrillary tangle pathology in midbrain (e.g., raphe nuclei, substantia nigra) and pontine (e.g., locus coeruleus, pedunculopontine nucleus) nuclei with increasing Braak stage, implicating the disruption of ascending neurotransmitter systems in the manifestation of atypical AD symptoms such as sleep-wake dysregulation, attentional/dysexecutive deficits, and neuropsychiatric abnormalities [9–11]. These histopathological approaches are supported by *in vivo* neuroimaging studies observing reduced midbrain and pontine volumes in clinically-diagnosed AD compared to cognitively normal older adults [12–16]. A shape analysis of the brainstem in AD patients and normal controls demonstrated deformation of a dorsal rostral brainstem region, and a recent voxel-wise study of the brainstem in AD and controls similarly showed differences in the dorsal rostral brainstem [12,16]. However, these studies were limited by relatively small sample sizes and a lack of sub-regional analyses. Furthermore, brainstem volumetric differences remain unexamined in biomarker-confirmed AD populations, the prodromal mild cognitive impairment (MCI) stage of disease, or the asymptomatic preclinical stage in cognitively normal individuals who eventually develop AD dementia. The present study aimed to address the dearth of knowledge regarding *in vivo* brainstem imaging in AD by quantifying brainstem subregions in a large, longitudinal study of MCI and AD dementia patients, conducting a sub-study in biomarker-confirmed AD cases, and examining the potential utility of brainstem volumetrics in predicting development of AD dementia in initially asymptomatic individuals.

MATERIALS AND METHODS

Study Participants

Participant data were drawn from the Alzheimer's Disease Neuroimaging Initiative (ADNI) database (adni.loni.usc.edu). The ADNI began in 2003 to test whether serial MRI, positron emission tomography, biofluid markers, and clinical and neuropsychological assessment can be combined to measure progression of MCI and early AD. For up-to-date information, see www.adni-info.org. Inclusion criteria for ADNI consisted of participants ages 55–90 years (inclusive), available study partner capable of accompanying participant to visits, Geriatric Depression Scale score < 6, Hachinski Ischemic Score ≥ 4 , stability of permitted medications for 4 weeks, adequate visual and auditory abilities for neuropsychological testing, adequate general health with no diseases expected to interfere with study participation, minimum of 6th grade education or equivalent work history, and fluency in English or Spanish. Exclusion criteria consisted of significant co-morbid neurological disease, history of substance abuse within the past 2 years, and history of significant head trauma. All participants received baseline clinical diagnoses of cognitively normal (CN), MCI, or AD dementia according to ADNI diagnostic criteria, which have been previously described [17]. This study was conducted in compliance with the Declaration of Helsinki and was approved at all sites by local Institutional Review Boards. All participants or legal representatives of participants gave written informed consent prior to participation in the study. For the present study, participant data consisted of 1,629 older adults enrolled in ADNI-1, ADNI-GO, or ADNI-2 with complete baseline data for all variables of interest (demographics, neuropsychological testing, baseline structural MRI). Age, sex, years of education, and apolipoprotein (*APOE*) $\epsilon 4$ carrier status were included as demographic variables. Further information regarding *APOE* $\epsilon 4$ genotyping is online (<http://adni.loni.usc.edu/data-samples/data-types/genetic-data/>).

Cluster Analysis

Due to the previously noted susceptibility of ADNI MCI diagnoses to false positives, all participants with baseline ADNI diagnoses of MCI were entered into a cluster analysis to resolve potential misclassifications [18–20]. First, a consistently cognitively normal reference group was formed from participants who were ADNI-diagnosed CN and remained CN for the length of their participation in the study ($n = 383$). Next, linear regression models were run within this group to predict cognitive performance from age and education for six neuropsychological tests (Rey Auditory Verbal Learning Test delayed memory recall, Rey Auditory Verbal Learning Test delayed memory recognition, Animal fluency, Boston Naming Test, Trail Making Test Parts A & B) across three cognitive domains (memory, language, executive function). Resulting regression coefficients were then used to calculate expected performance of MCI participants on the six neuropsychological tests based on their age and education. Finally, age- and education-adjusted z-scores (calculated based on their observed versus expected performance) were used in a hierarchical cluster analysis (Ward's method & forced 4-cluster solution) in line with prior studies to reclassify MCI participants into four previously described diagnostic groups: a cluster-derived cognitively normal group, amnesic MCI, dysnomic MCI, and dysexecutive MCI. The cluster-derived cognitively normal group was combined with ADNI-diagnosed cognitively normal individuals to form

the CN group for the present study, while the three MCI subtypes were combined into a single neuropsychologically-confirmed MCI group. ADNI-diagnosed AD dementia represented the AD group.

Neuroimaging Acquisition and Analyses

ADNI participants underwent MRI scanning on Siemens, GE, or Phillips scanners at 1.5T or 3T magnet strength. T1-weighted structural images were acquired using either a volumetric magnetization prepared rapid gradient-echo sequence (MPRAGE) or a sagittal 3D inversion-recovery prepared spoiled gradient echo imaging pulse sequence (IR-SPGR). Specific parameters for each sequence are available to view online (<http://adni.loni.usc.edu/methods/mri-tool/mri-analysis/>). Combining data from 1.5T and 3T magnetic field strengths has been previously shown to be feasible by the ADNI investigators and independent researchers, and we accordingly merged MRI scans from both 1.5T and 3T field strengths [21,22]. For all study participants, baseline T1-weighted images were first downloaded from the ADNI database (<http://adni.loni.usc.edu>) in raw NIfTI format prior to any processing. Using the “Display” function in SPM12 (<http://www.fil.ion.ucl.ac.uk/spm/>) within MATLAB (MATLAB R2018a, MathWorks Inc., Natick, MA, USA) on macOS, each T1-weighted image was individually checked for image quality and manually aligned and rotated to ensure AC-PC (anterior commissure-posterior commissure) alignment. Aligned images were processed through the voxel-based morphometry (VBM) pipeline in SPM12, which has been described in detail [23]. Briefly, each AC-PC aligned T1-weighted image was segmented into grey matter, white matter, and CSF tissue classes using SPM12’s unified segmentation procedure, followed by the creation of a study-specific DARTEL template [24–26]. Segmented images were then iteratively aligned to the DARTEL template, spatially normalized, modulated, and smoothed with an 8 mm full-width at half-maximum isotropic Gaussian kernel. Resulting smoothed, modulated, and warped tissue segmentations were used in subsequent analyses.

