

BMJ Open Sino Longitudinal Study on Cognitive Decline (SILCODE): protocol for a Chinese longitudinal observational study to develop risk prediction models of conversion to mild cognitive impairment in individuals with subjective cognitive decline

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ABSTRACT

Introduction Understanding the biological mechanism of subjective cognitive decline (SCD) in preclinical Alzheimer's disease (AD) and identifying those who will soon convert to mild cognitive impairment (MCI) are critical for developing appropriate strategies for early diagnosis and intervention of AD. We present the study protocol of the Sino Longitudinal Study on Cognitive Decline (SILCODE), a longitudinal observational study focusing on SCD in the context of AD.

Methods and analysis Within SILCODE, approximately 800 subjects with SCD who are between 50 and 79 years old will be recruited through standardised public advertisements or memory clinics. They will undergo extensive assessment, including clinical and neuropsychological assessments, blood sample collection for plasma beta-amyloid and ApoE genotype, urine samples collection for AD7c-NTP, and multimodal MRI scans (structural MRI, diffusion tensor imaging, resting-state functional MRI and optional task-based functional MRI) as well as optional glucose metabolism and amyloid positron emission tomography. Subjects will be contacted by telephone every 3 months and interviewed, on average, every 15 months for 5 years. The study endpoint is the development of mild cognitive impairment or dementia. Jak & Bondi's actuarial neuropsychological method will be used for diagnosis of MCI. The least absolute shrinkage and selection operator logistic regression model followed by the sub-distribution hazard function model with death as a competing risk will be constructed to establish risk prediction models.

Ethics and dissemination The ethics committee of the Xuanwu Hospital of Capital Medical University has approved this study protocol (ID: [2017]046). The results will be published in peer-reviewed journals and presented at national and international scientific conferences.

Trial registration number NCT03370744; Pre-results.

INTRODUCTION

Alzheimer's disease (AD) is the most common neurodegenerative disease and one

Strengths and limitations of this study

- A strength of this study is the use of multiple markers (clinical, blood, urine and imaging markers) to establish models predicting conversion from subjective cognitive decline (SCD) to mild cognitive impairment, which will provide references for clinical applications of SCD.
- This longitudinal study could enrich the understanding of SCD by examining a large-scale Chinese population because cultural factors have an impact on SCD.
- The relatively short follow-up duration (5 years) of this study allows longitudinal data to be analysed only in an exploratory way.
- The study results will be limited by the research environment in that individuals will be recruited from memory clinics and volunteer samples, without community-based or population-based samples.

of the greatest healthcare challenges of the 21st century.¹ It has been revealed that pathophysiology starts many years before the onset of clinical symptoms.² Several recent clinical trials of β -amyloid (A β)-lowering therapies in the mild or moderate dementia stage and even mild cognitive impairment stage have failed,^{3,4} further encouraging researchers to shift their focus to the preclinical stage of AD as a target for new treatments.⁵

Subjective cognitive decline (SCD) refers to self-perceived cognitive decline relative to a previously normal status, without impaired performance on standardised neuropsychological tests.^{6,7} Evidence from recent studies suggests that SCD may be one of the earliest symptomatic manifestations of AD.^{6,8} Previous

recall and orientation) will be administered during the call. Participants expressing significant cognitive decline on the telephone will be interviewed ahead of time for the next follow-up. During the follow-up visits with interviews, evaluations including clinical and neuropsychological examinations, blood tests, urine tests and multimodal MRI as well as optional glucose metabolism PET and t-fMRI will be conducted, on average, every 15 months. At each follow-up visit, diagnoses will be re-evaluated under the supervision of two neurologists. [Figure 1](#) provides the overall study outline.

Patient and public involvement

Patients and the public will not be involved in the development of the research question or the design of the study. Patients will not be involved in the recruitment of participants or the conduct of the study. The general results will be disseminated to participants through public education activities.

