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Alzheimer's disease-associated cortical atrophy does not differ between Chinese and whites

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Abstract

Purpose: To assess whether there are differences in Alzheimer's disease (AD)-associated atrophy regions in Chinese and white AD patients versus cognitively normal older adults, and to test whether associations between clinical severity and gray matter volume are similar or different across these ethnic groups in a cross-sectional analysis.

Methods: Chinese and white AD patients, individuals with mild cognitive impairment (MCI), and cognitively normal (CN) controls (n = 46 white and 48 Chinese) were clinically evaluated at an academic center within one year of MRI acquisition. Clinical severity was assessed using the Clinical Dementia Rating scale sum of boxes and cortical atrophy was measured via voxel-based morphometry as well as Freesurfer. Chinese and white cohorts were demographically matched for age, sex, and education.

Results: Clinical severity by diagnosis was similar across ethnicities. Chinese and white patient groups showed similar amounts of atrophy in the regions most affected in AD after accounting for demographic variables and head size. There was no significant difference between ethnic groups when compared by atrophy and clinical severity.

Conclusion: Our study suggests that Chinese and white AD patients, when matched demographically, are clinically and neuroanatomically similar on normalized measures of cortical atrophy and clinical severity.

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DISCLOSURE STATEMENT

The authors have no conflicts of interest to disclose.

Keywords

Alzheimer's disease; neuroimaging; diversity; Chinese Americans; clinical trial design

INTRODUCTION

Understanding the similarities and differences in Alzheimer's disease (AD) across ethnic groups (e.g., Asian versus European) could improve clinical evaluation and management of AD around the globe. Prior studies have found differences in AD risk across ethnicities; for example, older African Americans and Hispanics are more likely than older whites to have AD and other dementias¹ and may have longer survival compared with white AD patients². Interestingly, the AD genetic risk factor *APOE-ε4* has weaker effects in African Americans and Hispanics but stronger effects in Japanese compared to white individuals³. However, despite evidence for differences in AD risk across ethnicities and the ~3.4 million ethnic Chinese in the U.S.⁴, few studies of dementia have focused on Chinese-Americans and the similarities and differences to whites across the AD spectrum.

While neuropsychiatric symptoms (e.g., anxiety, delusions and depression)^{5,6} and some subscales of neuropsychological screens⁷ are different between Chinese and white AD populations, fewer studies have explored differences in imaging biomarkers between these ethnic groups. Two prior studies found morphological brain features and differences between cognitively normal (CN) English-speaking whites and Chinese-speaking Asians^{8,9}. We previously found that *APOE-ε4* is associated with lower volume in AD-affected regions in cognitively normal Chinese but not whites¹⁰. The goals of our study are two-fold: (1) to examine the association between clinical severity (assessed with the Clinical Dementia Rating Sum of Boxes [CDR-SB]) and gray matter (GM) volume loss in Chinese and white individuals and (2) to assess whether there are atrophy differences in AD-associated regions between the two groups across the AD spectrum. To ensure our volumetric findings were robust across neuroanatomically diverse individuals, we measured volume loss using two analytically different, independent image analysis pipelines. Understanding the similarities and differences between Chinese and white AD patients is important because any ethnic-specific differences could impact the interpretation of research or clinical trials utilizing diverse participants, particularly if clinical or neuroimaging measures are used to assess treatment outcome.

MATERIALS AND METHODS

Study Participants and Assessment

All participants were recruited from the San Francisco Bay Area as part of on-going studies of healthy aging and AD administered by the University of California, San Francisco (UCSF) Memory and Aging Center (MAC). All participants underwent a multi-step screening process requiring at least one in-person visit to the MAC or a MAC-sponsored location (Chinese participants were eligible to be seen at alternate locations described below). To be eligible for this study, a participant required a 1.5 Tesla (T) or 3T MRI scan at our center as well as clinical and cognitive assessments within one year of the MRI scan.

