Chemical genetic activation of the cholinergic basal forebrain hippocampal circuit rescues memory loss in Alzheimer’s disease

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Abstract

Background: The degeneration of the cholinergic circuit from the basal forebrain to the hippocampus contributes to memory loss in patients suffering from Alzheimer’s disease (AD). However, the internal relationships between the acetylcholine (Ach) cycle and memory decline during the early stages of AD currently remain unknown. Here, we investigate the mechanisms underlying the activation of the cholinergic circuit and its impact on learning and memory using APP/PS1 mice models.

Methods: Novel object recognition and Morris water maze tests were used to measure learning and memory function. Magnetic resonance spectrum (MRS) imaging was applied to longitudinally track changes in neurochemical metabolism in APP/PS1 mice aged 2, 4, 6, and 8 months. The number of neurons and the deposition of Aβ plaques were measured using Nissl, immunohistochemistry, and Thioflavin S staining. We then employed a chemogenetic strategy to selectively activate the cholinergic circuit from the medial septal nucleus (MS) and the vertical limb of the diagonal band nucleus (VDB) on the basal forebrain to the hippocampus. MRS and immunoblotting techniques were used to measure the neurochemical metabolism levels and cholinergic-related proteins, respectively.

Results: We found that the levels of choline (Cho) in the basal forebrain were markedly higher compared to other brain regions and that its decrease along with N-acetyl aspartate (NAA) levels in the hippocampus was accompanied by memory deficits in APP/PS1 mice aged 4, 6, and 8 months. In terms of pathology, we observed that the deposition of Aβ plaques gradually aggravated throughout the cerebral cortex and hippocampus in APP/PS1 mice aged 6 and 8 months, while no Aβ deposition was detected in the basal forebrain. In contrast, the activity of choline acetyltransferase (ChAT) enzyme in the basal forebrain was decreased at 6 months of age and the cholinergic neurons were lost in the basal forebrain at 8 months of age. In addition, the activation of the cholinergic circuit from the MS and VDB to the hippocampus using chemical genetics is able to improve learning and reduce memory impairment in APP/PS1 mice. Similarly, the levels of Cho in the basal forebrain; NAA in the hippocampus, as well as the expression of ChAT and vesicular acetylcholine transporter (vAchT) in the basal forebrain; and muscarinic acetylcholine receptor 2 (CHRM2) in the hippocampus all increased.

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Conclusions: These findings demonstrate that the neurochemical Cho and NAA of the cholinergic circuit can be used as biomarkers to enable the early diagnosis of AD. In addition, memory impairment in APP/PS1 mice can be attenuated using chemical genetics-driven Ach cycle activity of the cholinergic circuit.

Keywords: Alzheimer’s disease, Learning and memory, Chemical genetics, Cholinergic neural circuit, Neurochemical metabolism

Background

Alzheimer’s disease (AD) is a progressive and fatal neurodegenerative disease that typically occurs in people aged 65 years of age or more. AD typically manifests by a progressive loss of cognitive function and daily activity and the onset of neuropsychiatric disorders [1, 2]. The World Alzheimer Report from 2018 estimated that approximately 50 million people worldwide suffer from dementia and that this number will likely to rise to 152 million by 2050, which represents more than a 3-fold increase compared to today. In addition, it is estimated that the global societal cost for the treatment of dementia reached $1 trillion in 2018 and will rise to $2 trillion by 2030.

The degeneration of the cholinergic function is one of the main pathologies in patients suffering from AD, and the deposition of amyloid β (Aβ) plaque deposition and neurofibrillar tangles are thought to be responsible for cognitive impairment associated with the disease [3]. Autopsy reports on Alzheimer’s patients show that cholinergic neurons decrease by 70–80% in the basal forebrain, while choline acetyltransferase (ChAT) activity is reduced by 50–85% in the cerebral cortex and the hippocampus [4, 5]. The emergence of magnetic resonance imaging (MRI) technology made it possible to study brain structure and the integrity of neuro fibers and uncover atrophies in the basal forebrain and the interruption of cholinergic neurological connectivity between the basal forebrain and the hippocampus [6–8]. Over the past decades, it has been proposed that as AD advances, the degeneration of the cholinergic system is characterized by a decrease in acetylcholine (Ach), ChAT, and an enhancement of acetylcholine esterase (AchE) in the basal forebrain, cortex, and hippocampus which is known as the cholinergic hypothesis [9, 10]. Most importantly, the cholinergic system has become a therapeutic target for AD. Presently, the Food and Drug Administration (FDA) has approved 4 kinds of AchE inhibitors (tacrine, donepezil, rivastigmine, galanthamine) for the treatment of AD. However, these drugs are not effective to cure this disease and cause side effects such as nausea, vomiting, diarrhea, or anorexia [11–13]. Moreover, the use of AchE inhibitors is not recommended for the treatment of moderate AD until patients score below 20 points in the Mini-Mental State Examination (MMSE), which has a maximum of 30 points [14]. Accordingly, despite the ability of AchE inhibitors to delay cognitive decline and improve neuropsychiatric symptoms, it is nonetheless necessary to re-recognize the role played by the cholinergic system in AD progression, especially during the early stages of the disease.