Region-of-interest (ROI) masks extracted whole brainstem, midbrain, pons, and locus coeruleus (LC) volumes (Supplementary Fig. 1). A previously established ROI mask defined by the grey and white matter tissue maps from the ICBM152 template was used to assess whole brainstem volumes comprising the pons, medulla, and midbrain (Supplementary Fig. 1A) [27–30]. ROI masks for the midbrain and pons were obtained from an atlas created as part of a study establishing a probabilistic Bayesian segmentation procedure for automated delineation of the brainstem and its sub-regions, and these masks have been validated in clinical populations (e.g., progressive supranuclear palsy, corticobasal syndrome) known to experience atrophy of these regions (Supplementary Fig. 1B–C) [31,32]. To approximate LC volume, we used a previously created ROI mask derived by averaging coordinates for peak voxels of functional activity and neuromelanin sensitivity from two prior studies that localized the LC on functional MRI and T1-weighted turbo spin echo MRI scans (Supplementary Fig. 1D) (<https://rcweb.dartmouth.edu/CANlab/brainstemwiki/doku.php/lc.html>) [33,34]. Volumes for brainstem ROIs were calculated by summing grey and white matter voxel values from the VBM-processed images; both grey and white matter were included for the brainstem due to the mixed tissue classifications that make up the structure [35]. Total intracranial volume (TIV) was calculated as the sum of all voxels across the grey

matter, white matter, and CSF segmented maps. Volume extractions for TIV and all ROIs were performed using built-in SPM12 functions (e.g. “spm_summarise”) and the “get_totals” script (http://www0.cs.ucl.ac.uk/staff/gridgway/vbm/get_totals.m). To correct for head size, ROI volumes were normalized via simple division by TIV, a widely used method for volumetric normalization [14,36,37]. Additional volume normalizations for midbrain, pons, and LC ROIs were performed via division by whole brainstem volume to determine specificity of observed volumetric differences. Normalized volumes were subsequently multiplied by a factor of 10^3 (whole brainstem, midbrain, pons) or 10^4 (LC) to facilitate ease of comparisons.

In addition to ROI analyses, voxel-wise two sample t-tests examined morphometric differences between groups. White matter maps were selected due to SPM’s predominant classification of the brainstem as white matter, and prior work has demonstrated the feasibility of detecting brainstem abnormalities with VBM in clinical populations [15,16,32]. VBM maps were statistically compared between groups using two-sample t-tests with age, sex, education, *APOE* e4 status, and TIV included as covariates. Additional models replaced the TIV covariate with TIV-normalized pons volume or TIV-normalized brainstem volume to address potential concerns of regional specificity. An explicit mask combining the midbrain and pons restricted analyses to rostral brainstem structures. A height threshold of $p < .05$ with family-wise error (FWE) correction for multiple comparisons was used and resulting maps were inspected for significant clusters representing groupwise volumetric differences. Where noted below, less stringent height thresholds of uncorrected $p < .05$ and $p < .01$ were used for exploratory analyses in cases where no differences were observed at FWE-corrected $p < .05$. VBM analyses listed below compared clinically-diagnosed groups, rather than biomarker-confirmed groups, to preserve subject numbers and ensure sufficient statistical power.

CSF Biomarkers

A subset of ADNI participants underwent fasting lumbar puncture at baseline, and levels of amyloid β 1–42 ($A\beta_{1-42}$) and phosphorylated tau (pTau) were quantified using the automated Roche Elecsys $A\beta_{1-42}$ CSF and Elecsys phosphotau (181P) CSF electrochemiluminescence immunoassays at the UPenn Biomarker Research Laboratory [38]. Participants were categorized as amyloid-positive with values of $A\beta_{1-42}$ below 980 pg/ml and as amyloid-negative with values at 980 pg/ml and above. They were also categorized as pTau-positive with values of pTau_{181p} at or above 21.8 pg/ml and as pTau-negative with values below 21.8 pg/ml (see <http://adni.loni.usc.edu/methods/> for more information) [39]. Participants who were both amyloid-positive and pTau-positive were considered biomarker-confirmed AD or MCI due to AD (MCI n= 203; AD n = 160) [39,40].

Statistical Analyses

All continuous variables were checked for normality via skewness and kurtosis. Substantial departure from normality was noted for the Trails A, Trails B, and Boston Naming Test variables, and a \log_{10} transformation was applied. The Boston Naming Test variable was reflected prior to \log_{10} transformation to avoid undefined values. ANOVA with Tukey’s post-hoc test was used to test group differences in age and education. χ^2 test was used to test

group differences in sex and *APOE* ϵ 4 status. ANCOVA with age, sex, education, and *APOE* ϵ 4 status as covariates tested group differences in neuroimaging variables, with post-hoc LSD tests used for pairwise group comparisons.

Longitudinal analyses were restricted to cognitively normal participants with at least one follow-up timepoint of clinical diagnostic data. Proportional hazards survival analyses were conducted via Cox regressions to determine the value of brainstem ROI volumes in predicting progression from normal cognition to clinically-diagnosed AD dementia. In Cox regression models, continuous brainstem ROI volume was first entered as a sole predictor variable, with AD dementia diagnosis as event of interest and months to diagnosis as time variable. Next, for models that were significant with a single predictor variable, relevant covariates of age, sex, education, and *APOE* ϵ 4 status were added in a second block. Hazard ratios (HR) and 95% confidence intervals (CI) were reported from all Cox regression models with brainstem ROI volumes entered as continuous predictor variables. Median splits were performed solely for visualization of risk profiles at high versus low volumes, with survival curves plotted for groups split by high or low brainstem ROI volume. To address potential bias introduced by selective attrition, all Cox regression models were initially run with variable follow-up (6–120 months) and repeated with a fixed 48-month follow-up. Additional Cox regression analyses assessed value of brainstem ROI volumes in predicting progression from ADNI-diagnosed normal cognition to ADNI-diagnosed MCI.

To address the issue of multiple comparisons, false discovery rate (FDR) correction via the Benjamini-Hochberg procedure was applied to omnibus tests for all planned comparisons, including diagnostic group and biomarker-split group comparisons of ROI volumes, post-hoc pairwise tests, and Cox regressions of converters/non-converters to dementia [41]. Results were considered significant at FDR-corrected threshold of .05 and .10. For all other unplanned comparisons (e.g., demographic variables), results were considered significant at Bonferroni adjusted $p < .05$. All statistical analyses were performed in SPSS (IBM SPSS Statistics 25, IBM, Armonk, NY, USA) and Prism (GraphPad Prism 7, GraphPad Software Inc., San Diego, CA, USA).

RESULTS

Demographics

Baseline demographic data are displayed for clinically-diagnosed and biomarker-confirmed groups in Table 1 and for AD dementia converters and non-converters in Table 2.

