

Preclinical Alzheimer's disease has been characterized by the presence of normal cognitive function and abnormal levels of cerebrospinal fluid (CSF) biomarkers. The preclinical stage is typically followed by mild cognitive impairment, which progresses to clinically apparent dementia in some persons. Neuropathologic abnormalities and changes in biomarker levels can begin 15 to 20 years before clinical manifestations of Alzheimer's disease.

Changes in CSF biomarkers such as levels of amyloid-beta (A β), total tau, phosphorylated tau 181, and neurofilament light chain (NfL) have been indicators in preclinical Alzheimer's disease⁸ that become abnormal sequentially rather than simultaneously.⁹ Some previous studies of the sequential appearance of changes in CSF biomarkers have involved persons with autosomal dominant Alzheimer's disease, which accounts for only a small proportion of Alzheimer's disease cases, and these studies have typically used an estimated number of years before the onset of Alzheimer's disease symptoms to define the timeline of biomarker changes.⁴⁰⁻¹⁴

Determination of the sequence of these changes in sporadic Alzheimer's disease is challenging because a person's clinical course, beginning with normal cognition and progressing to Alzheimer's disease, cannot be predicted. Most studies regarding biomarkers in sporadic Alzheimer's disease have been cross-sectional and may not have reflected alterations of biomarkers over the period from a normal cognitive state to Alzheimer's disease. Longitudinal studies, such as the Alzheimer's Disease Neuroimaging Initiative, have advanced our understanding of preclinical sporadic Alzheimer's disease by exploring these biomarker changes.⁴⁶⁻¹⁹

However, a limitation of these studies has been the underrepresentation of Asian populations, which potentially has limited the generalizability of the results. In addition, the relatively short follow-up periods in previous studies do not reflect the lengthy trajectory over decades of biomarker alterations leading to the onset of Alzheimer's disease. We examined a cohort of participants from one of the nested studies in the China Cognition and Aging Study (COAST) with a goal of estimating the trajectory of changes in several CSF biomarkers in 6.4 (w) 12 (t) to 6.2 1 > 459 (i) - 7.8 (s) s i

tions, including lumbar puncture, and follow-up. range from 0 to 25, with higher scores indicating better memory abilities. The protocol was approved by the ethics committee of Xuanwu Hospital, Capital Medical University, and the study was conducted in accordance with the principles of the Declaration of Helsinki. Participants were not compensated for participating in the study. The sponsors had no role in the study design; collection, analysis, interpretation of the data; or the writing of the report.

Nested Case–Control Approach

Our study required that participants be observed for more than 15 years but not more than 20 years. They would undergo at least three assessments during which the diagnosis was made, and an intermediate follow-up visit between the two. The CDR–SB scores were independently assessed by physicians who were unaware of other cognitive tests in 2000, of whom 32,061 were eligible and enrolled in the current substudy. Of those enrolled, 30,272 were included (6435 discontinued the study, 3172 were unreachable, 10,470 had died, 2759 were cognitively impaired, 4514 were excluded for health-related reasons, 1228 had fewer than three assessments, and 1694 were excluded for other reasons), leaving 1789 participants enrolled.

At the last follow-up, 695 participants received a diagnosis of Alzheimer's disease and 1094 remained cognitively normal (as assessed by tests as described below). After propensity-score matching on the basis of age, sex, and education level, 648 (93.2%) of the participants with Alzheimer's disease were successfully matched in a 1:1 ratio with participants who remained cognitively normal at the last follow-up, and these two groups form the basis for the current report.