During the screening process, all participants underwent a neurologic exam, detailed cognitive assessment^{11,12}, and medical history. The tests included in the cognitive assessment varied depending upon whether each study participant chose to undergo testing in English or Chinese (Mandarin or Cantonese). For those testing in English, the cognitive assessment included all tests administered as part of the National Alzheimer's Disease Coordinating Center (NACC) Uniform Data Set (UDS)¹³. For participants testing in Chinese, the cognitive assessment included the Chinese version of the Cognitive Abilities Screening Instrument (CASI-C)¹⁴, which assesses memory, language, visuospatial, and executive function. Using the CASI-C, it is possible to derive a Chinese language Mini-Mental State Exam (MMSE) score. Beyond the CASI-C, participants who chose to test in Chinese underwent the Clinical Dementia Rating Scale (CDR)¹⁵; Geriatric Depression Scale (GDS)¹⁶; the Chinese Version of the Verbal Learning Test (CVVLT)¹⁷ or the Common Objects Memory Test (COMT)¹⁸; Stroop Color Word Test¹⁹; a design fluency task from the Ruff Figural Fluency Test (RFFT)²⁰; Filled Dots trial of the Delis-Kaplan Executive Function System Design Fluency (D-KEFS DF)²¹; Benson figure²²; digit span forward and backward²³; Color Trails Form A, trials 1 (CTT-1) and 2 (CTT-2)²⁴; and category verbal fluency test (animals, vegetables)²⁵. Each participant had a study partner who was interviewed to help evaluate the participant's functional abilities. Informants also completed the CDR, a well-validated test sensitive to the changes most often seen in AD²⁶ and a clinically useful staging instrument²⁷. The CDR has been shown to be valid in studies of Asian individuals²⁸. A multidisciplinary team composed of a neurologist, neuropsychologist, and nurse then reviewed each participant's data and determined the participant's clinical diagnosis. Mild cognitive impairment (MCI) was diagnosed according to criteria as described in Petersen *et al*²⁹. AD was diagnosed according to criteria as described in McKhann *et al*³⁰.

Chinese participants were recruited through the Chinese Outreach portion of the MAC Alzheimer's Disease Research Center (ADRC). Chinese participants were evaluated at the MAC or at two alternate locations in San Francisco's Chinatown neighborhood. Chinese participants were consented and received cognitive tests in their preferred language (Mandarin, Cantonese, or English). All printed materials were also available in Chinese.

Chinese and white study participants were selected to maximize similarities across demographic variables (sex, age, education) by diagnosis to reduce confounding from factors outside of race that may contribute to brain volume and clinical severity. We first included all Chinese participants with imaging data available that passed post-processing quality control and then selected white participants to best match the available Chinese cohort. All participants or proxies provided IRB-approved, written informed consent prior to participation, and all tests were approved by the University of California, San Francisco Committee on Human Research.

Structural Image Acquisition

Study participants underwent MRI scanning within one year of CDR administration and had at least one T1-weighted MR image available for analysis. Individuals were scanned at the UCSF Neuroscience Imaging Center (NIC) or at the UCSF Veterans Affairs Medical Center

(SFVA). Scans from the UCSF NIC were acquired using a 3.0 Tesla Siemens (Siemens, Iselin, NJ) TIM Trio scanner equipped with a 12-channel head coil using a magnetization prepared rapid gradient echo (MPRAGE) sequence (160 sagittal slices; slice thickness, 1.0 mm; field of view (FOV), $256 \times 230 \text{ mm}^2$; matrix 256×230 ; voxel size, $1.0 \times 1.0 \times \text{mm}^3$; repetition time (TR), 2,300 ms; echo time (TE), 2.98 ms; flip angle, 9°). Scans from the SFVA were acquired on a 1.5 Tesla Siemens Magnetom VISION system (Siemens, Iselin, NJ) equipped with a quadrature head coil using an MPRAGE sequence (164 coronal slices; slice thickness, 1.5mm; FOV, $256 \times 256 \text{ mm}^2$; matrix, 256×256 ; voxel size, $1.0 \times 1.5 \times 1.0 \text{ mm}^3$, TR, 10 ms; TE, 4 ms; flip angle, 15°).

SPM Image Processing—All T1-weighted MR images were visually inspected for movement and other artifacts prior to analysis. The images were then bias corrected and segmented into GM, white matter (WM), and cerebrospinal fluid (CSF) using Statistical Parametric Mapping (SPM) 12³¹ under the developers' recommended (default) preprocessing parameters. Each segmented image was manually validated for accurate segmentation. The segmented scans were then warped to a cohort-specific template using the Diffeomorphic Anatomical Registration Through Exponentiated Lie Algebra (DARTEL) tool box³². DARTEL-processed GM images were then linearly co-registered to the MNI152 template and smoothed using an 8 mm FWHM kernel. All Chinese and white participants' images were preprocessed together to minimize batch effects due to differences in morphology.

Freesurfer (FS) Image Processing—All images were processed using Freesurfer version 5.3 using the default settings. Prior to skull stripping, all images were bias corrected³³. The Freesurfer pipeline estimates white matter and cortical surfaces, which enables the estimation of cortical thickness across multiple brain regions³⁴. The standard Freesurfer segmentation is also able to estimate volumes of subcortical regions³⁵. Following initial surface estimations, we inspected each patient's segmentation visually and corrected errors by hand. Of the 94 individuals included in the study, 4 Freesurfer segmentations did not pass quality control after two separate attempts to segment and edit the image, leaving 90 individuals with Freesurfer data available for analysis. Following segmentation, we extracted regional cortical thicknesses from participant images using the Desikan-Killiany cortical atlas³⁶. In brief, the atlas contains 34 gyally-defined cortical regions of interest (ROIs). Given the strong relationship between atrophy of the hippocampus and amygdala in AD³⁷, we also extracted volumes for these subcortical ROIs. We used ROIs from both hemispheres, yielding a total of 72 ROIs.