The cholinergic system is a major neuromodulator of the brain and is distributed on a number of different nuclei. Among these, the medial septal nucleus (MS) and the vertical limb of the diagonal band nucleus (VDB) in the basal forebrain represent an important source of cholinergic neurons and send the majority (up to 90%) of cholinergic projections to the hippocampus to regulate episodic memory [15, 16]. Previous studies have demonstrated that acetylcholine released from the basal forebrain causes an oscillation in the theta waves by targeting the muscarinic acetylcholine receptor M (CHRM) of hippocampal pyramidal neurons, which promotes learning and memory function [17, 18]. Another study revealed that electrical or optogenetic stimulation in the MS region can activate cholinergic input in the hippocampus and enhance the synaptic plasticity of the hippocampal Schaeffer collateral pathway that promotes learning and memory in wild type mice [19]. These studies suggest that the cholinergic circuit from the MS/VDB of the basal forebrain projection to the hippocampus can modulate the processing of episodic memory. However, cholinergic circuits are impaired by neurofibrillary tangles and chemical phenotypic changes in cholinergic neurons as AD progresses [15]. Magnetic resonance spectroscopy (MRS) is a technique that quantifies the chemical changes associated with impaired neuronal integrity and cholinergic activity because of disease [20]. N-Acetyl aspartate (NAA) is a neurochemical metabolite which is detected by MRS and considered a chemical marker of neural integrity, indicating a regional decline in the hippocampus and posterior cingulate gyrus in patients suffering from AD [21]. Choline (Cho), the precursor of Ach, decreases in the basal forebrain and the hippocampus of patients with AD [22, 23]. Previously, we used animal MRI/MRS measurements and showed that NAA declines in the hippocampus in APP/PS1 mice during moderate-severe pathological stages [24], suggesting that abnormal AD changes in cholinergic metabolism impair cholinergic activity and the integrity of cholinergic neurons in the basal forebrain and hippocampus.
In light of this, it is possible to conclude that the degeneration of the cholinergic circuit from the basal forebrain to the hippocampus contributes to memory loss in AD, even though the early cholinergic activity and metabolic changes associated with memory dysfunction in AD remain poorly understood. Here, we examined memory loss improvement in APP/PS1 mice using longitudinal tracking of the early cholinergic changes in the basal forebrain and hippocampus by $^1$H MRS following memory performance evaluation and chemical genetic activation of the cholinergic circuit from MS/VDB to the hippocampus. We provide insights that deepen our understanding of cholinergic dysfunction during the early stages of AD and that can help in developing more effective therapeutic interventions to improve AD-related cognitive impairment.

Materials and methods

Animals and ethics

Male APPswe/PSEN1dE9 mice and female wild-type mice were purchased from the Model Animal Research Center of Nanjing University (SYXXK2013-009). The animals were fed and cultured in the SPF laboratory of the Experimental Animal Center of Fujian University of Traditional Chinese Medicine. For the current experiment, we obtained a mouse line by crossing APP/PS1 males with wild type females. The animals were raised by the Experimental Animal Center of Fujian University of Traditional Chinese Medicine [license number: SYXK (Min) 2019-0007]. The mice were housed in groups of 3–5 mice/cage under a condition of 12-h light-dark cycle, consistent ambient temperature (21–25°C) and humidity (60–70%). All animal experiments were conducted following the Guidelines for Animal Experimentation of the Fujian University of Traditional Chinese Medicine approved by the Animal Experimental Ethics Committee of the Fujian University of Traditional Chinese Medicine (Fuzhou, Fujian province, China).

Experimental protocol

This study was divided into two experiments. In experiment I, we detected the neurochemical substance alterations in APP/PS1 mice with the progression of AD. Specifically, we evaluated learning memory ability, and the neurochemical substances NAA, Cho, and mI in the basal forebrain and the hippocampus were detected in WT and AD groups at 2, 4, 6, and 8 months of age. These evaluations were accompanied by behavioral tests and MRS imaging. Each mouse group contained a total of 7 individuals. In experiment II, we chose 4-month-old APP/PS1 mice based on the results of experiment I to reveal the treatment effects of chemical genetic technology activating the cholinergic neural circuit from the MS/VDB to the hippocampus. Specifically, 4-month-old APP/PS1 mice were randomly divided into a CNO group and a Saline group, each composed of 7 individuals. Thirty days after the injection of the hM3dq virus, the mice in the CNO group were injected with a clozapine-N-oxide (CNO) solution to activate the cholinergic neural circuit, while the mice in the Saline group were injected with the same volume of saline to serve as controls. We measured the learning memory ability and neurochemical metabolism of the two groups before and after CNO/saline injection.