Brainstem Region-of-interest Analyses

Relative to the cognitively normal group, participants with neuropsychologically-confirmed MCI or clinically-diagnosed AD dementia had smaller TIV-normalized whole brainstem [$F(2,1615) = 7.13, p = .001, \eta_p^2 = .009$], midbrain [$F(2,1617) = 16.80, p < .001, \eta_p^2 = .02$], and pons [$F(2,1616) = 3.26, p = .039, \eta_p^2 = .004$] volumes (Table 1; Fig. 1A–C; Supplementary Table 1). Diagnostic groups did not differ in TIV. When normalizing brainstem substructures to whole brainstem volume, participants with AD dementia had smaller midbrain relative to whole brainstem volume [$F(2,1617) = 5.70, p = .003, \eta_p^2$

= .007] and larger pons relative to whole brainstem volume [$F(2,1620) = 7.12, p = .001, \eta_p^2 = .009$] as compared to the cognitively normal group (Supplementary Table 2A,D). When constraining analyses to biomarker-confirmed groups, AD dementia participants exhibited smaller TIV-normalized midbrain [$F(2,1150) = 4.58, p = .01, \eta_p^2 = .008$] volumes than cognitively normal participants, and there were no differences in whole brainstem or pons volumes (all p 's > .10) (Table 1; Fig. 1D–F; Supplementary Table 1). When normalizing brainstem substructures to whole brainstem volume, biomarker-confirmed AD dementia participants had smaller midbrain relative to whole brainstem volume [$F(2,1150) = 5.07, p = .006, \eta_p^2 = .009$] as compared to the cognitively normal group and larger pons relative to whole brainstem volume [$F(2,1152) = 7.84, p < .001, \eta_p^2 = .013$] as compared to the cognitively normal and MCI groups (Supplementary Table 2B,D). Cognitively normal participants who progressed to AD dementia (converters) had smaller baseline TIV-normalized midbrain [$F(1,775) = 8.68, p = .003, \eta_p^2 = .011$] volumes than those who did not progress to dementia (non-converters), and no differences were observed in TIV, TIV-normalized whole brainstem volume, TIV-normalized pons volume, or brainstem-normalized ROIs (Table 2; Fig. 2A–C; Supplementary Table 2C; see Supplementary Table 3 for life table). With FDR limited to 0.05, Cox regression analyses of relationships between baseline TIV-normalized ROI volumes and progression to dementia demonstrated that smaller baseline midbrain volume was associated with higher risk of progression to dementia (HR 3.24, 95% CI [1.51, 6.96], $p = .003$) (Fig. 2E). With FDR limited to 0.10, smaller baseline whole brainstem volume was associated with higher risk of progression to dementia (HR = 1.24, 95% CI [1.02, 1.52], $p = .033$) (Fig. 2D). Cox regression with baseline pons volume as predictor was not significant, nor were repeated Cox regression models with baseline brainstem-normalized ROI volumes as predictor variables (Fig. 2F; Supplementary Table 2E). Repeated Cox regression models with fixed 48-month follow-up periods confirmed that smaller baseline midbrain volume was associated with higher risk of progression to dementia (HR 3.14, 95% CI [1.47, 6.69], $p = .003$), however relationships for brainstem and pons were not significant (Supplementary Table 4A–B).

ROI analyses centered on the LC indicated that participants with neuropsychologically-confirmed MCI or clinically-diagnosed AD dementia had smaller TIV-normalized LC [$F(2,1616) = 4.50, p = .011, \eta_p^2 = .006$] volumes relative to the cognitively normal group, while no differences were observed when constraining analyses to biomarker-confirmed groups or when normalizing to whole brainstem volume (Table 1; Fig. 3C–D; Supplementary Table 1; Supplementary Table 2A). Baseline LC ROI volumes did not differ between AD dementia converters and non-converters at baseline when normalizing to TIV or to whole brainstem volume (Fig. 4A; Table 2; Supplementary Table 2C). With FDR limited to 0.10, Cox regression analyses demonstrated that smaller baseline LC volume conferred higher risk of progression to dementia (HR 9.10, 95% CI [1.20, 69.22], $p = .033$) (Fig. 4B). Baseline LC volume was not predictive of progression to dementia in repeated Cox regression models with fixed 48-month follow-up period or in models with brainstem-normalized LC volume (Supplementary Table 2E; Supplementary Table 4).

ROI analyses were additionally repeated with raw ROI volumes (as opposed to TIV-normalized volumes) and TIV included as a covariate in statistical models; this approach did not affect any results. In a risk analysis examining ADNI-diagnosed CN individuals who

progressed to an ADNI diagnosis of MCI, baseline midbrain volume was associated with higher risk of MCI diagnosis (HR 2.26, 95% CI [1.20, 4.27], $p = .012$) (Supplementary Table 5A–B).

Brainstem-Masked VBM Analyses

Brainstem-masked VBM analyses of the entire MCI sample, regardless of biomarker positivity, compared to cognitively normal participants indicated smaller regional volume of clusters overlapping the bilateral LC and bilateral clusters in the anterolateral midbrain (Fig. 3A; Supplementary Table 6A). Patterns of regional volume difference between AD dementia and cognitively normal participants similarly indicated smaller clusters overlapping bilateral LC, anterolateral midbrain, and dorsal rostral pontine regions (Fig. 3B; Supplementary Table 6B). All VBM findings were significant at an FWE-corrected height threshold of $p < .05$. Brainstem-masked VBM analyses within cognitively normal participants showed smaller regional volume of bilateral clusters corresponding to the anatomical distribution of the LC in AD dementia converters compared to non-converters, the only clusters that remained significant at an uncorrected height threshold of $p < .01$ (Fig. 4C; Supplementary Table 7A). When observed at a less stringent threshold of uncorrected $p < .05$, the clusters of interest extended caudally, further overlapping the anatomical distribution of the LC (Supplementary Table 7B; Supplementary Fig. 2). We compared MNI coordinates from our voxel-wise analyses and found specific overlap with prior VBM studies of the brainstem and with studies that have localized the structure of the LC on neuromelanin-sensitive T1-weighted sequences (Supplementary Table 8). In order to demonstrate that voxel-wise findings were not a reflection of overall pons or overall brainstem volume difference, VBM analyses were repeated with covariates for TIV-normalized pons volume and TIV-normalized whole brainstem volume in place of the TIV covariate. Clusters overlapping the LC remained significant at FWE-corrected $p < .05$ in AD compared to CN with pons covariate, and at uncorrected $p < .05$ in AD compared to CN with whole brainstem covariate, as well as MCI compared to CN with pons covariate and whole brainstem covariate (Supplementary Figs. 3–4). Similarly, clusters overlapping the LC remained significant at uncorrected $p < .05$ in AD dementia converters compared to non-converters when controlling for TIV-normalized pons and whole brainstem volumes (Supplementary Figs. 5–6).