Statistical Analyses

Comparing Clinical Characteristics—We first assessed whether clinical impairment varied by diagnosis in Chinese and white study participants across the AD spectrum. We used linear regression to compare CDR-SB scores by diagnosis and ethnic grouping, adjusting for age, sex, and education.

Voxel Based Morphometry (VBM)—All voxel-based testing was conducted using SPM. An aggregate region of AD-associated atrophy was generated using general linear models.

The model was fit to each voxel and tested the relationship between GM tissue density and an individual's CDR-SB score, under the one-tailed assumption that clinical impairment in MCI and AD patients would be associated with lower volume compared to controls. In this analysis, covariates included race (Chinese or white), sex, education, total intracranial volume (TIV), and scan type (1.5T or 3T). Chinese and white individuals were analyzed together to ensure identification of a 'common' set of regions across all study participants that could then be assessed at the individual level. For consideration as a statistically meaningful finding, a cluster of associated voxels was required to be significant at a family-wise error (FWE) corrected p-value ≤ 0.05 and have a cluster size greater than 50 voxels. GM volumes were then extracted from significant clusters and summed to make a composite AD ROI. The composite AD ROI was then used in secondary analyses as the outcome variable in a linear model, with race (Chinese or white), CDR-SB score, age, sex, education, TIV, and scan type (1.5T or 3T) as predictors.

Cortical Thickness Analysis—Using linear regression models, we assessed whether regional cortical thickness was associated with CDR-SB score across all Desikan-Killiany ROIs as well as hippocampus and amygdala. Covariates included race (Chinese or white), sex, education, and scan type (1.5T or 3T). TIV was included as a covariate when analyzing hippocampal and amygdalar volumes. For these analyses, we defined statistical significance using the Bonferroni technique at a p-value of 0.00069 ($0.05 / 72$ ROIs).

The top regions of atrophy identified in regional analyses were utilized in secondary analyses with the same covariates.

All statistical analyses were performed using R unless otherwise stated.

RESULTS

Cohort Composition

Ninety-four individuals were eligible for this study and were seen between 2005 and 2016 (Table 1). Nearly half of the participants ($n=46$) were white and the remainder were Chinese ($n=48$). Comparisons by race in the overall cohort, CN group, and AD group revealed no significant differences in age, education, sex, CDR-SB score, CDR total score, MMSE score, or scan type (all $p>0.05$). In the MCI group, there were no significant differences by race in age, education, sex, CDR score, and MMSE score (Table 1), but there were significant differences between Chinese and whites in CDR-SB score and scan type.

Chinese and white cohorts are clinically similar in MCI and AD

We first tested whether there were clinical differences between MCI and AD versus CN controls in the Chinese and white cohorts. There was a significant difference in clinical severity (measured by CDR-SB score) by diagnostic grouping in both the white (MCI $p=2.83\times 10^{-7}$; AD $p<2\times 10^{-16}$) and Chinese (MCI $p=0.19$; AD $p=3.17\times 10^{-16}$) cohorts as well as when the two races were combined together (MCI $p=1.41\times 10^{-4}$; AD $p<2\times 10^{-16}$). CDR-SB scores were not associated with age, race, sex, or education (see table Supplemental Digital Content 2, Figure 1).

VBM across the AD spectrum identifies common areas of regional atrophy in Chinese and white individuals

In the VBM analysis of all participants contrasting CN to MCI and AD individuals, three clusters were significant after multiple testing correction. As expected, both the right and left hippocampus, amygdala, and parahippocampal area showed significant atrophy along with the right middle temporal lobe ($p_{FWE} < 0.05$) (Figure 2A).

As expected, there were significant volumetric differences in the combined AD ROI by CDR-SB score in the Chinese ($p = 8.21 \times 10^{-5}$) and white ($p = 4.00 \times 10^{-5}$) cohorts as well as the combined cohort ($p = 4.73 \times 10^{-10}$) (Figure 2A). Volume within the AD ROI was also significantly associated with age, education, and TIV. Race, sex, and scan type were not significant predictors of volume (see Table, Supplemental Digital Content 2).

Cortical thickness analysis across the AD spectrum identifies common areas of regional atrophy in Chinese and white individuals

In the regional cortical thickness analysis across all participants comparing CN to MCI and AD, 19 regions were significantly associated with CDR-SB score ($p < 0.00069$). As expected, and consistent with the VBM analysis, both the right and left hippocampus as well as amygdalar volumes were associated with CDR-SB scores. Additionally, the bilateral entorhinal, middle temporal, paracentral gyrus, precuneus, and supramarginal thicknesses as well as left banks of the superior temporal sulcus, right caudal middle frontal, right inferior parietal, right precentral, right superior frontal, and right superior parietal thicknesses were associated with CDR-SB scores (Figure 2B). Full analysis results for all 72 ROI are available in Supplemental Digital Content 3.