New object recognition

The new object recognition (NOR) apparatus consisted of a white box (50×50×50cm). Object “A” was the same as object “a,” which were regarded as familiar objects. In addition, there was an unfamiliar object “B.” A camera was installed directly above the apparatus to record the activities of the mice, including the time each mouse explored each of these objects. Before the NOR test, each mouse was placed into the empty box to explore freely for a total of 5 min and adapt to the “new” environment. In the familiar stage, objects “A” and “a” were put in two adjacent corners of the NOR box, after which each mouse was put into the box to freely explore for 10 min. The test stage was conducted 1 or 24 h after the familiar stage. In this stage, the object “a” was replaced by the unfamiliar object “B” after which each mouse was put into the box to freely explore for 5 min. The apparatus was wiped with 75% alcohol to eliminate the odor after each test. The time that each mouse spent exploring the familiar object (TF) and the unfamiliar object (TN) was recorded separately. The recognition index (RI) for the unfamiliar object was defined as $RI = TN/(TF + TN) \times 100%$.

Morris water maze test

The Morris water maze apparatus consisted of a white circular pool with a diameter of 120 cm and a height of 50 cm. The water pool was divided into four equal quadrants, numbered from 1 to 4; a platform diameter of 7.5 cm was placed in the third quadrant, 2 cm under the surface of the water, which was maintained at a temperature of 24±1°C. During the positioning navigation stage, each mouse was placed into the water from four different quadrants each day. This stage lasted a total of 4 days. The time was recorded as the escape latency when the mouse found the platform within 90s and lasted for 3s. Otherwise, we recorded the escape latency as 90s and the mouse would be guided to the platform for 15s to learn. After the positioning navigation stage, we performed a space exploration test at day 5. In this stage, the platform was removed and each mouse was put into the water from the first quadrant and allowed to explore freely for 90s. All activities were recorded, including the time spent in each quadrant, the time of crossing the platform, and the swimming trajectory.
Magnetic resonance spectroscopy
Magnetic resonance spectroscopy (MRS) was performed using a Bruker Biospec 7.0T (70/20USR MRS scanner, Bruker Biospin, Germany). The animals were anesthetized using mixed gas (1.5~2% isoflurane and 30% oxygen). The body temperature of the mice was kept constant using a circulating hot water pump, while their respiratory frequency was constantly monitored. We used location imaging to ensure the head of each mouse was at the center of the image and then employed T2WI images to determine the volumes of interest (VOI). The basal forebrain and the hippocampus were selected as the VOI for scanning with the following parameters: TR=1500ms and TE=144ms. The positions of the neurochemical substances were shown as NAA 2.02 parts per million (ppm), Cho 3.20 ppm, mI 3.60ppm, and Cr 3.05 ppm. After data collection, we used the software package TOPSPIN (v3.1, Bruker Biospin, Germany) to process the spectral data. A quantum estimation (QUEST) method was used to calculate the ratio of the area under the curve of NAA and Cr (NAA/Cr), which represents the chemical metabolic levels of NAA. The chemical metabolic levels of Cho (Cho/Cr) and mI (mI/Cr) were calculated in the same fashion.

Designer receptors exclusively activated by designer drugs (DREADDs)
In experiment II, we injected adeno-associated virus encoding hM3Dq into the MS and VDB nuclei to selectively express the hM3Dq receptors in their respective cholinergic neurons. Detailed procedures
are described below. The mice in the CNO and Saline groups were anesthetized with 1% pentobarbital sodium (0.05 g/kg i.p.) and then positioned in a stereotaxic apparatus (RWD, Shenzhen, China). The rAAV-ChAT-cre-2a-eGFP-WPRE-pA and rAAV-ef1a-DIO-hM3Dq-mCherry-WPRE-pA (1 × 10^{12} particles/ml, Wuhan Institute of Physics and Mathematics, Chinese Academy of Sciences) viruses were mixed in a ratio of 1:3. The mixed virus solution was injected (400nl) into the MS/VDB region (AP +1.10 mm, ML 0.00 mm, DV −4.80 mm; relative to the bregma; AP, ML, and DV denote anteroposterior, mediolateral, and dorsoventral distance from the bregma, respectively) for 10 min using a 10-μl microsyringe. All injections were followed by an additional 5 min to allow for the virus to diffuse before removing the microsyringe. One month after injecting the virus, the clozapine N-oxide solution (0.33 mg/ml dissolved in saline) was administered intraperitoneally 30 min before behavioral tests and MRS imaging were performed in the CNO group (0.2 ml/20g). The same volume of saline was administered to the Saline group.

Thioflavin and Nissl staining
In experiment I, we detected pathological changes in the basal forebrain and hippocampus in 6- and 8-month-old APP/PS1 mice. The WT and AD groups were anesthetized with 1% pentobarbital sodium (0.05 g/kg i.p.). Saline and 4% paraformaldehyde were perfused from the left ventricle to,
respectively, wash and fix the brain tissue. Then coronal slices of the brain tissue were prepared for pathological staining, using 5μm per section. To perform Thioflavin S staining, the tissues were placed in 0.3% Thioflavin S solution (Thioflavin S, Sigma, T1892) and then differentiated in 50% ethanol. Nissl staining was performed with Nissl Staining Solution (Nissl Staining Solution (Cresyl Violet), Solarbio, G3410) according to the manufacturer’s instructions.