Sample size calculation

The formula proposed by Hsieh and Lavori²⁹ was used to estimate the sample size in our study with the software PASS V.15.0.5. According to a previous meta-analysis,⁹ the cumulative conversion proportion was 24.4% over a mean follow-up period of 4.1 years. We assumed that the power ($1-\beta$) was 80%, α was 0.05 and the rate of loss to follow-up was 20%. Due to the lack of statistical data on SCD, we calculated the sample size with data reported by Licher *et al*³⁰ and Pinto *et al*³¹ (HR=1.03, SD=8.2, R²=0.127) from cognitively normal elderly subjects for the association between age and cognitive decline. This calculation rendered a total sample size of 762. Considering the need for internal validation, we will recruit at least 800 subjects.

Recruitment of participants

Participants will be consecutively recruited through standardised public advertisements and through referrals from general physicians, memory clinics, or informants. Residents who meet the inclusion criteria will be recruited. Written informed consent will be acquired from each subject before enrolment.

Diagnostic criteria for SCD, MCI and dementia

SCD is defined by the research criteria for pre-MCI (SCD) proposed by Jessen *et al* in 2014⁶: (1) Self-experienced persistent decline in cognitive capacity in comparison with a previously normal status and unrelated to an acute event; (2) Normal age-adjusted, gender-adjusted and education-adjusted performance on standardised cognitive tests.

MCI is defined by an actuarial neuropsychological method proposed by Jak and Bondi.³² Participants are considered to have MCI if they meet any one of the following three criteria and fail to meet the criteria for dementia: (1) having impaired scores (defined as >1 SD below the age-corrected normative means) on both measures in at least one cognitive domain (memory, language, or speed/executive function); (2) having impaired scores in each of the three cognitive domains sampled (memory, language, or speed/executive function); (3) the Functional Activities Questionnaire (FAQ) ≥ 9 . The normative means in our study are from Guo and his team in a Chinese population.^{33–36} Measures and normative means are shown in [table 1](#).

The diagnosis of AD dementia is based on the Diagnostic and Statistical Manual of Mental Disorders, fifth edition, and the diagnostic guidelines for dementia due to AD delivered by the National Institute on Ageing–Alzheimer’s Association workgroups,³⁷ and a total CDR score ≥ 1 .

Inclusion/exclusion criteria

The inclusion criteria are as follows: (1) 50–79 years old, right-handed and Mandarin-speaking subjects; (2) presence of self-perceived continuous cognitive decline compared with previous normal status and unrelated to an acute event; (3) concerns (worries) associated with memory complaint; and (4) failure to meet the criteria for MCI or dementia.

The exclusion criteria are as follows: (1) History of stroke. (2) Current major psychiatric diagnoses such as severe depression and anxiety. When mild or moderate symptoms of psychiatric disorders are suspected, patients will not be excluded.³⁸ They will be evaluated by a psychiatrist to determine whether the psychiatric diagnoses

Table 1 Cognitive domains, neuropsychological tests and normative means

Cognitive domains	Neuropsychological tests	Normative means
Memory	Auditory Verbal Learning Test—long delayed memory ³³	50–59 years old: 5; 60–69 years old: 4; 70–79 years old: 3
	Auditory Verbal Learning Test—recognition ³³	50–59 years old: 20; 60–69 years old: 19; 70–79 years old: 18
Language	Animal Fluency Test ³⁵	Junior middle school: 12; high school: 13; college: 14
	30-item Boston Naming Test ³⁴	Junior middle school: 20; high school: 21; college: 22
Speed/executive	Shape Trail Test Part A ³⁶	50–59 years old: 70s; 60–69 years old: 80s; 70–79 years old: 100s
	Shape Trail Test Part B ³⁶	50–59 years old: 180s; 60–69 years old: 200s; 70–79 years old: 240s

are the cause of SCD. (3) Other neurological conditions that could cause cognitive decline (eg, brain tumours, Parkinson's disease, encephalitis, or epilepsy) rather than AD spectrum disorders. (4) Other diseases that could cause cognitive decline (eg, thyroid dysfunction, severe anaemia, syphilis, or HIV). (5) History of psychosis or congenital mental developmental delay. (6) Cognitive decline caused by traumatic brain injury. (7) Inability to complete the study protocol or presence of contraindications for MRI.

Clinical progression

At each follow-up visit, diagnoses will be re-evaluated under the supervision of two neurologists. The main outcome measure of clinical progression in our study is conversion to MCI or AD dementia. The time to event is the interval between the baseline neuropsychological tests and the date at which the MCI is first diagnosed. For those who bypass the diagnosis of MCI during follow-up, the time between the baseline neuropsychological tests and first diagnosis of dementia is defined as the time to event in our study.