Diagnosis of Cognitive Status

The cognitive status of participants was determined at baseline and at each follow-up with the use of three scales. Participants were considered to have no cognitive impairment if they had a score of 27 or higher on the Mini–Mental State Examination (MMSE; range, 0 to 30, with higher scores indicating better performance). Scores of 12 or higher on the Logical Memory Test (LMT), a modification of the episodic memory section of the Wechsler Memory Scale–Revised (scores

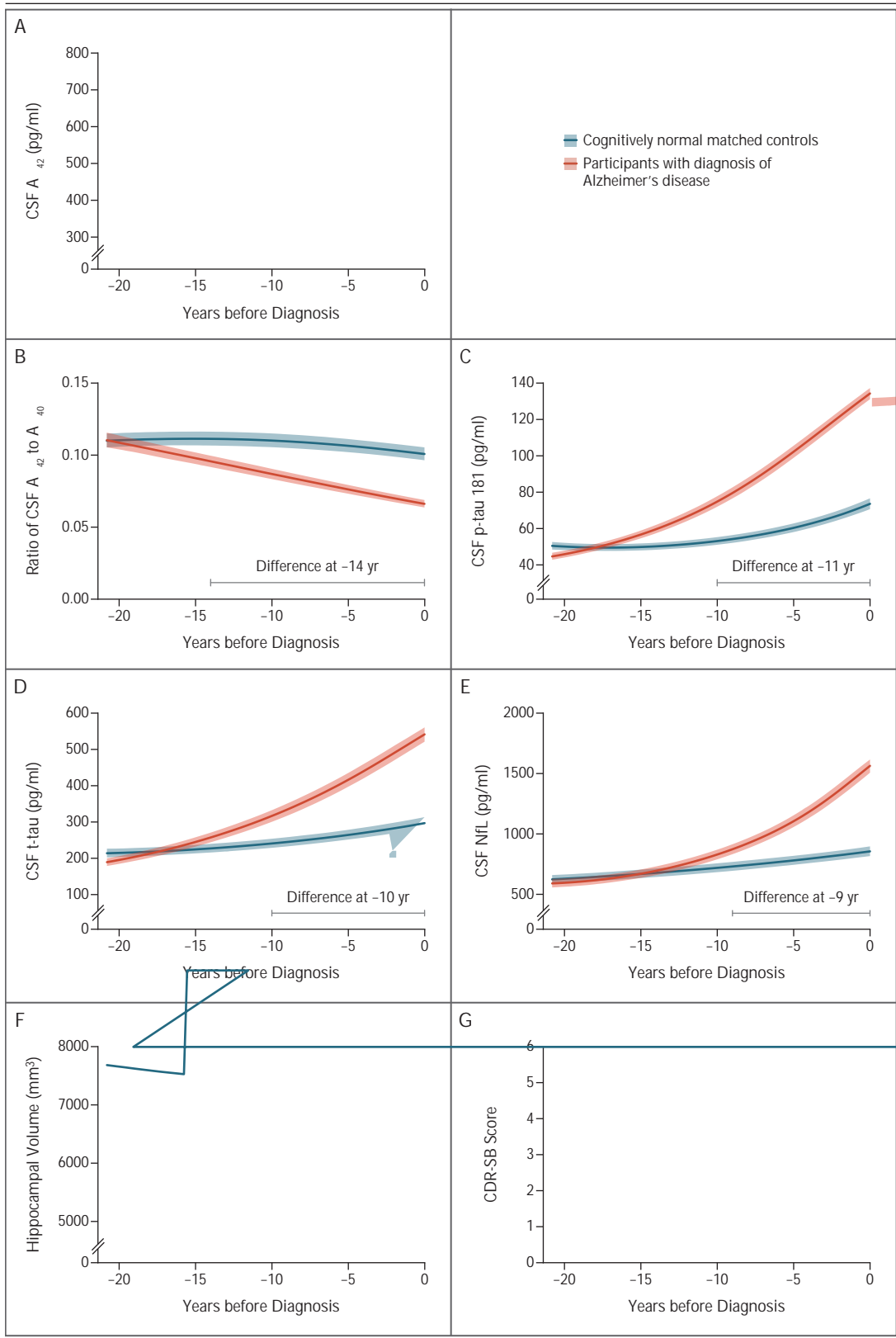
Table 1.Characteristic of the Participants at Baseline.*