Immunohistochemistry staining

Immunohistochemical staining was performed by standard methods. Paraffin-embedded tissues were sliced into 5-μm sections. Citric acid buffer was used at 95°C for 1–2 h for antigen retrieval. Then the sections were incubated in 0.3% H₂O₂ for 10 min. Following this procedure, the sections were washed in PBS for 10 min and then blocked in 3% bovine serum albumin (BSA) for another 30 min. Next, the sections were incubated with primary antibody goat-anti-ChAT (Sigma, AB144P, 1:100) for 48 h at 4 °C. And then the sections were washed in PBS for 10 min and incubated with secondary antibody biotinylated-anti-goat (Vector Laboratories, PK4005, 1:100) for 2 h. Then the brain sections were washed in PBS for 5 min and incubated with ABC-kit solution (Vector Laboratories, PK-4005) for 2 h. After being washed in PBS for 10 min, the brain sections were stained in DAB staining solution for 3 min and then quickly washed in PBS. Finally, the

![Fig. 3 Material metabolism of WT and AD groups at the 2nd and 4th month as measured by ¹H-MRS. A, B Material metabolism in the basal forebrain (A) and hippocampus (B) of WT and AD groups at the 2nd month. C, D Material metabolism in the basal forebrain (C) and hippocampus (D) of WT and AD groups at the 4th month.](image-url)
sections were mounted with neutral resin. The average number of ChAT-positive neurons in the basal forebrain was calculated manually and the average optical density value was calculated by ImageJ.

Western blotting

In experiment II, cholinergic-related proteins of the basal forebrain and hippocampus in the CNO and Saline groups were detected through Western blotting after DREADDs intervention. The brain tissues of the basal forebrain and hippocampus were readily put on ice. RIPA lysate was used to extract the protein, and a BCA assay was then applied for protein quantification. Brain homogenates were resolved by SDS–polyacrylamide gel electrophoresis (SDS-PAGE) and transferred to activated PVDF membranes. The membranes were blocked with 5% skimmed milk powder in pure water. The following antibodies were incubated overnight at 4°C: ChAT (1:5000; Abcam, 181023), AchE (1:1000; Abcam, 183591), ChT1 (1:5000; Abcam, 154186), vAchT (1:1000; Sigma, SAB4200559), CHRM2 (1:5000; Abcam, 109226), and GAPDH (1:8000; Lab, G0100). The membranes were washed in TBST and incubated with an HRP-conjugated secondary antibody (1:5000) for 1h at room temperature. The protein bands were visualized using enhanced chemiluminescence and

Fig. 4 Material metabolism of WT and AD groups at the 6th and 8th month as measured by 1H-MRS. A, B Material metabolism in the basal forebrain (A) and hippocampus (B) of WT and AD groups at the 6th month. C, D Material metabolism in the basal forebrain (C) and hippocampus (D) of WT and AD groups at the 8th month. *P<0.05, **P<0.01
imaged with a Bio-Image Analysis system (Bio-Rad Laboratories, Inc.).

**Verification of MS/VDB to the hippocampus circuit and expression of hM3Dq**

In experiment II, the mice in the CNO and Saline groups were sacrificed after behavioral tests were performed. We observed histological verification of the MS/VDB-hippocampus circuit and evaluated the expression of hM3Dq receptors. The mice in the CNO and Saline groups were anesthetized with 1% pentobarbital sodium (0.05 g/kg i.p.). Saline and 4% paraformaldehyde were perfused from the left ventricle to, respectively, wash and fix the brain tissue. We then prepared coronal slices (30μm per slice) of the brain tissue for confocal laser imaging.

**Statistical analysis**

The data were analyzed using the SPSS22.0 software, and the results expressed as the mean±SEM. The escape latency data acquired in the Morris water maze were analyzed by repeated measurements of variance analysis, and others were analyzed using Student t-tests. The difference was considered as significant when $P < 0.05$.

**Results**

**Longitudinal tracking of early memory impairment processes in APP/PS1 mice**

The NOR and Morris water maze tests were used to evaluate object recognition memory and spatial learning memory at the 2nd, 4th, 6th, and 8th month, respectively. The results of the NOR test showed the recognition index gradually decreased with age in AD mice. Specifically, we found no significant difference between the two groups on the 1-h and 24-h recognition index (RI) at the 2nd month (Fig. 1A, B). However, 4-, 6-, and 8-month-old AD mice showed a decreased recognition index in comparison to age-matched wild type mice (Figs. 1B and 2A, B). In regard to the positioning navigation trial in the Morris water maze, repeated-measures ANOVA showed that the escape latency of the two groups on the 2nd, 4th, 6th, and 8th month displayed a downward trend as training time increased. Comparisons between the groups showed
that 6- and 8-month-old AD mice exhibited significantly longer escape latencies compared to wild type mice of the same age (Figs. 1C, D and 2C, D). In the space exploration test, we observed that both platform crossing time and percentage of time spent in the 3rd quadrant decreased in 4-, 6-, and 8-month-old mice compared with wild type mice of the same age. No changes were observed between the two groups on the 2nd month (Figs. 1E–G and 2E–G). These results suggest that the learning memory of AD mice is already impaired at the 4th month and intensifies as the disease progresses.