Assessments of subjective cognitive functioning

A semi-structured interview used by the Multicenter German Center for Neurodegenerative Diseases (DZNE)-Longitudinal Cognitive Impairment and Dementia Study (DELCODE) study will be employed in this project to evaluate the details of SCD.¹⁸ It includes information about the onset time, concerns, comparison with others, and the history of visiting a physician not only for the memory domain but also for language, attention and executive control. We will also require informant reports in the evaluation of the self-reported information as suggested by the Subjective Cognitive Decline Initiative Working Group.^{7 39}

For quantitative assessment of the severity of SCD, we will use a newly developed SCD questionnaire including nine reliable SCD items (box 1), which are characterised by different domains, such as global memory functioning

and daily activities ability.^{23 26} In addition, the 12-item ECog²⁷ will also be applied to all participants to measure SCD severity. These questionnaires will not be inclusion criteria and will be considered only for statistical analysis.

Clinical, risk factor and neuropsychological assessments

The clinical assessments include a structured medical history, physical examination, routine laboratory tests (including blood tests, biochemical tests, thyroid function tests, vitamin B12 status, folic acid status, *Treponema pallidum*-specific antibodies and HIV antibodies) and MRI (T1 and FLAIR) for medial temporal atrophy scale and Fazekas score.

For risk factors of AD, all subjects and their informants will provide the following information: family history, vascular risk factors, occupation/retirement, socioeconomic status, cigarettes and alcohol use, and nutrition style. Additionally, the Pittsburgh sleep quality index,⁴⁰ Rapid Eye Movement Sleep Behaviour Disorder Screening Questionnaire⁴¹ and the Epworth Sleepiness Scale⁴² will be used for sleep quality assessment.

The neuropsychological tests are selected for comparability with other similar ongoing studies (eg, DELCODE).¹⁸ To avoid deviations in the evaluation, the neuropsychologists performing the cognitive tests will be trained according to standard guidelines. The kappa coefficient (Fleiss' kappa) will be used to measure the assessment agreement among the neuropsychologists. Neuropsychological results will be double-checked. The neuropsychological battery (table 1) measures different cognitive domains including episodic memory (Auditory Verbal Learning Test-HuaShan version),³³ language (Animal Fluency Test,³⁵ 30-item Boston Naming Test³⁴ and speed/executive function (Shape Trail Test Parts A and B.³⁶ In addition, the SILCODE study will implement tests for global cognition, daily life ability and neuropsychiatric assessment, including MoCA-B,²⁸ the memory and executive screening (MES),⁴³ the CDR, FAQ, the 15-item short form of the Geriatric Depression Scale, the Hamilton Anxiety Scale, the Hamilton Depression Scale and the Neuropsychiatric Inventory.

Box 1 Items included in subjective cognitive decline questionnaire²⁶

Items

- ▶ Do you think you have problems with your memory?
- ▶ Do you have difficulty remembering a conversation from a few days ago?
- ▶ Do you have complaints about your memory in the last 2 years?
- ▶ How often is the following a problem for you: Personal dates?
- ▶ How often is the following a problem for you: Phone numbers you use frequently?
- ▶ On a whole, do you think that you have problems remembering things that you want to do or say?
- ▶ How often is the following a problem for you: Going to the store and forgetting what you wanted to buy?
- ▶ Do you think that your memory is worse than 5 years ago?
- ▶ Do you feel you are forgetting where things were placed?

Blood tests

ApoE genotype

Single nucleotide polymorphisms (SNPs) rs7412 and rs429358 form the ApoE $\epsilon 2/\epsilon 3/\epsilon 4$ haplotype. ApoE will be genotyped using the standard Sanger sequencing method (Sangon, Shanghai, China) using the following primers: 5'-ACGCGGGCACGGCTGTCCAAGG-3' (forward) and 5'-GGCGCTCGCGGATGGCGCTGA-3' (reverse). ApoE will be amplified using the following conditions: 1 cycle of 98°C for 10s, 35 cycles of 72°C for 5s, 1 cycle of 72°C for 5 min. PCR was performed in a final volume of 30 μ l, containing 10 pmol of forward and reverse primers and 50 ng of genomic DNA template, using PrimeSTAR HS DNA Polymerase with GC Buffer (Takara Bio, Kusatsu, Shiga, Japan).

