

A Peptide That Binds Specifically to the β-Amyloid of Alzheimer's Disease: Selection and Assessment of Anti-β-Amyloid Neurotoxic Effects

Fang Wang¹, Xian-Ling Zhou¹, Qi-Gang Yang¹, Wen-Hua Xu¹, Fei Wang¹, Yong-Ping Chen², Gui-Hai Chen¹*

1 Department of Neurology, The First Affiliated Hospital of Anhui Medical University, Hefei, People's Republic of China, 2 Department of Biomedical Engineering, Johns Hopkins University, Baltimore, Maryland, United States of America

Abstract

The accumulation of the amyloid- β peptide (A β) into amyloid plaques, an essential event in Alzheimer's disease (AD) pathogenesis, has caused researchers to seek compounds that physiologically bind A β and modulate its aggregation and neurotoxicity. In order to develop new A β -specific peptides for AD, a randomized 12-mer peptide library with A β_{1-10} as the target was used to identify peptides in the present study. After three rounds of selection, specific phages were screened, and their binding affinities to A β_{1-10} were found to be highly specific. Finally, a special peptide was synthesized according to the sequences of the selected phages. In addition, the effects of the special peptide on A β aggregation and A β -mediated neurotoxicity *in vitro* and *in vivo* were assessed. The results show that the special peptide not only inhibited the aggregation of A β into plaques, but it also alleviated A β -induced PC12 cell viability and apoptosis at appropriate concentrations as assessed by the cell counting kit-8 assay and propidium iodide staining. Moreover, the special peptide exhibited a protective effect against A β -induced learning and memory deficits in rats, as determined by the Morris water maze task. In conclusion, we selected a peptide that specifically binds A β_{1-10} and can modulate A β aggregation and A β -induced neuronal damage. This opens up possibilities for the development of a novel therapeutic approach for the treatment of AD.

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* E-mail: chenguihai1964@163.com

Introduction

Alzheimer's disease (AD) is a highly prevalent neurodegenerative disorder and the leading cause of dementia in the elderly [1]. The characteristic symptoms of AD patients, including progressive cognitive impairment, memory loss, and behavioral deficits, are closely related to pathologic changes in the brain [2]. Senile plaques, a key pathological feature of AD, are essentially composed of the amyloid-beta (A β) peptide. A β is 39–43 residues long and is generated by two successive proteolytic cleavages of the amyloid precursor protein [3]. AD cases are thought to be chiefly associated with the apparent failures in regulating A β production and clearance, leading to increased levels of AB and consequent neurotoxicity. Neurotoxic Aß is initially released as a monomer; molecular interactions then cause it to aggregate into oligomers, fibrils, and plaques in AD brains [4]. The most aggregation-prone form, $A\beta_{1-49}$, which is the predominant and initial species deposited in the brain parenchyma, is considered to be the major pathogenic form in AD [5]. Oligomers are the most toxic Aβ species [6,7]. However, protofibrillar and fibrillar aggregates including senile plaques are also toxic [8,9].

Although $A\beta$ aggregation leading to deposition is a critical event in AD [10], the factors that affect $A\beta$ aggregation and accumulation

are not completely characterized. It is widely accepted that a considerable number of environmental factors as well as some intrinsic properties of $A\beta$ can work in concert to cause $A\beta$ deposition and aggregation. These factors can influence the thermodynamic stability of the various accessible conformations of Aβ that potentially cause AD. Recent evidence suggests that key subdomains within A β affect its propensity toward aggregation. The N-terminal domain of Aß seems to play an important role in the transition from soluble aggregates to insoluble plaques and acts as a regulatory site controlling both the solubilization and disaggregation process of the A β molecule [11], especially the 10 N-terminal residues of AB [12]. Intriguingly, site-directed antibodies towards the N-terminal residues 3-6 can relieve amyloid burden in the brain of an AD mouse model and improve their ability to perform cognitive tasks [13]. Meanwhile, a few studies suggest that some other regions of AB also play important roles in aggregation, including residues 17-20, 26-30, 30-35, and 39-41 [12,14,15,16]. In terms of therapeutic development, drugs locking these key regions with high specificity can affect the dynamics of the entire $A\beta$ molecule, preventing Aβ self-aggregation and enabling the resolubilization of previously formed aggregates.