Variable	Cognitively Normal (N=648)	Alzheimer's Disease (N=648)
Age — yr	61.3±4.1	61.2±4.1
Sex — no. (%)		
Male	328 (50.6)	327 (50.5)
Female	320 (49.4)	321 (49.5)
Education, level and total yr — no. of participants (%)		
Primary school, 6 yr	25 (3.9)	28 (4.3)
Middle school, 7–9 yr	28 (4.3)	26 (4.0)
High school, 10–12 yr	212 (32.7)	210 (32.4)
University, 13–17 yr	334 (51.5)	335 (51.7)
Postgraduate, 18–21 yr	49 (7.6)	49 (7.6)
APOE status — no. (%)		
Noncarrier	516 (79.6)	407 (62.8)
Carrier	132 (20.4)	241 (37.2)
Heterozygous	121 (18.7)	183 (28.2)
Homozygous	11 (1.7)	58 (9.0)
Cognitive score†		
MMSE	29.5±1.0	29.4±1.2
CDR-SB	0	0
LMT	16.8±0.6	16.8±0.7
Biomarker values		
Ratio of A ₄₂ to A ₄₀	0.1±0.0	0.1±0.0
Total tau — pg/ml	219.2±52.6	214.6±41.2
Phosphorylated tau 181 — pg/ml	48.8±9.9	48.4±7.8
Neurofilament light chain — pg/ml	633.9±139.3	645.4±140.4
Hippocampal volume — mm ³	7708.3±621.8	7683.0±645.5

* Plus–minus values are means ±SD. A denotes amyloid-beta, and APOE apolipoprotein E gene.

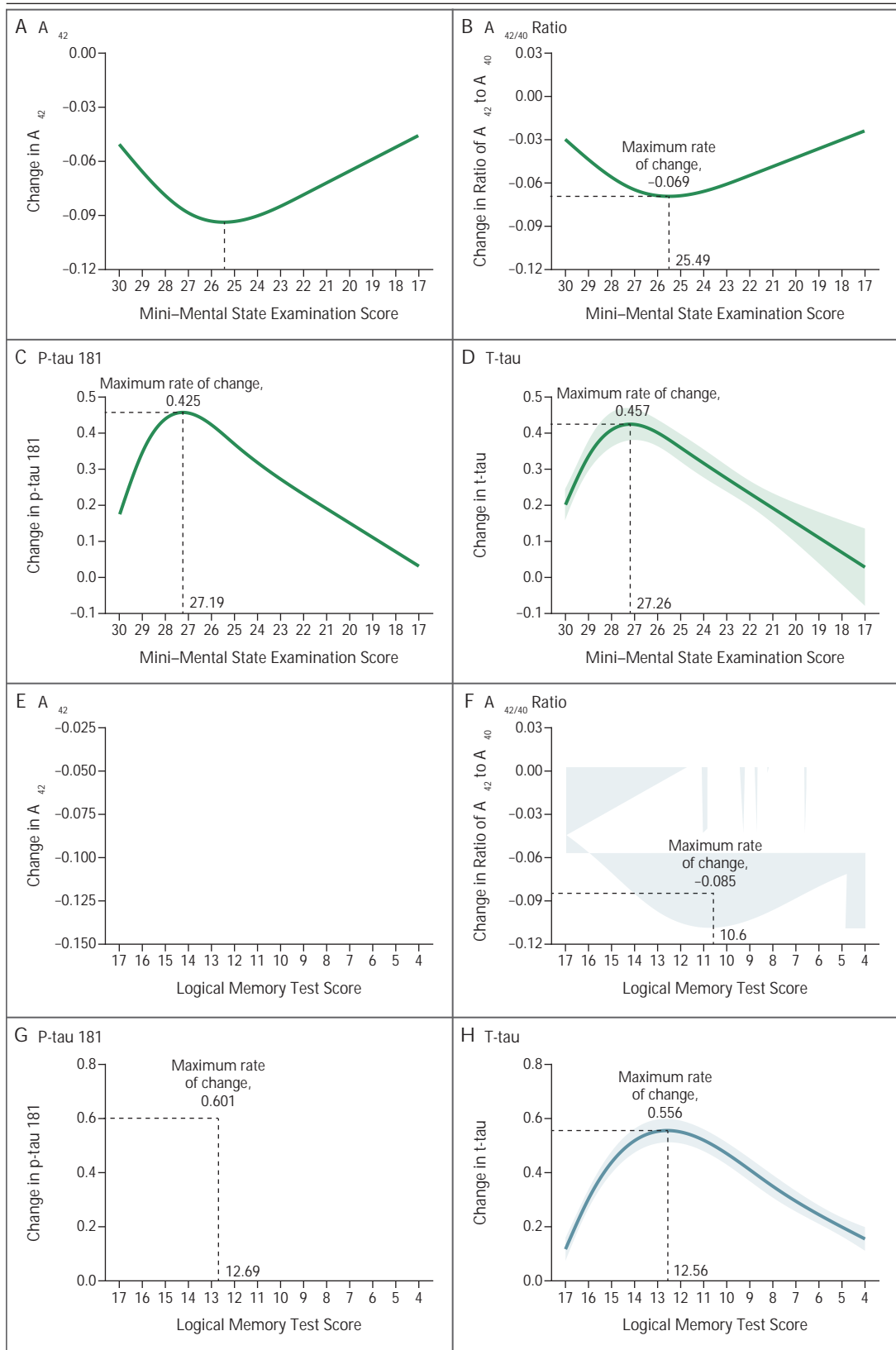
† Scores on the Mini–Mental State Examination (MMSE) range from 0 to 30, with higher scores indicating better performance. Totals for the Clinical Dementia Rating–Sum of Boxes (CDR-SB) range from 0 to 18, with higher values indicating greater cognitive impairment. Scores for the Logical Memory Test (LMT) range from 0 to 25, with higher scores indicating better memory abilities.