Plasma β -amyloid

Recently, some studies found that plasma amyloid level was correlated with cognitive capacity and cerebrospinal fluid amyloid protein.^{44 45} In particular, when predicting brain amyloid deposition, the accuracy of the composite biomarker (amyloid- β precursor protein 669–711/ $A\beta$ 1–42 and $A\beta$ 1–40/ $A\beta$ 1–42 ratios) could be approximately equal to 90%.⁴⁶ Therefore, we would like to identify whether the plasma β -amyloid is also associated with cognitive decline in SCD individuals. Plasma $A\beta$ will be determined using a commercially available kit, V-PLEX $A\beta$ Peptide Panel 1 (6E10) Kit (K15200E) (Mesoscale Diagnostics, Rockville, Maryland, USA). $A\beta$ peptide levels from each blood draw will be measured in duplicate using the same aliquot.

Urine tests

Ten millilitres of first morning voided urine will be collected from each subject and then immediately refrigerated. An ELISA kit will be used to detect the protein level of AD7c-NTP in the urine specimens.

Imaging protocol

MRI data will be acquired using an integrated simultaneous 3.0 T TOF PET/MR (SIGNA PET/MR, GE Healthcare, Milwaukee, Wisconsin, USA) at the Xuanwu Hospital of Capital Medical University. For each participant, simultaneous PET and 3.0 T MRI data will be obtained. Brain MR images will be inspected by an experienced neuroradiologist.

PET: All participants will be invited for optional [¹⁸F] florbetapir (AV-45) and [¹⁸F] fluorodeoxyglucose (FDG) PET in 3-dimensional acquisition mode. The time duration between the FDG-PET and AV-45 PET is at least 3 days to eliminate the effect of the first tracer. For FDG-PET, each subject will be instructed to fast for at least 6 hours and must have a confirmed serum glucose level below 8 mmol/L; 35 min dynamic scan is acquired approximately 40 min after an intravenous injection of 3.7 MBq/kg of 18F-FDG. For $A\beta$ -PET, a 35 min dynamic scan is acquired approximately 40 min after an intravenous injection of 7–10 mCi [¹⁸F] florbetapir. The PET data are acquired using a TOF-OSEM algorithm (time-of-flight ordered subset expectation maximisation) with the following parameters: eight iterations, 32 subsets matrix=192×192, field of view (FOV)=350×350, half-width height=3.

SMRI: Parameters for T1-weighted 3D brain structural images are as follows: SPGR sequence, FOV=256×256 mm², matrix=256×256, slice thickness=1 mm, gap=0, slice number=192, repetition time (TR)=6.9 ms, echo time (TE)=2.98 ms, inversion time (TI)=450 ms, flip angle=12°, voxel size=1×1×1 mm³.

DTI: DTI data are obtained with a single-shot spin-echo diffusion-weighted echo planar imaging (EPI) sequence with the following parameters: FOV=224×224 mm², data matrix=112×112, slice thickness=2 mm, gap=0, slice number=70, slice order=interleaved, TR=16 500 ms,

TE=95.6 ms, 30 gradient directions and 5 b0 images (b=1000 s/mm²), voxel size=2×2×2 mm³.

Rs-fMRI: A single-shot gradient-echo EPI sequence is used for rs-fMRI with the following parameters: scan duration=8 min, FOV=224×224 mm², data matrix=64×64, slice thickness=4.0 mm, gap=1.0 mm, slice number=28, slice order=interleaved, TR=2000 ms, TE=30 ms, flip angle=90°, voxel size=3.5×3.5×4 mm³.

T-fMRI: All participants will also be invited for a block design fMRI paradigm to measure somatosensory mismatch negativity (MMN). We will use a custom-designed air jet pressure stimulation device, which can apply 5.0 bar pressure somatosensory stimulation to the right index or middle finger. The index and middle finger of the right hand will be fixed in place with two pieces of adhesive tapes to restrict any movement during the experiment. The length of the two air pipes from the air compressor to the apertures will be 6.0 m, and the diameter of the two stimulation apertures will be 1.0 mm. In addition, the distance between the aperture and the centre of the finger pad will be fixed at 5.0 mm for all subjects. Subjects will be instructed to watch a video (without sound) presented on a custom designed screen and to ignore stimulation to the fingers.