Longitudinal tracking of early neurochemical metabolic abnormalities of the cholinergic circuit in APP/PS1 mice
The VOI of MS and VDB nuclei of the basal forebrain and hippocampus were selected to examine $^1$H-MRS imaging in WT and AD mice groups at different ages. In $^1$H-MRS, NAA is regarded a marker of neuronal integrity while Cho is a marker of membrane synthesis and degradation and is also a potential marker for cholinergic neuron. The $^1$H-MRS results showed that the ratio of Cho/Cr in the basal forebrain is significantly higher than that of the hippocampus due to the presence of abundant cholinergic neurons in the basal forebrain (Figs. 3 and 4). A comparison between the WT and AD groups revealed the ratio of Cho/Cr in the basal forebrain decreased in 6- and 8-month-old APP/PS1 mice compared to wild types of the same age (Fig. 4A, C). We observed no difference between the two groups in the Cho/Cr ratio in the hippocampus at different ages (Figs. 3B, D and 4B, D). In contrast, the NAA/Cr ratio in the hippocampus declined at the 6th and 8th months in APP/PS1 mice compared to wild type individuals, while in the basal forebrain this decrease was only evident in 8-month-old APP/PS1 mice (Fig. 4). No changes of mI/Cr in basal forebrain and hippocampus were detected in APP/PS1 mice (Figs. 3 and 4).

Aβ deposition in the brain and neuronal loss in APP/PS1 mice
The pathological changes of the basal forebrain and hippocampus in APP/PS1 mice were observed...
using Thioflavin, Nissl staining, and immunohistochemistry staining. Thioflavin staining showed that Aβ-immunoreactive plaques were distributed throughout the cerebral cortex and the hippocampus in 6-month-old APP/PS1 mice and that Aβ plaque deposition aggravated at 8 months of age (Figs. 5 and 6). However, there was no Aβ plaque deposition in the basal forebrain in 6-, 8- and even 13-month-old APP/PS1 mice (Fig. S1). Nonetheless, Nissl staining showed that neurons in the basal forebrain were sparsely arranged in 6-month-old APP/PS1 mice and that their number decreased in individuals 8 months of age compared to wild types. No significant neuron reduction was found in the hippocampus of 6- and 8-month-old APP/PS1 mice (Figs. 7 and 8). The results of immunohistochemistry staining showed that ChAT-positive neurons are mainly located in the forebrain structures, including caudate putamen, MS nucleus, and VDB nucleus (Fig. 9A, B). And no significant ChAT-positive neuron reduction was found in the MS and VDB nuclei of 6-month-old APP/PS1 mice. However, comparisons between the WT group and AD group showed that 6-month-old AD mice exhibited significantly decreased ChAT activity in MS and VDB nuclei (Fig. 9C, D).

**Chemical genetics stimulation of the cholinergic circuit and early memory deficits in APP/PS1 mice**

The results of experiment I indicated that learning memory and the function of cholinergic neural circuit were impaired in APP/PS1 mice aged 4–6 months. We therefore selected 4-month-old APP/PS1 mice for chemical genetic intervention (Fig. 10A). NOR and Morris water maze tests were used to examine the cognitive function of the CNO and Saline groups before and after chemical genetic intervention. No significant differences were found between the two groups in pre-intervention object recognition memory and spatial learning memory. Overall, chemical genetic intervention was able to ameliorate cognitive decline in the AD mouse model. Specifically, in the NOR test, we observed that the 24-h recognition index in the CNO group increased compared to
the Saline group after chemical genetic intervention. In the Morris water maze test, platform crossing times in the CNO group increased compared to the Saline group (Fig. 10B–G). To further confirm the effects of chemical genetic manipulation, we increased the times of chemical genetic intervention and found a magnification of the effects of chemical genetic intervention in both NOR and Morris water maze tests (Fig. S2). These results suggest the activation of the cholinergic neural circuit from MS/VDB to the hippocampus by chemical genetic technology might improve learning and memory abilities in APP/PS1 mice.

**Chemical genetic stimulation of the cholinergic circuit for treatment of early AD in APP/PS1 mice**

The neurochemical substances in the basal forebrain and hippocampus were measured by 1H-MRS before and after chemical genetic intervention. The results showed no significant differences in the ratio of Cho/Cr and NAA/Cr in the basal forebrain and hippocampus between the CNO and Saline groups before chemical genetic intervention (Fig. 11A, C). However, as shown in Fig. 11B, D, the ratio of Cho/Cr in the basal forebrain and NAA/Cr in the hippocampus increased after chemical genetic intervention. This suggests the activation of the cholinergic neural circuit from the MS/VDB to the hippocampus by chemical genetic technology might promote Cho and NAA synthesis in APP/PS1 mice.