We next tested whether Freesurfer-derived hippocampal volumes and entorhinal cortex thicknesses varied by diagnosis, as these regions had the most significant bilateral volumetric and thickness findings, respectively. Hippocampal volumes were significantly associated with CDR-SB score in the Chinese (right $p = 7.26 \times 10^{-5}$; left $p = 2.21 \times 10^{-5}$) and white cohorts (right $p = 0.005$; left $p = 0.006$), as well as the combined cohort (see Table, Supplemental Digital Content 2). Similarly, entorhinal cortex thicknesses were significantly associated with CDR-SB score in the Chinese (right $p = 6.22 \times 10^{-5}$; left $p = 0.004$) and white cohorts (right $p = 0.01$; left $p = 0.001$), as well as the combined cohort (see Table, Supplemental Digital Content 2). However, race was not a significant predictor of hippocampal volume or entorhinal cortex thickness after multiple testing correction ($p > 0.00069$).

Although not significant after multiple testing, Chinese participants showed greater entorhinal thicknesses than white participants (raw p -value < 0.05). To specifically test whether GM volume loss varied as a function of clinical progression and race, we added an interaction term to the VBM and FS models and found that volume loss did not significantly vary between races across the AD spectrum for the VBM-generated AD ROI or top Freesurfer regions (Table 2, Figure 3).

DISCUSSION

In this study, we compared GM loss with clinical measures of disease severity between demographically similar groups of Chinese and white CN, MCI and AD participants living in the San Francisco area. We found no significant differences in disease severity as measured by CDR-SB score. Similarly, using two independent image processing pipelines, we found no significant volumetric or cortical thickness differences in GM atrophy between Chinese and white participants across the AD spectrum. The correlation between volume loss and degree of clinical severity was the same in both Chinese and white individuals. Taken together, this suggests that Chinese and white AD patients, when matched demographically, look neuroanatomically and clinically similar on normalized measures of GM atrophy and clinical severity.

The incidence of AD in European populations was recently estimated at 11.08 cases per 1,000 person-years, and was 6.25 cases per 1,000 person-years in the most populated country in the world, China^{38,39}. While some cohort studies suggest that age-specific incidence of dementia has decreased in the past 20 years in European populations⁴⁰ and in the United States⁴¹, the prevalence and incidence of dementia has increased in China due to a growing elderly population, rising dementia awareness, and growing research in less developed regions³⁸. This increasing prevalence and incidence of dementia could have important ramifications given the accelerated growth of the oldest segments of the Chinese population⁴². Differences in cultural stigma, prediction of dementia risk, early and accurate diagnosis, and therapeutic treatment and response across populations all likely contribute to these differences in AD epidemiology in diverse groups. Despite the high numbers of Chinese-Americans in the U.S., this population has not been included in many studies of AD and other dementias. Taken together, these factors can lead to challenges in predicting the severity of cognitive impairment as well as providing appropriate diagnoses and treatments for people with AD, particularly in underserved groups and ethnic populations in developing countries.

In our study, CDR-SB scores were significantly associated with diagnosis and age, consistent with prior studies in different Asian populations^{43,44} and providing further evidence that the CDR-SB is a reliable as well as valid classification tool to evaluate severity of dementia and is sensitive to cognitive decline in diverse populations⁴⁵. Our neuroimaging analyses identified several brain regions that showed GM volume loss in both Chinese and white AD patients. As expected, these regions included many regions classically associated with AD, such as the hippocampus, amygdala, parahippocampal regions, and temporal cortex^{46,47}.

Previous studies suggest that MRI volumetric data can aid clinical evaluation as a complementary diagnostic tool⁴⁸. Indeed, two studies conducted separately in Chinese and white cohorts have examined atrophy patterns in AD and MCI patients and consistently demonstrated that hippocampal volume derived from MRI scans predicts future cognitive decline and conversion to AD among individuals with MCI^{46,49}. GM volume loss in medial temporal lobe, hippocampus, and parahippocampus was also shown to accurately distinguish AD from normal controls and other neurodegenerative diseases in Chinese individuals^{50,51},

as well as other populations^{48,49}. Our results suggest that volume loss in the AD-associated ROIs we identified using both VBM and Freesurfer — which included hippocampal and parahippocampal structures — could be a good predictor for clinical progression from MCI to AD in diverse populations.