DISCUSSION

The current study found that older adults with biomarker-confirmed dementia due to AD exhibited smaller midbrain volumes than cognitively normal individuals. Furthermore, smaller midbrain volumes were observed in cognitively normal older adults who later went on to develop AD dementia compared to those who did not progress to dementia, and lower baseline brainstem, midbrain, and LC volumes were predictive of future progression to AD dementia diagnosis. These findings confirm prior MRI studies implicating brainstem volumetric differences in clinically-diagnosed AD dementia and further clarify that these differences are observable earlier in AD pathophysiological processes [12,15,31]. A brainstem-masked analysis using voxel-level methods revealed further brainstem differences between neuropsychologically-confirmed MCI and clinically-diagnosed dementia due to AD compared to cognitively normal individuals in a small cluster corresponding with the

anatomical location of the LC along the midbrain-pontine axis. Our findings provide preliminary *in vivo* evidence of structural brainstem abnormalities detectible on traditional MRI sequences, mirroring neuropathological studies that have localized early AD pathology to brainstem nuclei [42,43]. Taken together, the patterns of brainstem volumetric differences across clinically-diagnosed and biomarker-confirmed AD groups suggest early brainstem pathology in the midbrain and LC is observable on MRI, and this pathology is predictive of clinical progression from the earliest preclinical phase of the disease.

A strength of the present study was the use of VBM in addition to ROI-based volumetrics to compare diagnostic groups and constrain analyses to brainstem substructures. Our finding of reduced LC volume in MCI and AD dementia compared to cognitively normal individuals provides a volumetric analogue to previous studies showing reduced LC neuromelanin contrast ratios in AD and MCI relative to cognitively normal individuals [44,45]. To our knowledge, no prior studies have outlined proxy estimates of LC volume on traditional T1-weighted images and instead have focused on neuromelanin-sensitive T1-weighted scans. Our study represents the first known effort to evaluate LC integrity with volumetrics as opposed to neuromelanin contrast ratio. Despite the difficulty in quantifying the volume of a nucleus as small as the LC on structural MRI, groupwise differences in LC volume were in the expected direction, with smaller LC volume seen in cognitive impairment and predicting future cognitive impairment. Clusters corresponding to the bilateral LC resulting from our voxel-wise analyses overlap with coordinates reported from prior studies using neuromelanin-sensitive T1-weighted FSE scans, demonstrating the potential utility of traditional T1-weighted scans in detecting LC volumetric differences [34,46,47]. Our analyses indicated gross detectible differences in the midbrain while voxel-level analyses revealed sub-regional differences in midbrain and pontine regions adjacent to nuclei known to degenerate with advancing Braak stage [43]. Our midbrain-specific findings in biomarker-confirmed AD cases emphasize that volumetric differences are detectible in a structure known to undergo selective neuronal loss in the earliest precortical stages of AD and may be indicative of compromised optimal regulation of various arousal-related systems (e.g., serotonergic, glutamatergic, cholinergic, noradrenergic) in AD progression. [1,43,48]. Future *ex vivo* neuropathological studies will be helpful in clarifying whether our findings represent a proxy measure of neurodegeneration, synaptic loss, axonal deterioration, or a different neuropathological marker, but in any case our approach has established detectible differences in brainstem MRI in very early stage AD.

Given that smaller midbrain volume in cognitively normal older adults predicts future dementia, it is possible that greater premorbid midbrain volume confers a degree of protection against insidious tau deposition and consequent disease progression. This is consistent with the neural reserve literature suggesting the LC as a site of reserve due to its known involvement in higher-order executive cognitive processes, neuroplasticity, memory, and arousal, and our findings support a critical role of the LC in protecting against the deleterious effects and corresponding clinical consequences of increasing AD pathology [49–51]. Alternatively, our findings could be interpreted as suggesting brainstem regions are selectively vulnerable to neurodegenerative disease. As we demonstrated, reduced premorbid structural integrity of the midbrain and LC bestows greater risk for cognitive decline and disease progression, in line with studies observing reduced LC neuromelanin

contrast ratios in AD dementia and MCI populations compared to cognitively normal individuals, inverse correlations between LC neuromelanin contrast ratio and A β pathological burden, and general declines in LC neuromelanin contrast with age [44,45,52,53]. Prior studies have examined LC degeneration in the early stages of AD pathophysiology in postmortem tissue, and the present study provides preliminary evidence that these differences may be apparent *in vivo* with widely used structural neuroimaging techniques [1,2,54,55]. The LC's role as the primary site of noradrenergic production and regulation has increased its potential as a biomarker for neurodegenerative disease, and our findings suggest that gross anatomical differences quantified on T1-weighted scans may be useful when assessing preclinical and prodromal populations in contexts where more advanced imaging sequences are unavailable [56,57].

Alternative pathways of dysfunction must also be considered when examining a nucleus as functionally diverse as the LC. Recent animal studies have suggested that LC integrity is linked to optimal regulation of cerebral blood flow, with degeneration of the LC-norepinephrine projection system predicting downstream vascular consequences in AD-related regions [58,59]. Multimodal neuroimaging and biomarker studies of well-characterized clinical populations are needed to disentangle whether subcortical brainstem nuclei represent sites of selective vulnerability or resilience to AD pathological burden. In the context of ongoing debate over whether subcortical regional pathology in the LC and other brainstem nuclei represent the initial site of tau seeding and hyperphosphorylated tau deposition in AD, our study cannot argue one way or the other due to the lack of histopathological analysis and Braak staging. However, the observation of early brainstem volume differences in MCI due to AD and cognitively normal older adults at risk for future AD dementia diagnosis clearly emphasizes the importance of integrating *in vivo* neuroimaging studies with histopathological studies to continue characterizing the disease-related progression of tau pathology.

The present study is not without several caveats. One limitation is the estimation of LC volume from structural T1 images via approximated ROI masks. The LC is a notoriously difficult nucleus to localize on images and neuromelanin-sensitive T1-weighted fast spin echo or turbo spin echo sequences are thought to best visualize and capture the integrity of the LC [34,47,49]. These specialized scans leverage the natural accumulation of neuromelanin in noradrenergic cells of the LC over the lifespan, which has paramagnetic T1-shortening effects, and apply targeted scan parameters to capture this neuromelanin concentration as hyperintensities visible in the pons [46,60]. The present study used T1-weighted volumetric sequences and an LC mask centered on average peak voxels from multiple studies to approximate LC volume. Future studies will need to compare this methodology to neuromelanin-sensitive sequences [33,34]. Although we demonstrated groupwise differences among diagnostic groups in ROI volume of LC when normalizing to TIV, this finding did not remain significant when normalizing to whole brainstem volume. This may represent artifact specific to our methodology, as various processing steps including warping to template space and spatial smoothing make it difficult to ensure regional specificity in a mixed tissues structure such as the brainstem. VBM methodologies contain inherent limitations for evaluating grey/white matter contrast in a mixed tissue class structure such as the brainstem, and it has been previously noted that VBM may have a

limited capacity to detect subtle changes in white matter regions that are largely homogenous in nature [61,62]. However, prior studies have used VBM analysis of white matter maps to successfully detect volume loss within brainstem substructures in disease populations known to experience specific volume loss within the brainstem (progressive supranuclear palsy and corticobasal degeneration), as well as in Alzheimer's disease compared to healthy controls [15,32]. Our study further adds to this literature by providing additional support for the feasibility of this approach within the greater context of the limitations VBM analyses pose. Additionally, regional findings from our voxel-wise analyses remained significant when correcting for total pons volume or total brainstem volume. Nevertheless, future studies should conduct similar analyses in cohorts with specialized neuromelanin-sensitive MRI sequences designed to specifically assess LC structural integrity [57].