Brain Volumetric Imaging participants. Left and right hippocampal volumes were summed to assess the degree of brain atrophy. Structural magnetic resonance imaging of the brain was performed with 3.0-T scanners (Siemens) with a 20-channel phased-array head-neck coil. The absolute volume of each region of interest was determined with the use of FreeSurfer software, version 5.3.0 (Table S2). The relationship between hippocampal volume and diagnosis was assessed with the use of par Surfer software, version 5.3.0 (Table S2). The relationship between hippocampal volume and diagnosis was assessed with the use of par Surfer software, version 5.3.0 (Table S2). The relationship between hippocampal volume and diagnosis was assessed with the use of par Surfer software, version 5.3.0 (Table S2). The relationship between hippocampal volume and diagnosis was assessed with the use of par Surfer software, version 5.3.0 (Table S2).



We used the R software, version 4.3.1, *lcmm* package (R Foundation for Statistical Computing) to establish latent-class mixed models for the estimation of the trajectories of each biomarker over time (see the Supplementary Methods-section).^{26,27} These models incorporated quadratic functions of retrospective time and were adjusted for case–control status, covariates (e.g., age, sex, education level, and *POE* status), and their interactions with time and time squared. Within-participant correlations were accounted for by correlated random intercepts and slopes of time and time squares. Spline functions were integrated into the models to capture potential variations in biomarker trajectories over time. The final models with the optimal number of knots were determined with the use of the Akaike and Bayesian information criteria.²⁸

Using the R software *mvtnorm* package for Wald tests, we evaluated the differences in biomarkers between participants with Alzheimer's disease and cognitively normal participants for



matched cognitively normal group. The study included only Han Chinese persons. Within both groups, men slightly outnumbered women. Baseline CSF biomarker levels, cognitive scores, and hippocampal volumes were similar in the two groups. Participants in whom Alzheimer's disease ultimately developed were more likely than their matched controls to be carriers of the $\epsilon 4$ allele (37.2% vs. 20.4%). Overall, the participants had a level of education that slightly surpassed the educational norms for the general population of older adults in China. The representative

ease; in the control group, the rate of change appeared to have flatter trajectories (Fig. S5).



Biomarker Changes Preceding Alzheimer's Disease

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