Furthermore, the detection of cholinergic-related proteins in the basal forebrain and hippocampus confirms the function of the cholinergic neural circuit from the MS/VDB to the hippocampus after chemical genetic intervention. Western blotting showed that the CNO group exhibited significant increases in the amount of choline acetyltransferase (ChAT) and vesicular acetylcholine transporter (vAchT) in the basal forebrain after chemical genetic intervention. However, we found no significant differences in the expression of acetylcholinesterase (AchE), ChT1, and muscarinic acetylcholine receptor 2 (CHRM2) in the basal forebrain when comparing the CNO and Saline groups after chemical genetic intervention (Fig. 12A). A comparison of the hippocampus between these two groups further revealed that
the expression of muscarinic receptor 2 (CHRM2) was enhanced in the CNO group following chemical genetic intervention. However, no significant differences in the expression of ChAT, AchE, ChT1, and vAchT were found between the groups after chemical genetic intervention (Fig. 12B).

Validation of the cholinergic projection from MS/VDB to the hippocampus
We found that the recombinant adeno-associated virus with ChAT promoter-eGFP was expressed in the nuclei of MS/VDB and projected to the hippocampus (Fig. 13A), which demonstrates this region of the brain receives cholinergic projections from the basal forebrain, mainly stemming from the nuclei of MS and VDB.

The recombinant adeno-associated virus expressing hM3Dq-mCherry colocalized with ChAT-eGFP in the nuclei of MS and VDB in the basal forebrain (Fig. 13B, C), suggesting the chemical genetic excitatory receptor hM3Dq was successfully expressed in cholinergic neurons of the basal forebrain.

Discussion
The cholinergic hypothesis was first put forward more than 30 years ago and assumes that memory deficits of AD are associated with an irreversible deficiency in cholinergic function [10]. Abundant evidence from previous studies showed that the basal forebrain neuronal projections are important for ensuring cognitive functions, such as selective attention, conscious awareness, sensory processing, and learning memory [25, 26]. In addition, it is well established that the cholinergic projection which arises from the MS and VDB nuclei of the basal forebrain to the hippocampus is associated with learning and memory ability [27–30] and that this cholinergic neural circuit is damaged as a result of AD [31]. However, currently available cholinesterase inhibitors, such as donepezil, galantamine, rivastigmine, or tacrine, which were developed following the cholinergic hypothesis, are only able to slightly reduce the short-term cognitive decline in patients suffering from AD and have high withdrawal rates due to adverse effects [32]. The powerful regulation of specific neurons by chemical genetic technology
has been widely utilized to substantiate the involvement of physiological mechanisms in behavioral outcomes [33, 34]. To the best of our knowledge, our current study is the first to perform chemical genetic technology in the cholinergic neural circuit to stimulate the release of acetylcholine in APP/PS1 mice. This approach differs from the use of cholinesterase inhibitors which slow down the degradation of acetylcholine by inhibiting the activity of acetylcholinesterase. Firstly, we used MRS longitudinal tracking of neurochemical substances to reveal the metabolic changes in the MS and VDB nuclei of the basal forebrain and hippocampus in APP/PS1 mice. Surprisingly, the peak of Cho compounds in the basal forebrain was significantly higher than those in the hippocampus and other brain regions (data not shown), which makes this a potential marker of cholinergic neurons that are present in the basal forebrain. The levels of Cho decreased with the degeneration and loss of basal forebrain cholinergic neurons. Furthermore, we found that the activation of cholinergic basal forebrain neurons in MS and VDB nuclei by chemical genetics is capable of increasing Cho in the basal forebrain and is accompanied by an improvement in learning and memory behavior. Therefore, our study demonstrates that the neurochemicals Cho and NAA of the cholinergic circuit might be used as biomarkers for early AD diagnosis.

Magnetic resonance spectroscopy (MRS) is a useful neuroimaging tool to evaluate changes in neurochemical metabolism during AD. The peak in the amount of Cho shown in the MRS mainly results from the presence of glycerophosphocholine and phosphocholine in the cytoplasm. Previous studies often attributed variations in Cho peaks to neuronal integrity and membrane turnover [35] and found that the amount of Cho in the posterior cingulate and inferior precunei increased in patients suffering from AD compared to healthy individuals [36–39]. In addition,
Another study found that abnormal elevations in Cho/Cr ratio were associated with poorer performances in cognition tests [40] and that people with a higher Cho/Cr ratio in the white matter above the lateral ventricles had a higher risk to develop Alzheimer’s disease within the following 4 years [41]. However, other studies reached contrasting conclusions. For example, a study used MRS to measure the Cho/Cr ratio in the brain and found it decreased in several brain regions in patients with AD, including the posterior cingulate, thalamus, frontotemporal areas, and basal ganglia [23]. A different study found the Cho/Cr ratio decreased in the hippocampus and posterior cingulate in patients suffering from AD [42]. Combined with MRS and \(^{18}\)F-fluoro-2-deoxy-d-glucose positron emission tomography (FDG-PET), decreases in the Cho/Cr ratio have been observed in the supraventricular white matter and the medial cortex of patients with AD. Further analysis found the level of Cho/Cr correlates with cognitive status and cerebral glucose metabolism [43]. In addition to this, the exogenous choline complex can be used as an exogenous source of choline that promotes the synthesis of acetylcholine. The combination of citicoline and cholinesterase inhibitors in the treatment of AD patients has been shown to be more effective in improving cognitive function and delaying disease progression than monotherapy using cholinesterase inhibitors [44, 45]. Surprisingly, we found that Cho levels in
the basal forebrain were significantly higher than in the hippocampus, which suggests that the Cho peak is a potential cholinergic marker reflecting the cholinergic function of basal forebrain neurons. The decline in Cho/Cr ratio in the basal forebrain was observed in 6- and 8-month-old APP/PS1 mice compared with wild type mice of similar age, but we found no changes in the hippocampus. It is well recognized that the decline in the NAA/Cr ratio is a common manifestation of Alzheimer’s disease [23, 46]. Our findings support this conclusion as the NAA/Cr ratio in the basal forebrain and hippocampus decreased in APP/PS1 mice.