Our data suggests that Chinese and white individuals do not show significant brain atrophy differences in the context of AD neurodegeneration. In our study, both cohorts showed similar patterns of clinical and neuroanatomical vulnerability, suggesting similar pathophysiological processes occur in both ethnic groups. We replicated this finding using two independent data processing pipelines (after normalizing regional volume by TIV, scan type, and demographic variables). While not definitively conclusive, our study has important implications for studies evaluating AD progression with measures such as CDR-SB score or hippocampal atrophy. Our findings suggest that clinical and neuroimaging data from Chinese and white individuals can be combined and analyzed as a single group without confounding due to differential disease effects by racial groupings. Despite differences in population-based genetic background, this suggests that the patterns of neurodegeneration are similar in both groups, supporting a common ‘trajectory’ of conversion from healthy aging to MCI and AD that may be amenable to similar types of interventions.

Our cohorts were clinically well characterized and we used this information to minimize bias across cohorts. This included comparing baseline assessments of CN controls, MCI, and AD patients and matching Chinese and white individuals within each diagnostic group by age, sex, education, and scan type to the greatest extent possible. To ensure our results were not driven by a single image processing modality, we performed our analyses using both VBM and Freesurfer. These two analytically different pipelines converged upon the same findings, suggesting our results are not driven by technical artifacts or biased by an individual pipeline’s reference atlas. In our study, scanner type was not associated with brain morphology, although variation introduced by the difference in distribution of the two scanner types across diagnostic and population subgroups cannot be wholly discounted. This is important for clinical practice and research when using 1.5T MRI⁵³.

The main caveat of our study is the limited sample size, raising the possibility that more varied environmental and/or genetic influences could mask effects in other populations in China or other European countries. Although the number of white research participants at our center is larger, we specifically designed our study to directly match all available Chinese participants with demographically similar white participants. One of the contributors to this limited sample size is the low participation rate of ethnic minorities in clinical research and intervention trials^{54,55}. This study is also limited by the comparatively small number of cognitive assessments for participants who chose to test in Chinese, some of which remain to be fully validated for neuroanatomic correlates in Mandarin and Cantonese-speaking individuals. It is possible that more detailed cognitive testing could reveal subtle differences between study groups. Furthermore, we did not have sufficient data to account for potential modifying contributions of lifestyle and/or environment across the Chinese-American cohort. Future work will benefit from broader samples, additional language-validated cognitive testing, and inclusion of systematic measures of lifestyle and

environmental factors that may modify trajectories in cognitive health and neurodegeneration.

Our study provides cross-sectional evidence that Chinese and white individuals across the AD spectrum show the same disease-associated patterns of neuroimaging and clinical severity. This research highlights the importance of implementing strategies to improve diversity in clinical research and therapeutic trials. Gaining a better understanding of ethnically diverse communities in AD clinical research will generate rich information about contributions of genetic variation, environment, diet, and culture among ethnic groups that may differentially contribute to AD risk and may influence future treatment decisions as therapeutic interventions become available. Evaluation of diverse aging populations will be critical for effectively addressing public health policy issues related to AD and in recruitment for clinical trials, and could ultimately inform the design and interpretation of intervention trials in multi-ethnic cohorts.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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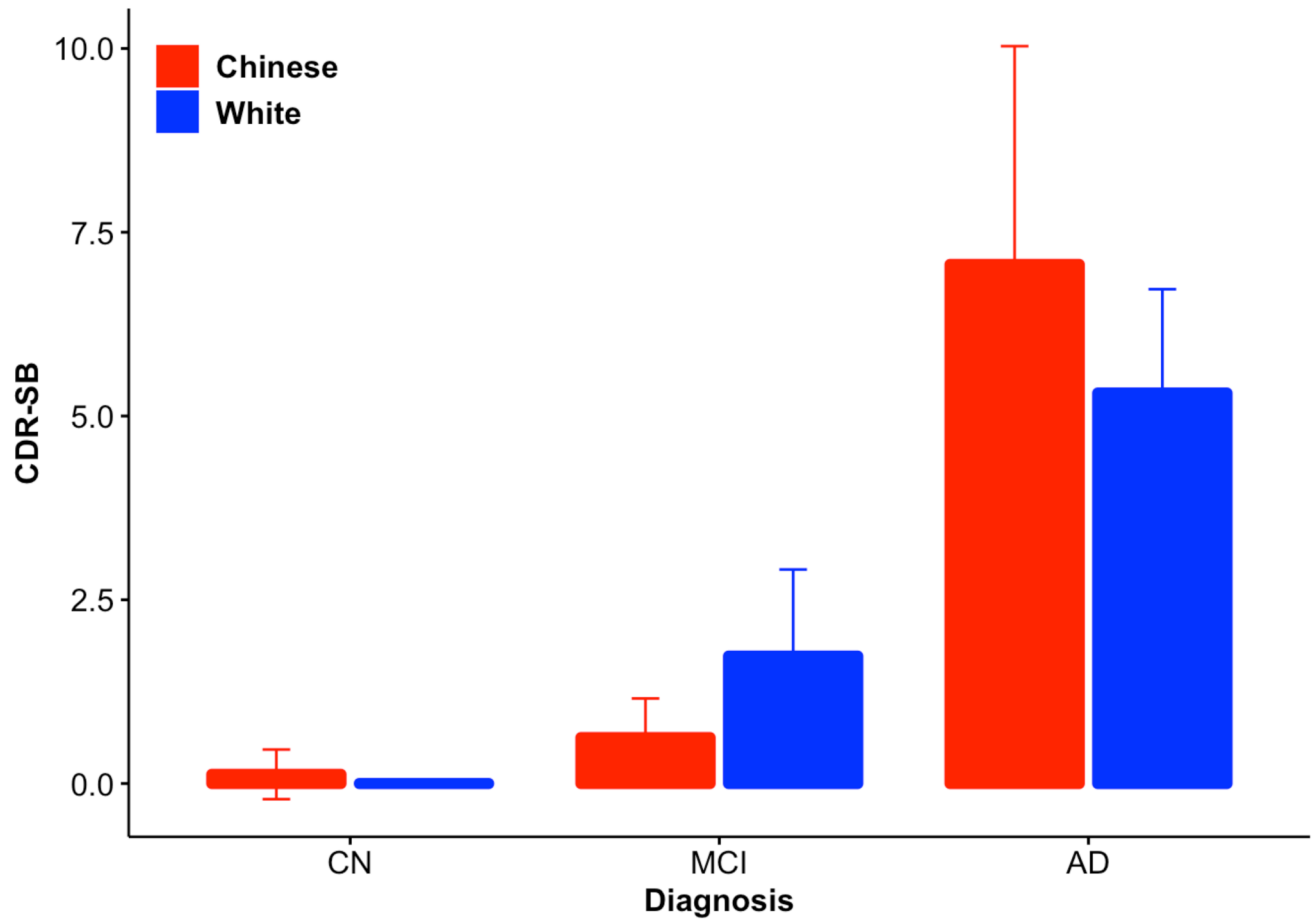