Compounds that block $A\beta$ aggregation may ultimately be clinically useful for treating AD [4,17]. Over the years, much effort

has been directed toward screening and designing compounds that inhibit the aggregation and toxicity of A β . It is reported that various compounds have inhibitory effects on the aggregation of A β , such as A β antibodies [18], protease (β - or γ -secretase) inhibitors [19], anti-inflammatory drugs [20], cinnamon extract [21]. However, the stability, safety, validity, cost, and development time limit the suitability of using these agents for different purposes. Recently, peptide-based drugs are now viable alternatives to biopharmaceuticals [22] and are comparable with antibodies in some cases. As drug candidates, peptides have several advantages over antibodies including lower manufacturing costs, higher activity per mass, lower royalty stack, greater stability, and a lower chance of unintended interactions with the immune system [22].

A number of peptides have been designed to bind and inhibit $A\beta$ based on the sequences and structures related to the selfassembling property of AB. Some of these peptides not only have especially strong anti-AB aggregation effects, but they can also inhibit $A\beta$ neurotoxicity in vitro. More importantly, a few peptides can also reduce cerebral amyloid deposition and attenuate ADtype cognitive deterioration. For example, Austen et al. designed peptide-based aggregation inhibitors containing the binding region (residues 16-20) and retro-inversion of these sequences; these peptides can also inhibit Aβ neurotoxicity in vitro [23,24,25]. Nmethylation peptides of regions corresponding to the amyloid self-recognition elements (e.g., residues 17-20) can prevent Aβ aggregation and inhibit Aβ-induced toxicity in vitro [26,27,28,29,30,31,32]. Moreover, two N-methylation peptides can reverse the Aβ-induced inhibition of long-term potentiation at remarkably low stoichiometry [32]. Nevertheless, there may be some potential limitations associated with some of these peptide therapies, including insolubility and toxicity [32]. The various βsheet breaker peptides and the amyloid sequence-derived pentapeptide LPYFDa effectively inhibit fibrillogenesis and the subsequent deposition of amyloids both in vitro and in vivo; furthermore, they can improve behavior in AD model animals [33,34,35,36,37]. D-amino acid peptides attenuate A β aggregation and cell toxicity, and reduce amyloid plaque load in transgenic mice [38,39,40]. Although the high dose necessary to observe these effects may preclude their use as a preventive or therapeutic drug [39], they may be suitable for use as probes for detecting amyloid plaques in living brains [38]. However, despite these promising results, very few aggregation inhibitors have reached clinical trials.

Phage display technology is a powerful method for identifying peptides that can target any type of biomolecule [41]. It is a technique in which bacteriophages are engineered to insert a foreign DNA fragment with their capsid proteins and hence express a peptide on their external surfaces. The peptides selected by this method tend to be directed toward biologically relevant sites on the surface of the target protein. Consequently, peptides derived from library screenings often modulate the target protein's activity both in vitro and in vivo and can be used as lead compounds in drug design as well as alternatives to antibodies for target validation in drug discovery [42]. Recently, 20- and 12-mer peptides were used to screen peptides specific to variable lengths or forms of A β [12,38,43,44].

As potential therapeutic agents, peptides that directly bind to $A\beta$ are still highly desirable. Although phage display with $A\beta_{1\text{--}10}$ as the target to find such peptides is not reported, we reasoned that this method might yield $A\beta_{1\text{--}10}\text{-specific}$ peptides. Here, we successfully identified such a peptide and investigated its properties.

Materials and Methods

Biopanning of M13 phage display library against $A\beta_{1-10}$

A randomized 12-mer peptide library presented on M13 phages (PhD-C7C; New England Biolabs, USA) was used for screening against Aβ₁₋₁₀ (Xi'an Huachen Biotechnology, China). The well of an ELISA plate (Shanghai Go On Chemical Ltd., China) was coated with a 100-µl aliquot of 100 µg/ml streptavidin (Sigma, USA) and 0.1 mol/l NaHCO₃ (pH 8.6) overnight at 4°C. A 10-µl aliquot of 2.0×10¹³ phage units of purified phage from the initial library with 100 μg/ml biotin-Aβ₁₋₁₀ (Xi'an Huachen Biotechnology, China) was solubilized in 400 µl phosphate-buffered saline (PBS) with 0.05% Tween-20 (PBST, pH 7.4). The mixed solution was subsequently incubated for 60 min at 37°C and overnight at 4° C to allow complete binding between phages and A β_{1-10} . The next day, the solution in the ELISA plate was discarded, and the mixed solution was added to the well for 60 min. The well was washed 5 times with PBST followed by 5 washes with glycine-HCl in order to eliminate the non-specific bound phages. Then, 0.1 mmol/l NHS-LC-Biotin (Sigma, USA) was added to the well and incubated with agitation for 10 min twice. Next, phages bound to the target were eluted with 0.2 mol/l glycine-HCl buffer (pH 2.2) with 0.1% bovine serum albumin (Sigma, USA). After 10 min, the solution was neutralized with 15 µl 1 M Tris–HCl buffer (pH 9.1).