Survival analyses indicated that baseline LC volume confers significant risk for dementia, but the large 95% CI suggests the stability of this prediction is highly variable, and predictive value was no longer significant when examining over a fixed 48-month follow-up. Additionally, the present study analyzed a sample with high rates of MCI and AD cases likely not present in real-world settings, suggesting that a similar analysis in a smaller sample would result in lower predictive value. Thus, the predictive value of the LC should be interpreted with caution and future studies in more representative samples will help to determine the stability of predictive values and may aim to make comparisons with regions well-established to experience atrophic changes in AD dementia (e.g., hippocampus, medial temporal lobe). Nevertheless, the close anatomical localization provided from the LC ROI mask, combined with the marked volumetric differences observed in VBM analyses, implicate midbrain and pontine regions adjacent to and overlapping the LC in preclinical AD. Given the clinical subjectivity of differential diagnoses in ADNI and the lack of pathological diagnostic confirmation in our cohort, it is possible that our findings in the clinically-defined groups were influenced by individuals with co-morbid subcortical dementias, primary age-related tauopathies, and other non-AD processes. We examined cross-sectional volumetric differences at baseline rather than progression of brain atrophy over time purposefully in an effort to comprehensively quantify baseline volumetrics. Future studies will aim to detail the longitudinal progression of brainstem substructure atrophy in each of our diagnostic groups. It should be noted that although our analysis predicting AD dementia used neuropsychologically-confirmed cluster analysis to refine baseline MCI diagnoses, our analysis predicting MCI used original ADNI MCI diagnoses. Future efforts should use neuropsychological data to inform serial MCI diagnoses to avoid potential misclassifications, as demonstrated by recent studies [63]. Additionally, our longitudinal risk analyses included clinically-diagnosed groups and were not performed in biomarker-confirmed groups due to limited sample sizes. Thus, longitudinal studies are needed to detail brainstem atrophy progression in biomarker-confirmed MCI and AD. Finally, the ADNI database comprises an ethnically homogenous, highly educated sample that is not necessarily representative of the general population, and future studies will need to replicate our findings in diverse cohorts.

Despite these limitations, our study demonstrates brainstem MRI abnormalities that are detectable in preclinical populations, highlighting the importance of considering the

brainstem when developing novel biomarkers and innovative therapeutics. Future studies should leverage additional neuroimaging modalities, including functional MRI, diffusion tensor imaging, and arterial spin labeling, to address issues of neurovascular coupling, structural white matter tract degeneration, and cerebral perfusion as they relate to the brainstem and disease progression. Future endeavors should also separate individuals by Braak stage and quantify these brainstem substructure volumes to examine whether brainstem atrophy progresses in temporal conjunction with the widely accepted pathophysiological staging advancement. Future MRI-based studies of dementia populations may aim to include brainstem volumetrics as outcome variables and address their potential utility as clinical trial endpoints [15,31,64]. In summary, we provide here preliminary evidence that *in vivo* visualization of brainstem substructure and LC-specific differences are detectible with widely used MRI sequences in preclinical and prodromal AD populations.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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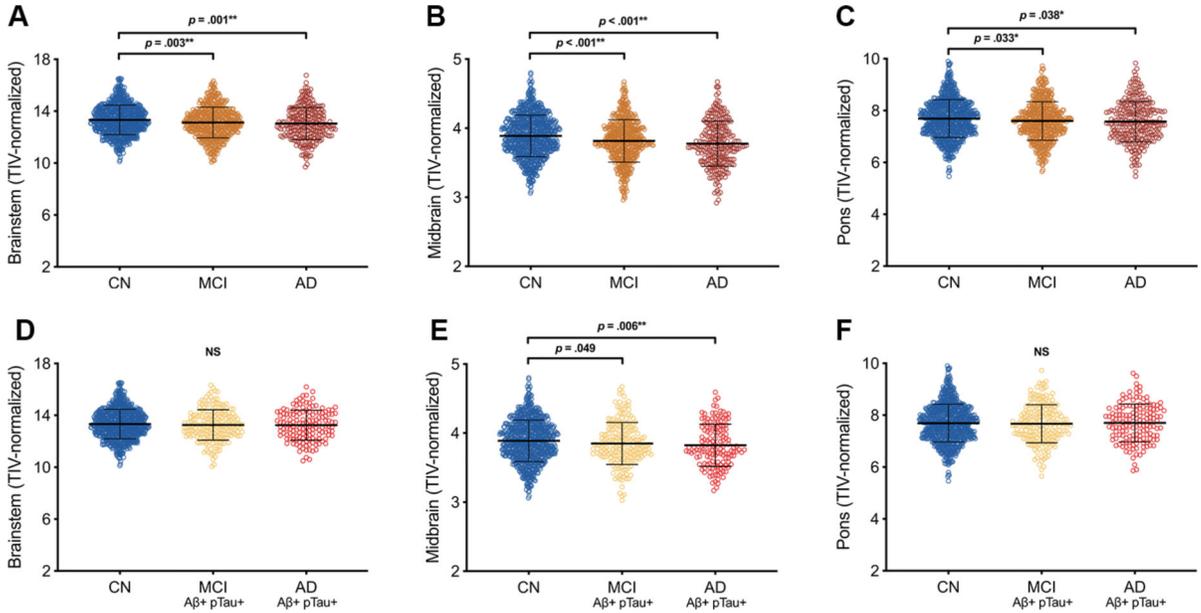


Figure 1. Brainstem gross regional volumes in cognitively normal, MCI, and Alzheimer’s disease participants

Relative to older adults who are cognitively normal (CN; blue dots), those with mild cognitive impairment (MCI; orange dots) and clinically-diagnosed Alzheimer’s disease dementia (AD; dark red dots) exhibit smaller volumes of overall brainstem (A), midbrain (B), and pons (C). In a subgroup with biomarker-confirmed AD pathology based on CSF Aβ₁₋₄₂ and pTau abnormalities, brainstem volume differences are specific to the midbrain in AD dementia (bright red dots) (D-F). All *p*-values reported are the results of post-hoc Fisher’s LSD pairwise comparisons following one-way ANCOVA controlling for age, sex, education, and *APOE* ε4 status. *Remains significant with FDR limited to .10; **Remains significant with FDR limited to .05 or .10. Error bars represent ± 1 standard deviation. Abbreviations: AD = Alzheimer’s disease; CN = cognitively normal; MCI = mild cognitive impairment; NS = non-significant.