Figure 1: CDR-SB score by race and diagnosis.

Clinical Dementia Rating Sum of Boxes (CDR-SB) scores were significantly different across diagnostic groups (see Table, Supplemental Digital Content 1), but showed no significant difference across races ($p=0.96$). The plotted points are best linear unbiased predictions (BLUPs). See Supplemental Digital Content 1 and the text for details. CN – Control; MCI – Mild Cognitive Impairment; AD – Alzheimer's disease.

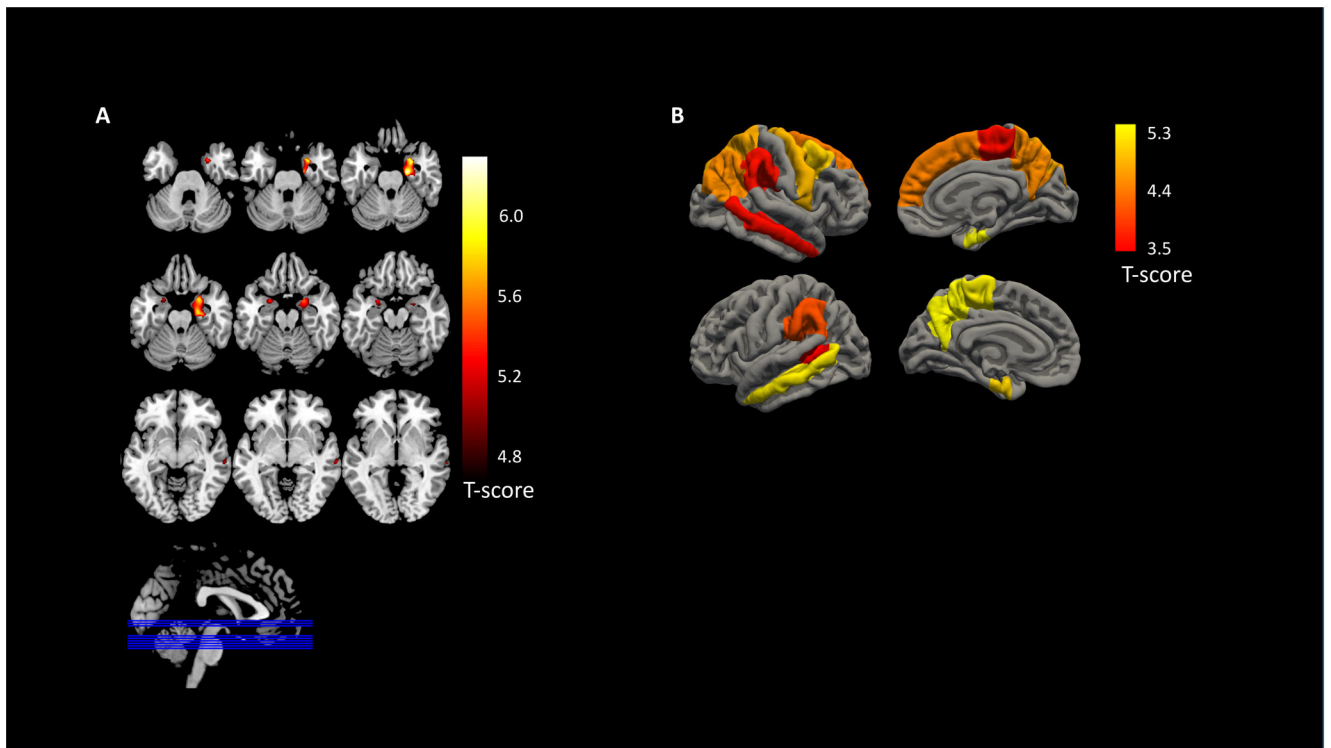


Figure 2: AD-associated atrophy measured by VBM and FS.

Atrophy varied significantly across the AD spectrum (measured by Clinical Dementia Rating Sum of Boxes (CDR-SB) scores). A) The voxel-based morphometry (VBM) regions of interest (ROI) included the right and left hippocampus, amygdala, and parahippocampal areas along with the right middle temporal lobe. B) The Freesurfer (FS)-based AD ROIs included the bilateral hippocampi and amygdalae volumes along with bilateral entorhinal, middle temporal, paracentral