After the round of biopanning, logarithmically growing *Escherichia coli* (ER2738 host strain; New England Biolabs, USA) was infected with a portion of the eluted phages. After amplification, the bacteria were removed by centrifugation, and the phages were purified by serial precipitation with a 20% PEG-8000/2.5 M NaCl solution overnight at 4°C. The phage pellet obtained by centrifugation was finally solubilized in TBS (50 mM Tris-HCl and 150 mM NaCl) supplemented with 0.02% NaN₃.

The titers of eluted phages were determined by serial dilution after the round of biopanning. Amplification was determined by counting the blue plaques obtained after *E. coli* infection and culturing on a selective medium containing isopropyl-beta-d-thiogalactoside (ICN; Biomedical, Belgium) and 5-bromo-4-chloro-3-indolyl-beta-d-galactopyranoside (Sigma, USA). The remaining phages were grown in bacterial culture overnight, pack-aggregated, expressed through co-infection with a helper phage, and precipitated from the bacterial supernatant. The precipitated phages were then used for subsequent rounds of panning until an obvious enrichment phenomenon appeared.

Assay to determine the affinity of the selected phages for $A\beta_{1-10}$

Real-time biomolecular interaction analysis (BIA; Amersham Pharmacia, USA) based on surface plasmon resonance was used to assess the affinity of the interaction between the selected phages and $A\beta_{1-10}$. In the binding specificity test, 100 µg/ml streptavidin and 0.1 mmol/l biotin were immobilized in flow cell 1, and 100 µg/ml streptavidin and 100 µg/ml biotin-A β_{1-10} in flow cell 2, by amine coupling onto the carboxylated dextran layer of a CM5 chip. The screened phage (1.0×10^{11}) phage units) was then injected over flow cells 1 and 2 at 5 µl/min for 6 min. The interaction was monitored as a change in the SPR signal. Greater differences in the response signals (RU) between cells 2 and 1 indicate higher affinity. The procedure of the competitive inhibition test was similar to that of the binding specificity test, except that the selected phage pre-combined with $A\beta_{1-10}$ was injected. BIA evaluation 3.2 software was used to assume one-toone binding. For the Biacore experiments, samples were run in duplicate, with HBS-EP (5 mM HEPES, 150 mM NaCl, 3.4 mM EDTA, and 0.005% surfactant P20 (pH 7.4) as the running buffer.

One surface of the chip in each experimental setup was activated and deactivated, and used as a reference. The response from the reference surface was subtracted from all curves when evaluating the results.

Sequencing of selected phage clones and synthesis of the special peptide

Phage single-stranded DNA was isolated using phenol/chloroform extraction [45]. The phage genome was sequenced using the Sanger method with two sequencing primers: 5-HOGTA TGG GAT TTT GCT AAACAA C-3 and 5-HOCCC TCA TAG TTA GCG TAA CG-3 (New England Biolabs, USA). The DNA sequence was analyzed on a CEQ 2000 XL DNA Analysis System (Beckman-Coulter, Belgium). The sequence was read automatically using the JaMBW 1.1 software.

The optimal monoclone phage displaying sequence was selected and synthesized by Xi'an Huachen Biotechnology Company, using standard solid-phase peptide synthesis. All peptides were purified and characterized by MALDI-TOF mass spectrometry.

Morphological changes in $A\beta_{1-42}$ aggregation under various conditions

 $A\beta_{1\text{-}42}$ was dissolved in sterile filtered H_2O at 1.0 mg/ml and was immediately diluted to 0.5 mg/ml (115 μM) in PBS with or without an equimolar concentration of the special peptide or selected phage $(1.0\times10^{11}$ phage units), $A\beta_{1\text{-}10}$, or $A\beta_{1\text{-}42}$ alone as a control. Aggregation was allowed to progress in an incubator for 7 days at 37 °C. Electron micrographs were taken using a JEOL 100CX transmission electron microscope with uranyl acetate negative staining.