The functional MRI paradigm will have a total duration of 9 min. In a block design, eight control blocks with 32 s duration and 8 MMN blocks (also 32 s duration) will be used. A constant stimulus onset asynchrony of 500 ms will be employed for both the MMN and control conditions resulting in 40 stimuli per block. In devising the MMN and control blocks, we will take advantage of an MMN paradigm in which two stimuli will be presented equiprobably and their presentation order will be varied. In MMN blocks, stimulation will be arranged to compose alternating ‘mini-sequences’ of stimulus 1 (index finger, 5.0 bar, 300 ms) and 2 (middle finger, 5.0 bar, 300 ms). Mini-sequences will be 2, 3, or 4 repetitions of a single stimulus (each sequence length is represented equiprobably), followed by a mini-sequence of the other stimuli, and so on. The number of trials in a given mini-sequence will vary such that the occurrence of a switch from stimulus 1 to stimulus 2 (and vice-versa) is irregular. As such, the switch trial stimuli will be designated as deviants and will elicit the MMN. This basic stimulation paradigm, which has proven to elicit robust MMNs, will be used for the MMN condition.^{47 48} For the matched control condition to the aforementioned MMN condition, the same two stimuli will be alternated sequentially to form a regular pattern (eg, stimulus 1, stimulus 2, stimulus 1, stimulus 2, etc.).

Imaging data analysis

PET data analysis. The AV-45 and FDG-PET scans will be preprocessed by Statistical Parametric Mapping V.12 (SPM 12; <https://www.fil.ion.ucl.ac.uk/spm/software/spm12>). The structural images will be individually registered to the averaged PET images. The unified segmentation method⁴⁹ will then be applied to all coregistered

structural images. The PET images will be spatially normalised to Montreal Neurological Institute (MNI) standard space by using the forward parameters estimated during the unified segmentation and smoothed with an 8mm full width at half maximum (FWHM) Gaussian kernel. Finally, the voxel-wise AV-45 and FDG-PET standardised uptake value ratio (SUVR) will be normalised by the whole cerebellum and pons as the reference regions, respectively. A global AV-45 PET SUVR value will be estimated using a composite of the prefrontal, orbitofrontal, parietal, temporal, anterior cingulate, and posterior cingulate and precuneus cortices.

SMRI data analysis. The cortical thickness analysis will be performed using the FreeSurfer image analysis suite, V.5.3 (<http://surfer.nmr.mgh.harvard.edu>). We will construct models of the boundaries between the grey matter (GM) and the white matter as well as the pial surface.⁵⁰ Cortical thickness measures will be obtained by calculating the distance between these surfaces. Automated reconstruction and labelling will be performed using the default 'recon-all' command line. All generated images will be visually inspected for image and segmentation quality and corrected manually if necessary. Subsequently, 'lobesStrict' will be performed to obtain a lobar annotation and consecutively imported for further statistical analyses. For the voxel-based morphometry analysis, structural images will be segmented into GM tissue using the new segment function within SPM12. The diffeomorphic anatomical registration through the exponentiated Lie algebra toolbox will be used to generate a reference template object of the sample that will be warped into a standard MNI space. The generated flow fields of each subject will be calculated and normalisation parameters will be then implemented to normalise the GM maps in the native space to the MNI space. During the normalisation process, the modulated images (local native amount of GM) will be preserved. Images will be spatially smoothed with an 8mm FWHM Gaussian kernel. Finally, for each individual, we will obtain a smoothed GM volumetric map. Additionally, the total intracranial volume for each individual will be estimated by summing the segmented GM, white matter and cerebral spinal fluid (CSF). Then, the smoothed GM images of every individual will be used for subsequent statistical analysis.