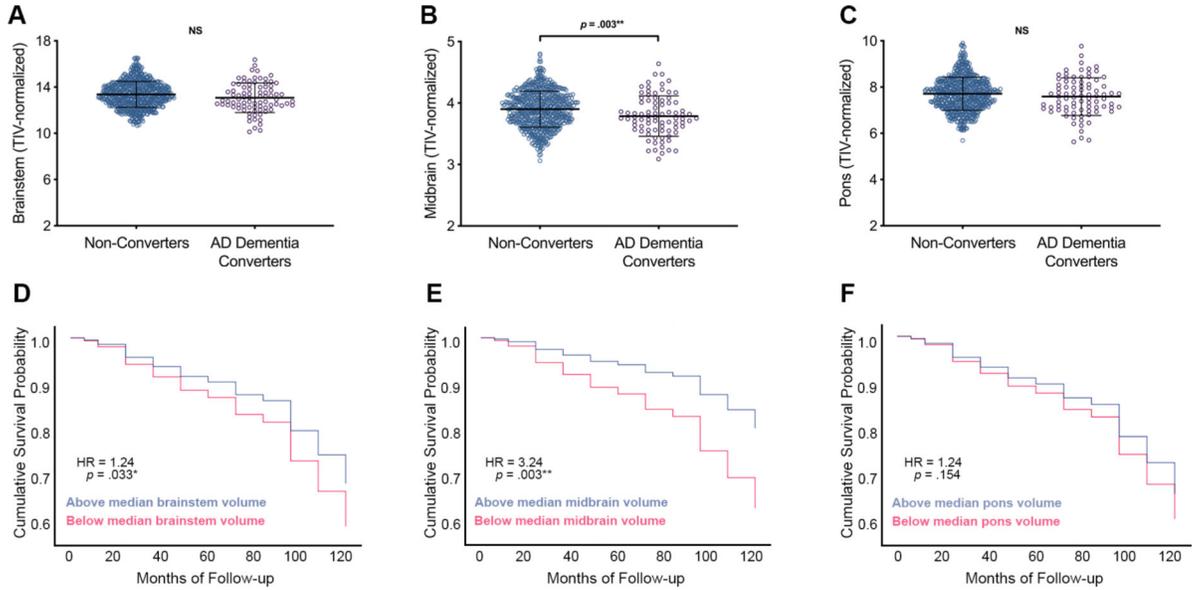


Figure 2. Brainstem gross regional volumes in progression to AD dementia

CN older adults who prospectively progress to dementia (AD dementia converters; purple dots) demonstrated smaller volume of the midbrain (B) at baseline relative to those who did not progress to dementia (non-converters; blue dots), while no differences were observed in whole brainstem (A) and pons (C). CN older adults with smaller baseline midbrain volumes (pink lines) are more likely to later develop cognitive impairment and receive a clinical diagnosis of AD dementia over 6 to 120 months of follow-up than those with larger baseline midbrain volumes (blue lines) (D-F). Hazard ratios (HR) and *p* values report results of proportional hazards Cox regressions with continuous brainstem ROI volume as predictor variable, AD dementia diagnosis as event of interest, and months to diagnosis as time variable (event cases = 83; censored cases = 702). ROI volume was first entered as a sole continuous predictor variable, and significant models added in covariates of age, sex, education, and *APOE* ϵ 4 status. For display purposes, median splits were performed on brainstem ROI volumes to show risk at high and low volumes. *Remains significant with FDR limited to .10; **Remains significant with FDR limited to .05 or .10. Abbreviations: AD = Alzheimer’s disease; CN = cognitively normal; NS = non-significant.