gyrus, precuneus, and supramarginal thicknesses as well as left banks of the superior temporal sulcus, right caudal middle frontal, right inferior parietal, right precentral, right superior frontal, and right superior parietal thicknesses.

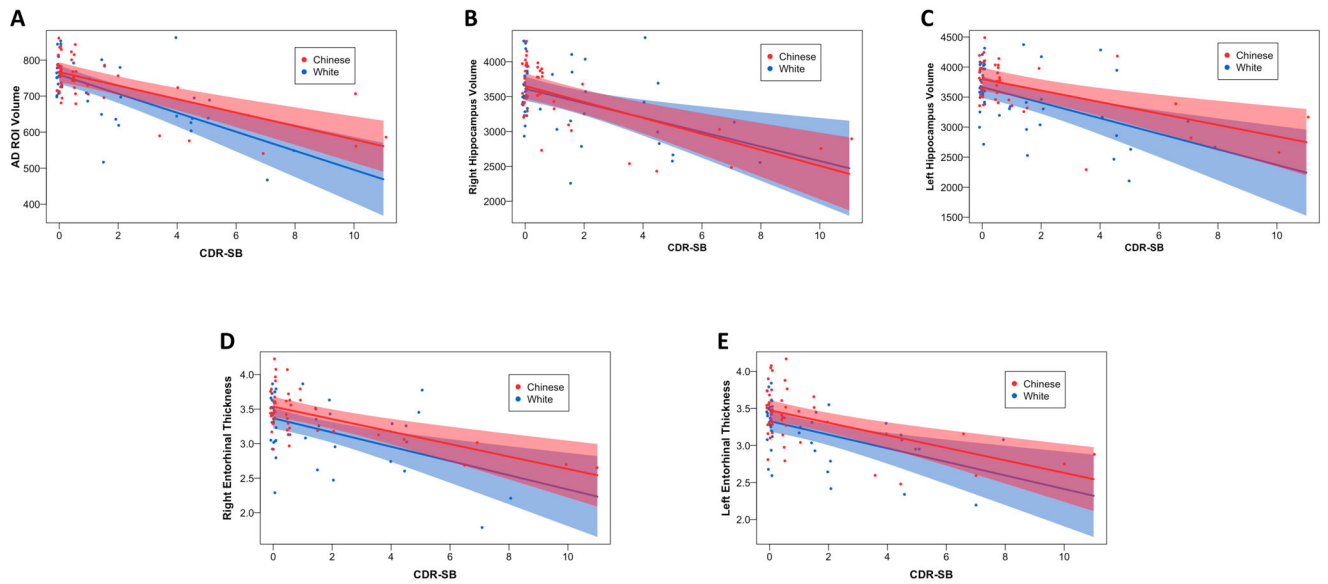


Figure 3: No difference in AD-associated volume loss between Chinese and white individuals across the AD spectrum.

Interaction analyses of Clinical Dementia Rating Sum of Boxes (CDR-SB) score by race versus volume or thickness for the top AD regions identified by voxel-based morphometry (VBM) and Freesurfer (FS) are shown. Volume was used for the VBM region of interest (ROI) along with FS hippocampal measurements (A-C) while thickness was used for the FS entorhinal cortex measurements (D-E). There was no significant association between AD-associated volume loss and race across the AD spectrum. The plotted points are partial residuals with 95% confidence bands provided in shading. See Table 2 for details.

Table 1.