DTI data analysis. Raw DTI data will be processed using the Oxford Centre for Functional MRI of the Brain (FMRIB) Software Library (FSL, <http://www.fmrib.ox.ac.uk/fsl>).^{51–53} Initially, the EddyCorrect tool will be used to correct eddy current distortions and motion artefacts by fine registration of the DTI images to a reference image (b0 image). The Brain Extraction Tool⁵³ will be used for creating brain masks of all subjects, and then a diffusion tensor will be modelled at each voxel by using the least-squares algorithm fitting tensor model within the DTI-FIT Tool.⁵⁴ Fractional anisotropy (FA), mean diffusivity (MD), radial diffusivity (RD) and axial diffusivity (AxD) values of each voxel will be calculated based on the eigenvalues of the tensor. Voxel-wise statistical analysis of the FA, MD,

RD and AxD data will be performed using tract-based spatial statistics.⁵⁵ All subjects' FA maps will be non-linearly coregistered to the FMRIB58_FA template with FSL's non-linear image registration algorithm. Then, the mean FA image will be obtained and thinned to create a mean FA skeleton representing the centre of all tracts common to all subjects. Each subject's aligned FA data will then be projected onto the FA skeleton to obtain their FA skeletons and deformation matrixes. With the deformation matrixes, the skeletonized AxD, MD and RD maps will be created for every individual by the `tbss_non_FA` tool. These maps will be used for subsequent statistical analysis.

Rs-fMRI data analysis. Rs-fMRI will be preprocessed using the Data Processing Assistant for Resting-state fMRI (DPASf; <http://www.rfmri.org/DPARSF>).⁵⁶ The first 10 volumes will be discarded for image stabilisation and the participant's to adaptation to the scanning. The remaining functional sequences will be first corrected for timing differences between each slice and motion effects. Next, the structural image will be coregistered to the mean functional image. Then, the transformed structural images will be segmented into GM, white matter and CSF. The motion-corrected functional volumes will be spatially normalised to MNI 152 standard space and resampled to 3 mm×3 mm×3 mm cubic voxels by using the normalisation parameters estimated during unified segmentation. The resulting images will further undergo spatial smoothing (Gaussian kernel with 8mm FWHM), linear detrending and temporal filtering (0.01–0.08Hz). To avoid overestimating regional homogeneity (ReHo) values, spatial smoothing will be conducted for individual ReHo maps rather than during data preprocessing. Finally, nuisance signals (including Friston 24-head motion parameters, the white matter and CSF) will be extracted and regressed out from the data to reduce the residual effects of non-neuronal factors. For the amplitude of low frequency fluctuation (ALFF) analysis,⁵⁷ the time series of each voxel will be transformed into the frequency domain using a fast Fourier transform. The square root of the power spectrum will be calculated and averaged across 0.01–0.08Hz. This averaged square root will be taken as the ALFF value for this voxel. For the ReHo analysis,⁵⁸ Kendall's coefficient of concordance (KCC) will be computed on the ranked time series of a given voxel with its 26 nearest neighbours. The resultant KCC will be taken as the ReHo values. The generated ALFF and ReHo images will be used for statistical analysis.

Network analysis. We will construct a structural cortical network based on GM volumes.⁵⁹ The nodes will be defined as brain regions corresponding to automated anatomic labelling (AAL) areas. The structural connections will be defined as statistical correlations between pairs of average GM volumes in our study. A structural connection will be considered to exist if the correlation coefficient is statistically significant. Before the correlation analysis, the effects of age, sex and total GM volume on the GM volume of regions will be adjusted. We will calculate Pearson correlation coefficients across

individuals between the average GM of every pair of regions, and then an interregional correlation matrix will be obtained for every individual. We will construct a white matter network based on fibre number. The network nodes will be defined as the 90 regions of interest corresponding to the AAL template. The weight of the network edge will be defined as the number of connected fibres with two regions. The fibre number will be calculated through DTI tractography as reported in our previous study.⁶⁰ To avoid spurious connections, a minimum threshold of fibre number (weight of the edge=10) will be used. We will construct a functional network based on the average time sequence. The network nodes will be defined as the 90 regions of interest corresponding to the AAL template. The network edge will be defined as the partial correlation coefficients between the average time sequence of two regions, and we will obtain an incidence matrix. Non-significant correlations will be excluded. The network analyses will be performed with the GRETNA toolbox (<http://www.nitrc.org/projects/gretna/>).⁶¹ The 'rich-club coefficient' is defined as the density of connections between rich-club nodes and rich-club regions in our study will be defined as the top 13 brain regions with the highest degree as reported in our previous study.⁶² A module is defined as a subset of nodes connected to the other nodes in the same module other than those outside the module.⁶³ We plan to use Newman's metric to measure the modularity⁶⁴ and maximise the modularity parameter Q by the algorithm proposed by Clauset *et al.*⁶⁵ The two parameters will be calculated for the structural cortical network, white matter network and functional network constructed for each individual. The values of the two measures will be calculated for every individual and used in the model construction.