According to the “Cholinergic hypothesis of Alzheimer’s disease,” the loss of basal forebrain cholinergic neurons is likely behind characteristic cognition dysfunction of Alzheimer’s disease. By combining 3D reconstruction technology and immunohistochemistry (3D-IHC), a study found five familial AD mutations in 5XFAD mice associated with cholinergic-positive neuron loss in the basal forebrain at 9 months compared to age-matched C57BL/6 control mice. However, this was not observed in the 1st, 3rd, and 6th months of age [47]. We have previously shown that the degeneration of cholinergic neurons in the basal forebrain in an APP/PS1 animal model of AD occurred during the 7th month, including MS and VDB nuclei [48]. Another study showed that morphological changes of cholinergic fibers appeared in the cortex and hippocampus as early as

**Fig. 12** DREADDs intervention might increase the expression of cholinergic-related proteins in APP/PS1 mice. A The expression of cholinergic-related proteins in the basal forebrain of the CNO and Saline groups after intervention. B The expression of cholinergic-related proteins in the hippocampus of the CNO and Saline groups after intervention. C The molecular weight of each cholinergic-related protein. *P<0.05
2–3 months of age in APP/PS1 mice. However, cholinergic neuron or ChAT activity remained unchanged in the forebrain structures with age [49]. Here, we applied Nissl staining to observe the arrangement and number of neurons present in the basal forebrain and hippocampus and found no significant reduction in 6-month-old APP/PS1 mice. In addition, we showed that neuron loss in the basal forebrain occurred at 8 months of age. The ChAT-specific staining results showed that the number of ChAT-positive in the MS and VDB nuclei in 6-month-old APP/PS1 mice was not decreased. However, the activity of ChAT enzyme is reduced. The “Amyloid cascade hypothesis” holds that Aβ-amyloid–containing plaques are the main cause behind the degeneration and death of peripheral neurons around senile plaques, as evidenced in numerous studies that detected Aβ deposition across many brain regions, including the cerebral cortex and the hippocampus [50, 51]. The results presented here are thus in accordance with previous observations. However, it is intriguing that we did not detect any Aβ plaques in the basal forebrain in 6-, 8-, and 13-month-old APP/PS1 mice, which suggests neuron degeneration in the basal forebrain is not associated with the presence of Aβ-amyloid–containing plaques. Several lines of evidence now support the idea that the

Fig. 13 Validation of the MS/VDB-HIP circuit and expression levels of hM3dq receptors. A The cholinergic neurons in the MS/VDB projected to the hippocampus (green). B, C AAV virus-transfected hM3Dq receptor was successfully expressed on cholinergic neurons in the basal forebrain (yellow, as the arrows show).
dysfunction or even the loss of basal forebrain cholinergic neuron is associated with abnormal NGF signaling [52].

The disruption of cholinergic neurons in the basal forebrain leads to learning and memory deficits [53], while their activation through cholinesterase inhibitors, deep brain stimulation, and electrical stimulation technology seemingly improve cognitive function in Alzheimer’s disease [9, 54, 55]. However, these experimental approaches applied failed to achieve a precise regulation of cholinergic neurons in specific brain regions. Chemical genetic technology has been successfully applied in 5XFAD mice to reduce the deposition of Aβ plaque [56] and has been applied to improve cognitive function in AD rat models by stimulating noradrenergic neurons in the locus coeruleus [57]. We employed chemical genetic technology to regulate the cholinergic neural circuit from MS/VDB nuclei to the hippocampus in APP/PS1 mice. The behavioral tests suggest that the activation of the cholinergic neural circuit from MS/VDB nuclei to the hippocampus might significantly improve learning and memory abilities in APP/PS1 mice and is accompanied by elevated Cho/Cr and NAA/Cr ratios in the basal forebrain and hippocampus, respectively. NAA is mainly synthesized in the mitochondria of neurons, and a decrease in acetyl-CoA directly leads to NAA reduction, since the acetyl group of acetyl-CoA represents the main substrate for NAA synthesis [58]. Previous studies showed that cholinergic neurons exert far-reaching effects on the synthesis of NAA by regulating the activity of acetyl-CoA [59], whereby we hypothesize that chemical genetic technology is able to enhance the activity of cholinergic neurons in the basal forebrain in order to accelerate NAA synthesis. Similarly, donepezil treatment improves the cognitive function of patients with AD by increasing NAA levels in the hippocampus, cortex, and other brain regions [60, 61]. To confirm the activation of the cholinergic neural circuit from MS/VDB nuclei to the hippocampus, we evaluated the presence of cholinergic-related proteins in the basal forebrain and hippocampus. We found that the expression of ChAT and vAchT in the basal forebrain, and CHRM2 in the hippocampus, was enhanced after chemical genetic intervention. This suggests that chemical genetics triggers cascade activities including ACh transport, release, and synaptic transmission from the basal forebrain to the hippocampus, which can improve early memory impairment in APP/PS1 mice (Fig. 14). These findings might deepen our understanding of the cholinergic hypothesis in early AD stages and provide a basis for a variety of applications, such as chemogenetics, optogenetics, and deep brain stimulation (DBS) treatment of early AD.