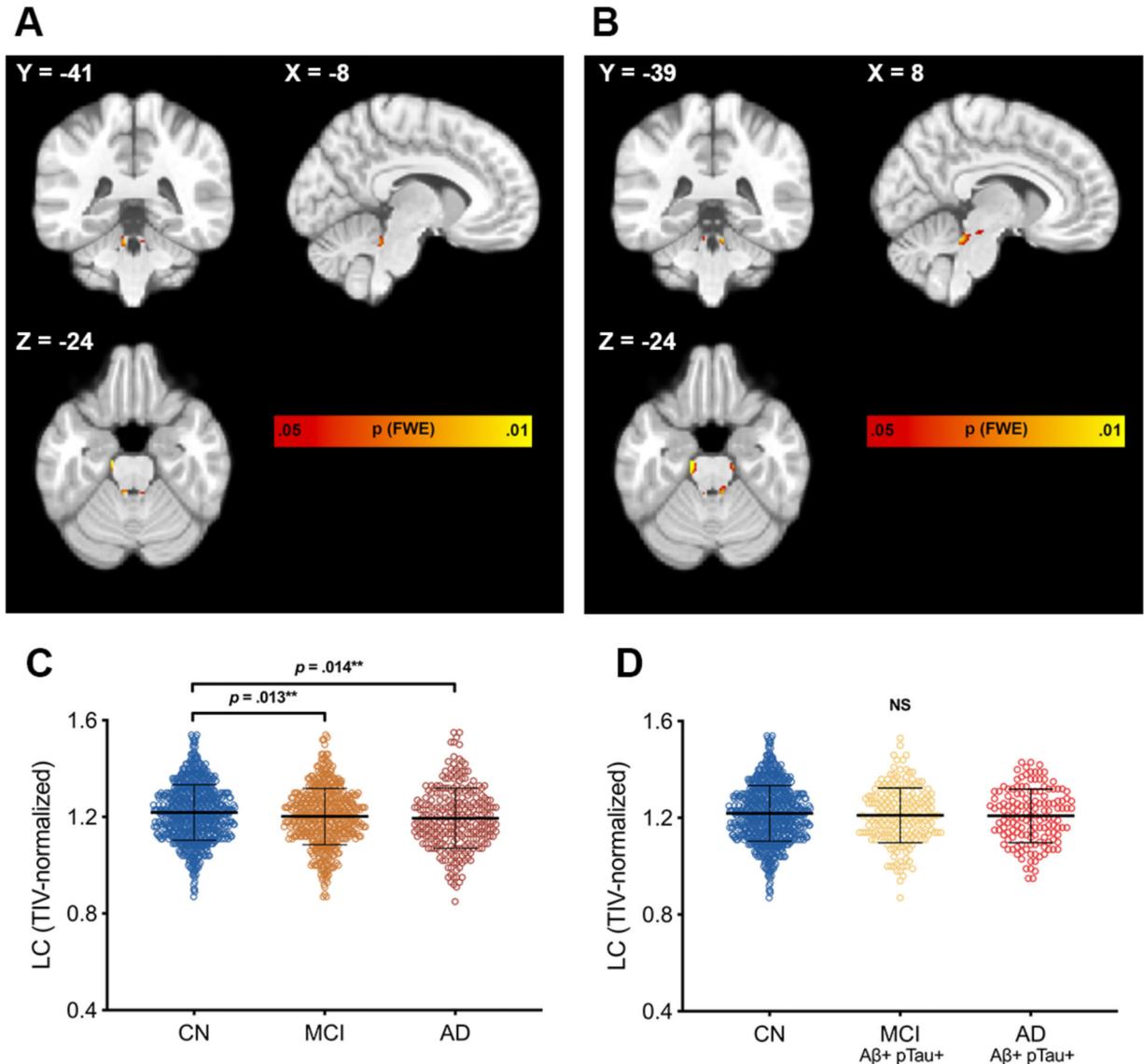


Figure 3. Brainstem-masked VBM comparisons and regional LC volumes

Brainstem-masked VBM analyses in the overall sample revealed a specific pattern of reduced regional dorsal rostral brainstem volume at baseline in MCI (A) and clinically-diagnosed AD (B) relative to those who are CN that closely corresponds to the anatomical distribution of the LC. VBM analyses consisted of two-sample t-tests with age, sex, education, *APOE* ϵ 4 status, and TIV as covariates. Results are displayed at FWE-corrected height threshold of $p < .05$, represented by color bar. Images are displayed in neurological orientation. LC ROI volumes, extracted using an LC atlas mask, (C) were smaller in MCI (orange dots) and AD (dark red dots) relative to cognitively normal older adults (blue dots). No significant differences were observed between biomarker-confirmed subgroups (D). p -values reported are the results of post-hoc Fisher's LSD pairwise comparisons following one-way ANCOVA controlling for age, sex, education, and *APOE* ϵ 4 status. *Remains significant with FDR limited to .10; **Remains significant with FDR limited to .05 or .10.

Error bars represent ± 1 standard deviation. Abbreviations: AD = Alzheimer's disease; CN = cognitively normal; MCI = mild cognitive impairment; NS = non-significant.

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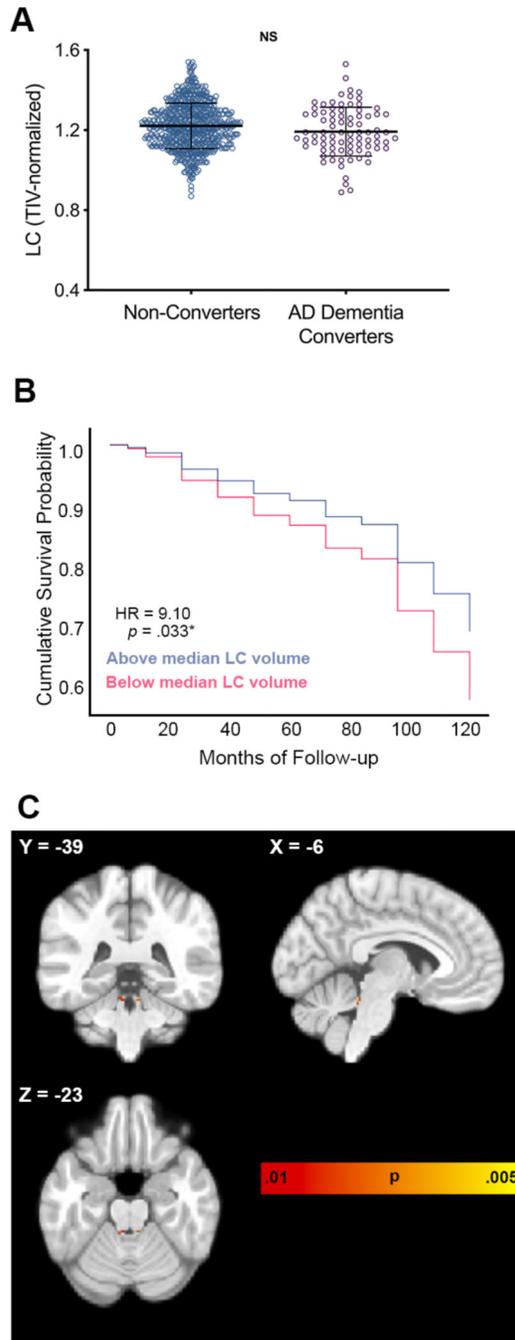


Figure 4. LC volume predicts progression to dementia in cognitively normal older adults
 AD dementia converters (purple dots) did not differ from non-converters (blue dots) in LC volume at baseline (A). CN older adults with smaller baseline LC volumes (pink lines) are more likely to later develop cognitive impairment and receive a clinical diagnosis of AD dementia over 6 to 120 months of follow-up than those with larger baseline midbrain volumes (blue lines) (B). Hazard ratios (HR) and p -values report results of proportional hazards Cox regressions with continuous brainstem ROI volume as predictor variable, AD dementia diagnosis as event of interest, and months to diagnosis as time variable (event

cases = 83; censored cases = 702). ROI volume was first entered as a sole continuous predictor variable, and significant models added in covariates of age, sex, education, and *APOE* ϵ 4 status. For display purposes, median split was performed on LC ROI volume to show risk at high and low volumes. Brainstem-masked VBM analyses revealed a specific pattern of reduced regional dorsal rostral brainstem volume at baseline in AD dementia converters compared to non-converters (C) that closely corresponds to the anatomical distribution of the LC. VBM analyses consisted of two-sample t-tests with age, sex, education, *APOE* ϵ 4 status, and TIV as covariates. VBM results are displayed at uncorrected height threshold of $p < .01$, represented by color bar. Images are displayed in neurological orientation. *Remains significant with FDR limited to .10. Error bars represent ± 1 standard deviation. Abbreviations: AD = Alzheimer's disease; CN = cognitively normal; L = left; NS = non-significant; R = right.

Table 1.

Baseline demographics and neuroimaging data for cognitively normal, MCI, and AD

	CN	MCI	AD	<i>F</i> or χ^2	<i>p</i> -value	η^2 or Cramer's V
Demographics						
<i>n</i>	814	542	273			
Age	73.49 (6.76)	73.54 (7.35)	75.12 (7.74)	5.78	.003	.007
Sex (M/F)	417/397	332/210	153/120	13.30	.001	.09
Education	16.29 (2.65)	15.85 (2.92)	15.23 (2.92)	14.20	<.001	.017
<i>APOE</i> ϵ 4 (0/1/2 ϵ 4)	536/246/32	249/221/72	87/132/54	138.36	<.001	.206
Neuroimaging ^{a,b}						
TIV	1499.35 (146.12)	1517.81 (157.29)	1504.57 (166.64)	0.47	.628	.001
Brainstem	13.33 (1.14)	13.13 (1.18)	13.05 (1.24)	7.13	.001	.009
Midbrain	3.89 (0.30)	3.82 (0.31)	3.78 (0.33)	16.80	<.001	.02
Pons	7.69 (0.73)	7.60 (0.74)	7.58 (0.78)	3.26	.039	.004
LC	1.22 (0.11)	1.20 (0.12)	1.19 (0.12)	4.50	.011	.006
Demographics						
	CN	MCI [A β +pTau+]	AD [A β +pTau+]			
<i>n</i>	814	202	145			
Age	73.49 (6.76)	73.61 (7.13)	73.91 (8.01)	0.23	.797	<.001
Sex (M/F)	417/397	112/90	78/67	1.31	.521	.034
Education	16.29 (2.65)	15.98 (2.86)	15.58 (2.72)	4.68	.009	.008
<i>APOE</i> ϵ 4 (0/1/2 ϵ 4)	536/246/32	57/104/41	31/77/37	197.6	<.001	.292
Neuroimaging ^{a,c}						
TIV	1499.35 (146.12)	1505.68 (166.34)	1508.55 (172.89)	1.05	.35	.002
Brainstem	13.33 (1.14)	13.26 (1.17)	13.24 (1.15)	0.98	.376	.002
Midbrain	3.89 (0.30)	3.85 (0.30)	3.82 (0.30)	4.58	.01	.008
Pons	7.69 (0.73)	7.67 (0.73)	7.70 (0.73)	0.15	.859	<.001
LC	1.22 (0.11)	1.21 (0.11)	1.21 (0.11)	0.76	.469	.001