T-fMRI data analysis. We will use SPM12 to process and analyse the t-fMRI data. The first 10 volumes will be discarded due to unsteady magnetization. First, the functional images from each run will be realigned. The structural image will then be coregistered to the first scan in the functional image, and the resulting coregistered structural image will be normalised to MNI 152 standard space. Finally, these spatially normalised functional images will be smoothed (Gaussian kernel with an 8 mm FWHM).

Statistical analysis

Baseline comparison between SCD converters and SCD non-converters

All data (demographic information, clinical data, risk factors, neuropsychologic tests, blood and urine biomarkers, multimodal MRI biomarkers, glucose mentalism and amyloid deposition) will be described. The Shapiro-Wilk test and Q-Q plots will be used to confirm the normality. All normally distributed continuous variables will be reported as the mean \pm SD. A comparison of the baseline between SCD converters and SCD non-converters will be performed with a two-sample t-test for continuous variables and a chi-squared test for

categorical variables. Dunnett's multiple comparison tests will be performed for comparison.

Statistical analyses of t-fMRI data will be subjected to a general linear model analysis. The MMN condition will serve as the predictor of interest and will be modelled with a boxcar function convolved with the canonical haemodynamic response function; the Control condition will serve as a baseline. The beta estimates from the individual general linear models enter a second level random effects analysis. Contrast maps will be created by applying paired t-tests comparing the MMN versus control condition for each group separately as well as a two-sample t-test for the between-groups comparison. Activation maps will be corrected ($p < 0.001$) by the false discovery rate approach implemented in SPM.

Longitudinal patterns in SCD converters

The longitudinal analysis in neuropsychological, plasma, urine and MRI variables in SCD-converters will be assessed. We will use the general linear mixed effects model to estimate the individual's change in each variable.

Statistical prediction models

We will use the competing risk regression model to detect the association between possible risk factors and endpoint. The endpoint event of our study is converting to MCI. Death before conversion to MCI is considered a competing risk in our study. The time to event is the date difference between the baseline neuropsychological tests and the date at which the MCI or dementia is first diagnosed.

Independent variables are listed in [table 2](#), including features of baseline clinical characteristics, blood, urine, MRI and PET biomarkers. Considering the overfitting effect when establishing the model, the least absolute shrinkage and selection operator (LASSO) model with penalty parameter tuning will be used for variable screening. It is a data mining method for shrinkage estimation and dimensionality reduction, overcoming processing difficulties caused by high-dimensional data, and estimating the parameters more accurately. Then, we will perform the sub-distribution hazard function model, which could evaluate hazards for the endpoint (MCI or dementia) and competing (death) events, to establish the final multivariate models. Ten-fold cross-validation is used to perform internal verification of the established models. The optimism-corrected C-statistics will be used to evaluate the performance of risk prediction models.

To better clarify the relationship between possible risk factors and endpoint events, some confounding factors will be adjusted. Four models will be established in our study. Model 1 will simply take demographic data, lifestyle, clinical assessment, neuropsychological assessments and SCD report into account. Model 2 will add the ApoE genotype, plasma A β and urine AD7c-NTP on the basis of model 1. Model 3 will take neuroimaging data into account on the basis of model 2 and model 4 will include amyloid-PET and FDG-PET on the basis of model 3.