Cohort description by ethnic and diagnostic groupings

	Overall Cohort			Cognitively Normal			Mild Cognitive Impairment			Alzheimer's Disease		
	White	Chinese	p-value	White	Chinese	p-value	White	Chinese	p-value	White	Chinese	p-value
N	46	48		23	24		15	16		8	8	
Age (Mean ± SD)	69 ± 9.2	69 ± 8.5	0.77	69 ± 8.7	69 ± 8.1	0.99	67 ± 9.5	66 ± 9.0	0.83	75.8 ± 8.2	74.5 ± 6.5	0.74
Female Sex (N; %)	25 (54.3%)	25 (52.1%)	0.99	13 (56.5%)	13 (54.2%)	1	9 (60%)	9 (56.2%)	1	3 (37.5%)	3 (37.5%)	1
CDR-SB Score (Mean ± SD)	1.5 ± 2.1	1.5 ± 2.8	0.94	0 ± 0	0.12 ± 0.3	0.08	1.7 ± 1.2	0.6 ± 0.5	0.002	5.31 ± 1.4	7.1 ± 3.0	0.15
CDR Score (Mean ± SD)	0.3 ± 0.4	0.4 ± 0.5	0.53	0 ± 0	0.06 ± 0.2	0.08	0.5 ± 0.1	0.4 ± 0.2	0.18	0.9 ± 0.2	1.3 ± 0.6	0.11
MMSE (Mean ± SD)	27.1 ± 3.7	26.9 ± 3.6	0.78	29.3 ± 0.9	28.9 ± 1.1	0.20	27.5 ± 1.6	27.5 ± 1.4	0.95	20.0 ± 2.5	19.9 ± 2.6	0.92
Education (Years; Mean ± SD)	15.2 ± 2.5	14.9 ± 3.3	0.52	15.3 ± 1.9	15.5 ± 2.9	0.75	15.5 ± 3.2	14.9 ± 2.8	0.59	14.6 ± 2.6	12.8 ± 4.5	0.33
3T MRI Scan (N; %)	36 (78.3%)	45 (93.8%)	0.06	22 (95.7%)	23 (95.8%)	1	9 (60%)	16 (100%)	0.02	5 (62.5%)	6 (75%)	1

Cohort demographic and clinical characteristics. MMSE: Mini-mental state examination; CDR-SB: Clinical Dementia Rating – Sum of Boxes score; CDR: Clinical Dementia Rating; T: Tesla; SD: Standard deviation.

Table 2.

Interaction analyses of AD regions of interest across the AD spectrum

	VBM ROI		FS Right Hippocampus		FS Left Hippocampus		FS Right Entorhinal		FS Left Entorhinal	
	$\beta \pm SE$	p-value	$\beta \pm SE$	p-value	$\beta \pm SE$	p-value	$\beta \pm SE$	p-value	$\beta \pm SE$	p-value
Age	-3.34 ± 0.79	6.33E-05	-37.17 ± 5.38	1.03E-09	-34.92 ± 5.67	2.64E-08	-0.02 ± 0.01	0.001	-0.01 ± 0.004	0.24
Sex	18.05 ± 17.49	0.30	0.76 ± 119.42	0.99	285.84 ± 125.69	0.03	0.02 ± 0.08	0.77	0.10 ± 0.08	0.21
Education	-6.83 ± 2.38	0.005	19.58 ± 18.66	0.30	-8.62 ± 19.63	0.66	-0.03 ± 0.02	0.08	0.001 ± 0.02	0.95
Scan Type	26.05 ± 22.38	0.25	-191.06 ± 151.43	0.21	-272.02 ± 159.375	0.09	0.05 ± 0.13	0.70	0.11 ± 0.12	0.38
TIV	405.45 ± 62.94	6.75E-09	6.38E-04 ± 3.667E-04	0.86	0.001 ± 3.860E-04	0.002	-	-	-	-
CDR-SB	-26.390 ± 4.94	7.49E-07	-103.56 ± 33.13	0.002	-129.48 ± 34.87	3.74E-04	-0.10 ± 0.03	5.34E-04	-0.10 ± 0.03	0.001
Race	7.11 ± 16.06	0.66	39.50 ± 107.07	0.71	132.67 ± 112.69	0.24	0.17 ± 0.09	0.07	0.15 ± 0.09	0.09
CDR-SB X Race	7.70 ± 5.73	0.18	10.91 ± 39.56	0.78	34.13 ± 41.64	0.41	0.01 ± 0.03	0.71	0.01 ± 0.03	0.83

There was no significant difference in AD-associated grey matter volume loss between Chinese and whites after accounting for age, sex, education, TIV, scan type, and race. For the interaction analyses, we used the same five regions of interest generated in our primary volumetric analyses, a voxel-based morphometry (VBM) aggregate region of interest (ROI) and Freesurfer volumetric measures (bilateral hippocampi and entorhinal cortices). ROI: regions of interest; TIV: total intracranial volume; CDR-SB: Clinical Dementia Rating – Sum of Boxes score; VBM: Voxel-based morphometry; FS: Freesurfer; β : Beta coefficient from regression model; SE: Standard error.