Mean (standard deviation) are shown for all variables except for sex and *APOE* ϵ 4. *F* or χ^2 are result of one-way ANOVA (age, education), chi-square test of independence (sex, *APOE* ϵ 4), or one-way ANCOVA (all other variables; covariates = age, sex, education, *APOE* ϵ 4). Effect sizes are displayed as η^2 (age, education, neuroimaging variables) or Cramer's V (sex, *APOE* ϵ 4). Results from pairwise comparisons are displayed in Supplementary Table 1. Omnibus *p*-values for planned comparisons (neuroimaging variables) remained significant when FDR rate was limited to 0.05 (exception: pons CN vs. MCI vs. AD) and 0.10 (all comparisons).

^aTIV is presented in milliliters. ROI volumes were normalized using the following equation: (ROI volume / TIV) \times 10³ for brainstem, midbrain, and pons or (ROI volume / TIV) \times 10⁴ for LC

^bROI neuroimaging variables were screened for outliers (\pm 3 standard deviations from group mean) prior to statistical analyses. Revised n's (CN/MCI/AD) by brainstem ROI are as follows: TIV(813/541/273); brainstem (809/541/272); midbrain (810/541/273); pons (810/541/272); LC (808/542/273)

^cRevised n's (CN/MCI [A β +pTau+]/AD [A β +pTau+]) by brainstem ROI are as follows: TIV(813/202/145); brainstem (809/202/145); midbrain (810/202/145); pons (810/202/145); LC (808/202/145)

Abbreviations: A β , amyloid β ; AD, Alzheimer's disease; *APOE* ϵ 4, apolipoprotein ϵ 4; CN, cognitively normal; FDR, false discovery rate; LC, locus coeruleus; MCI, mild cognitive impairment; pTau, phosphorylated tau; ROI, region-of-interest; TIV, total intracranial volume

Table 2.

Baseline demographics and neuroimaging data for cognitively normal participants who progressed to dementia (converters) and did not progress to dementia (non-converters)

	Non-converters	Converters	<i>F</i> or χ^2	<i>p</i> -value	η^2 or Cramer's V
Demographics					
<i>n</i>	702	83			
Age	73.38 (6.79)	75.17 (6.29)	5.18	.023	.007
Sex (M/F)	350/352	54/29	6.87	.009	.094
Education	16.36 (2.68)	15.71 (2.40)	4.50	.034	.006
<i>APOE</i> ϵ 4 (0/1/2 ϵ 4)	482/198/22	35/38/10	30.01	<.001	.196
Neuroimaging ^{a,b}					
TIV	1498.76 (147.91)	1520.87 (132.93)	0.16	.688	<.001
Brainstem	13.36 (1.12)	13.07 (1.28)	3.64	.057	.005
Midbrain	3.90 (0.29)	3.79 (0.33)	8.68	.003	.011
Pons	7.71 (0.72)	7.59 (0.82)	1.63	.202	.002
LC	1.22 (0.11)	1.19 (0.12)	2.96	.086	.004

Mean (standard deviation) are shown for all variables. *F* or χ^2 are result of one-way ANOVA (age, education), chi-square test of independence (sex, *APOE* ϵ 4), or one-way ANCOVA (all other variables; covariates = age, sex, education, *APOE* ϵ 4). Effect sizes are displayed as η^2 (age, education, neuroimaging variables) or Cramer's V (sex, *APOE* ϵ 4). All omnibus *p*-values for planned comparisons (neuroimaging variables) remained significant when FDR rate was limited to 0.05 and 0.10

^aTIV volume is presented in milliliters. ROI volumes were normalized using the following equation: (ROI volume / TIV) $\times 10^3$ for brainstem, midbrain, and pons or (ROI volume / TIV) $\times 10^4$ for LC

^bROI neuroimaging variables were screened for outliers (± 3 standard deviations from group mean) prior to statistical analyses. Revised *n*'s (non-converters/converters) by brainstem ROI are as follows: TIV (701/83), brainstem (697/83), midbrain (698/83), pons (697/83), LC (696/83)

Abbreviations: *APOE* ϵ 4, apolipoprotein ϵ 4; FDR, false discovery rate; LC, locus coeruleus; ROI, region-of-interest; TIV, total intracranial volume

Table 3.

Cox regression models predicting dementia risk from baseline brainstem volumes

	HR	95% CI	p-value
Model 1a			
Brainstem	1.22	[1.01, 1.48]	.043
Model 1b			
Brainstem	1.24	[1.02, 1.52]	.033
Age	1.03	[1.00, 1.07]	.085
Sex	0.68	[0.42, 1.09]	.105
Education	0.93	[0.86, 1.00]	.059
<i>APOE</i> ε4	2.76	[1.98, 3.84]	<.001
Model 2a			
Midbrain	3.05	[1.47, 6.32]	.003
Model 2b			
Midbrain	3.24	[1.51, 6.96]	.003
Age	1.03	[0.99, 1.06]	.167
Sex	0.70	[0.44, 1.12]	.139
Education	0.92	[0.85, 1.00]	.047
<i>APOE</i> ε4	2.81	[2.01, 3.92]	<.001
Model 3			
Pons	1.24	[0.92, 1.68]	.154
Model 4a			
LC	9.16	[1.32, 63.55]	.025
Model 4b			
LC	9.10	[1.20, 69.22]	.033
Age	1.03	[0.99, 1.07]	.107
Sex	0.67	[0.42, 1.07]	.091
Education	0.93	[0.86, 1.01]	.065
<i>APOE</i> ε4	2.74	[1.97, 3.82]	<.001

Brainstem ROI volumes were reflected prior to Cox regression analyses to ensure consistent directionality, with smaller volume predicting greater risk. *p*-values for brainstem ROI variables were significant with FDR limited to 0.05 (midbrain) and 0.10 (brainstem, midbrain, LC).