**Fig. 14** Activating the basal forebrain-hippocampal cholinergic neural circuit through chemical genetics might improve learning and memory in APP/PS1 mice by accelerating the choline cycle in the basal forebrain and the synthesis of NAA in the hippocampus.
Limitations
Our research confirmed that chemical genetics technology activates the basal forebrain-hippocampus cholinergic neural circuit and can improve learning and memory in APP/PS1 mice. However, the underlying mechanisms need further study. For example, long-term potentiation represents the electrophysiological basis of learning and memory, whereby it is possible that chemical genetics technology improves learning and memory due to synaptic plasticity. Since electrophysiological recordings were not performed in this study, this hypothesis deserves further exploration in the future. It is worth noting that the pathological changes we found in APP/PS1 model mice do not represent all the pathological changes seen in human AD.

Conclusion
Our study demonstrates that (1) cholinergic circuit Cho and NAA can be used as biomarkers for early AD and (2) chemical genetics-driven Ach cycle activities of the cholinergic circuit attenuate early memory impairment in APP/PS1 mice. Our findings help in clarifying the role of the cholinergic hypothesis in early AD.

Abbreviations
3D-IHC: 3D reconstruction technology and immunohistochemistry; 5XFAD: Five familial AD mutations; ACh: Acetylcholine; AChE: Acetylcholine esterase; AD: Alzheimer’s disease; AP: Anteroposterior; Aβ: Amyloid β; ChAT: Acetyltransferase; Cho: Choline; CHRM1: Muscarinic acetylcholine receptor 1; CHRM2: Muscarinic acetylcholine receptor 2; CNO: Clozapine-N-oxide; Cr: Creatine; DREADDs: Designers receptors exclusively activated by designer drugs; DV: Dorsoventral; FDG-PET: 18F-fluoro-2-deoxy-d-glucose positron emission tomography; HM3dsq: Human muscarinic acetylcholine M3 receptor agonist; mI: Myo-inositol; ML: Mediolateral; MMSE: Mini-Mental State Examination; mMRI: Magnetic resonance imaging; mRS: Magnetic resonance spectrum; MS: Medial septal nucleus; NAA: N-Acetyl aspartate; NOR: New object recognition; MRI: Magnetic resonance imaging; ppm: Part per million; RI: Recognition index; T2WI: T2-weighted imaging; TE: Echo time; TR: Repetition time; TF: Time that each mouse explores the familiar object; TN: Time that each mouse explores the unfamiliar object; vAchT: Echo time; VOI: Volumes of interest; WT: Wild type.

Supplementary Information
The online version contains supplementary material available at https://doi.org/10.1186/s13195-022-00994-w.

Acknowledgements
We would like to thank Prof. Shan Baoci at the Institute of High Energy Physics, Chinese Academy of Sciences, for help in the analysis of magnetic resonance spectroscopy.

Authors’ contributions
WL: Conceptualization, resources, data curation, formal analysis, investigation, methodology, writing—original draft, and writing—review and editing; JL, MY, SW: data curation, formal analysis, investigation, methodology, and writing—review and editing; XK, YD, HL: investigation, methodology and writing—review and editing; LC, JT: conceptualization, resources, and writing—review and editing. All authors read and approved the final version of the manuscript.

Funding
This study was supported by grants from the National Natural Science Foundation of China (ref. 81873353).

Availability of data and materials
The data generated in this study is included in the published article.

Declarations
Ethics approval and consent to participate
Experimental animals were raised at the Experimental Animal Center of the Fujian University of Traditional Chinese Medicine (Fuzhou, Fujian province, China) [license number: SYXX (Min) 2019-0007]. All animal experiments were conducted according to the Guidelines for Animal Experimentation of the Fujian University of Traditional Chinese Medicine and approved by their internal Animal Experimental Ethics Committee.

Consent for publication
All authors have read and agreed with the publication of the manuscript in its current form.

Competing interests
The authors declare that they have no competing interests.

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Received: 3 January 2022   Accepted: 27 March 2022
Published online: 13 April 2022

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