Cell toxicity assay

 $A\beta_{1-42}$ was dissolved in sterile filtered H_2O at $100~\mu M.$ The solution was incubated in an incubator at $37^{\circ}C$ for 7 days, so that $A\beta$ would go into the "aggregated phase," and stored at $4^{\circ}C$ before use. Low-differentiated PC12 cells (Cell Bank, China) were maintained in logarithmic-phase growth on poly-l-lysine-precoated (Sigma, USA) 100-mm dishes (Corning, USA) in DMEM (Gibco, USA) containing 10% fetal bovine serum, 100 units/ml penicillin, and 100 $\mu g/ml$ streptomycin. Cultures were incubated at $37^{\circ}C$ in a 5% CO $_2$ humidified atmosphere.

Different concentrations of aggregated $A\beta_{1-42}$ were tested to assess its effect on neuronal cell viability. After PC12 cells were incubated with $A\beta_{1-42}$ (0 (control), 0.01, 0.1, 1, 10, and 20 μ M) for 24 h, cell viability was measured using cell counting kit-8 (CCK-8; Beyotime Institute of Biotechnology, China).

To evaluate the role of the special peptide in $A\beta_{1-42}$ -mediated neurotoxicity, PC12 cells were incubated in the presence of 20 μ M $A\beta_{1-42}$ together with several final concentrations of the special peptide (0.0004, 0.004, 0.02, 0.1, 0.5, and 2.5 μ g/ μ l) for 24 h. To investigate whether the observed effects were due to the special peptide, PC12 cells were treated with the same final concentrations of the special peptide alone for 24 h and cell viability was subsequently evaluated.

The CCK-8 kit was used to count living cells. Briefly, $10~\mu l$ of the kit reagent was added to the cells treated as described above in 96-well plates and incubated for 3 h. Cell viability was assessed using the ELISA plate reader at 450 nm.

Apoptosis assessment with propidium iodide staining

Propidium iodide (PI) staining was performed to evaluate the status of apoptosis as follows. PC12 cells were incubated for 24 h with any one of the following three treatments: (1) without the

special peptide or aggregated $A\beta_{1-42}$ as a control, (2) with 20 μM $A\beta_{1-42}$ only, and (3) with both 20 μM $A\beta_{1-42}$ and the special peptide at concentrations of 0.0004, 0.004, 0.02, 0.1, 0.5, and 2.5 $\mu g/\mu l$. All samples were centrifuged for 10 min at $800\times g$, and the supernatants were discarded. After washing twice with PBS, the supernatants were fixed in 70% ethanol (100 μl) overnight at – 20°C. The cells were then centrifuged for 10 min at $1,200\times g$, washed twice with PBS, and stained with the DNA-specific fluorochrome propidium iodide (PI; Sigma, USA) at a terminal concentration of 50 $\mu g/\mu l$. The mixed cells were then filtered and incubated in the dark at 4°C for 30 min before flow cytometric analysis (Beckman-Coulter, USA). A total of 1,000 cells were counted to determine the percentage of cells exhibiting the morphological hallmarks of apoptosis such as DNA fragmentation, nuclear condensation, or segmentation.

Animals, surgery, and drug administration

Ethics statement: this study was approved by the Committee on the Ethics of Animal Experiments of Anhui Medical University (Permit Number: 12-2866). All animal procedures were carried out in strict accordance with the recommendations in the Guide for the Care and Use of Laboratory Animals of the National Institutes of Health. All surgery was performed under chloral hydrate, and all efforts were made to minimize suffering.

Forty experimentally naive male Sprague-Dawley rats (SD; Beijing Vital River Experimental Animal Technology, China), weighing 260-280 g at the beginning of the experiment were used. They were housed individually in a room maintained at 23°C with a 12-h light-dark cycle (lights on at 08:00 h) for the duration of the experiment. Rats were allowed free access to food and water except during experimental testing. Rats anesthetized with chloral hydrate (350 mg/kg, i.p.) were positioned in a stereotaxic instrument (RWD Life Science, China), and a cannula (RWD Life Science, China) was implanted into the left cerebral ventricle (A: 0.3 mm, L: 1.2 mm, V: 3.6). Rats were allowed 5 days of recovery after the surgery. The correct location of the cannula was checked by dissecting the brain following the completion of the experiments. Only animals with correctly placed cannulas were used in the evaluation of the experiments. Six rats in each group were used for the final analysis. The special peptide was dissolved in 0.9% saline solution before use. $A\beta_{1-42}$ (Sigma, USA) was also dissolved in 0.9% saline solution. The solution was then incubated at 37°C for 7 days to induce formation of aggregated Aβ and stored at 4°C before use. For intracerebroventricular administration in vivo, aggregated $A\beta_{1-42}$, the special peptide binding to the $A\beta_{1-10}$, or saline solution was infused with the aid of a mini-pump (RWD Life Science, China). The rats were randomly divided into three groups. (1) AD model group: the AD model rats were established as described previously [46]. After surgery, the rats were injected with 3 μ l aggregated A β_{1-42} (1 μ g/ μ l, 0.8 μ l/min) on days 5, 8, and 11, and with 3 µl saline solution (0.6 µl/min) on the other days, days 5–14 (2) The special peptide treatment group with three subgroups: AD model rats were infused daily with 3 µl special peptide at different final concentrations (0.110, 0.522, and 2.610 µg/µl at 0.6 µl/min) for 14 consecutive days from day 5 onwards. (3) Control group: the rats were infused daily with 3 µl saline solution (0.6 µl/min) alone throughout the same period. Upon completion of behavioral testing, the rats were sacrificed by decapitation.