Table 2 Independent variable list**Types**

Demographic data	Age; education level; family history of dementia; socioeconomic information
Lifestyle	Nicotine and alcohol use; sleep quality (PSQI, RBDSQ, ESS); nutrition style
Clinical assessment	Vascular risk factors (hypertension, hyperlipemia, diabetes, coronary heart disease, Fazekas score); medial temporal lobe atrophy scale; HAMA; HAMD; GDS
Neuropsychological assessment	AVLT-D; AVLT-R; STT-A; STT-B; BNT; AFT
SCD report	Onset time; comparison with others; SCD-Q9; ECoG; consistency of SCD; inform report
Blood tests	ApoE genotype; Plasma β -amyloid
Urine tests	Level of AD7c-NTP
MRI data	Cortical thickness; grey matter volume; FA; MD; RD; AxD; ALFF; ReHo; network characteristic (rich-club coefficient, modality); MMN-related haemodynamic responses
PET data	Global SUVR of FDG-PET and A β -PET

AFT, animal fluency test; ALFF, amplitude of low-frequency fluctuations; AVLT, Auditory Verbal Learning Test; AxD, axial diffusivity; A β -PET, β -amyloid positron emission tomography; BNT, Boston naming test; ECoG, Everyday Cognition; ESS, Epworth Sleepiness Scale; FA, fractional anisotropy; FDG-PET, fluorodeoxyglucose positron emission tomography; GDS, Geriatric Depression Scale; HAMA, the Hamilton Anxiety Scale; HAMD, the Hamilton Depression Scale; MD, mean diffusivity; MMN, mismatch negativity; PET, positron emission tomography; PSQI, Pittsburgh sleep quality index; RBDSQ, Rapid Eye Movement Sleep Behaviour Disorder Screening Questionnaire; RD, radial diffusivity; ReHo, regional homogeneity; SCD, subjective cognitive decline; STT, Shape Trails Test; SUVR, standardised uptake volume ratios.

The HRs with 95% confidence intervals are determined. A two-sided $p < 0.05$ is defined as statistical significance. All statistical tests were performed using R statistical software. The 'glmnet' package will be used for the LASSO model analysis, and the 'cmprsk' package will be performed for the sub-distribution hazard function model.

DISCUSSION

The current study evaluates the characteristics of SCD patients presenting cognitive decline within 5 years. Furthermore, we will construct risk forecast models based on different combinations of potential predictors to achieve early diagnosis in the preclinical stage of AD.

Considering the long preclinical stage of AD and compensated cognitive function in the SCD stage, the application of a refined SCD approach as an enrichment strategy for clinical trials focusing on preclinical AD shows great promise. SILCODE is a longitudinal study, and we

will focus on features of SCD converters with multi-perspective analysis. In particular, we would like to establish an integrated diagnostic system for the early detection and prediction of SCD progressing at the individual level. We aim to provide scientific evidence for a more effective diagnosis of SCD due to AD and explore the underlying mechanism.

In this project, plasma A β , urine AD7c-NTP, and multi-modal MRI will also be included in the predicting models. Plasma and urine biomarkers are non-invasive and feasible in clinical practice. Previous studies have found their diagnostic efficiency in MCI and dementia,^{46 66} but in SCD individuals, there are few pieces of evidence. Therefore, this project also focuses on new biomarkers and tries to identify their usage in the clinical scene. Additionally, through this project, we would like to reveal the longitudinal patterns of involved clinical and MRI biomarkers, especially the temporal sequences, which may provide a basis for us to understand the changes during disease progression.

The stage model theory proposes that human memory can be divided into three stages: sensory memory, short-term memory, and long-term memory.⁶⁷ The MMN has been proposed as an objective measure of the existence of auditory (visual, somatosensory) sensory memory traces.⁶⁸ Previous studies have suggested that MMN may have the potential to measure the age-related changes⁴⁷ or improve the diagnostic value for the early diagnosis of AD.⁶⁹ Although many more SCD studies have concentrated on short-term memory and long-term memory,^{18 70 71} there are very few published studies on the sensory memory stage. SILCODE will investigate the changes in the initial memory stage in the SCD population and the predictive value of somatosensory MMN for disease progression in an exploratory way.

Finally, cultural factors were found to impact SCD by a cross-cultural comparison between the USA and China,²¹ which indicates the need for SCD research in China. SILCODE is a longitudinal study in China, and the implementation of SILCODE would provide characteristics of SCD in China, which contribute to the harmonisation of the SCD concept across cultural borders.

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