Morris water maze task

The apparatus and procedure were slightly modified as described previously [47]. Briefly, the water maze apparatus consisted of a circular pool 150 cm in diameter, 60 cm deep, and

filled daily to a height of 30 cm with fresh tap water (21~22°C). A black escape platform (10 cm diameter, 24 cm height) was placed in one of the 4 quadrants of the pool (in target quadrant 2) and submerged approximately 1.5 cm below the water surface. Each rat's swimming performance was monitored by a video camera linked to a computer-based image analyzer (Chinese Academy of Medical Sciences, China). Place learning was tested for 10 consecutive days. Each rat was trained to find the platform with 4 successive trials a day. The sequence of water-entering points differed each day, but the location of the platform was constant. The time taken to find the platform (escape latency), swimming speed, and the distance traveled were measured and averaged over 4 trials. If a rat failed to find the platform within 60 s, it was placed on the platform manually; regardless of whether the rat found the platform, it was kept there for 30 s. Each animal was subsequently returned to its cage until the next trial. After all of the rats completed 1 trial, the next trial began and followed the same order of rats. On day 10 of training, an additional trial was given as a probe trial by removing the platform. The animals were placed in the quadrant where the platform was previously located in the fourth starting position and were put through a single test-free swimming period for 60 s. The percentage of distance and time spent in the quadrant where the platform was previously located (target quadrant) were used as measures of spatial memory.

Statistical analysis

The data followed a normal distribution, and are presented as the mean \pm standard error of the mean (SEM). For the place learning test in the Morris water maze task (MWM task), analysis was performed using a two-way repeated-measures analysis of variance (ANOVA) with day and treatment as independent variables. For the other tests, statistical analysis was performed using ANOVA with groups as the independent variable. Posthoc analysis using Tukey test was used to compare results for different days and groups. Statistical significance was set at $P{<}0.05$.

Results

Selection of specific phages binding to $A\beta_{1-10}$

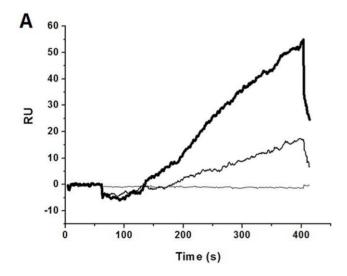
After three rounds of screening, the eluted phage titers progressively increased after each round of panning (Ps < 0.01), suggesting that non-specific phages were eluted and specific phages were amplified.

Affinity of $A\beta_{1-10}$ for the selected phages

In the binding specificity test, as reaction time increased, the positive difference of RU between flow cells 2 and 1 gradually increased (Fig. 1A), indicating that the combined power of the selected phages and streptavidin–biotin–Aβ₁₋₁₀ was significantly stronger than that of the selected phages and streptavidin-biotin. This result suggests that high binding specificity and affinity exist between the selected phages and $A\beta_{1-10}$. In the competitive inhibition test, as reaction time increased, the positive difference in RU between flow cells 2 and 1 also gradually increased (Fig. 1A). This result indicates that the bonding force between the selected phages precombined with $A\beta_{1-10}$ and streptavidin–biotin– $A\beta_{1-10}$ in cell 2 was also higher than that between the selected phages and streptavidinbiotin in flow cell 1. However, the positive difference between cells 2 and 1 in the competitive inhibition test was lower than that in the binding specificity test for all corresponding time points; this is because the binding site of the selected phages is partly locked by the pre-combined $A\beta_{1-10}$. Together, the results suggest that the affinity between the selected phages and $A\beta_{1\text{--}10}$ is highly specific.

Sequencing of the selected phage clones and synthesis of the special peptide

The DNA of the 20 clones selected on the basis of their affinity for $A\beta_{1-10}$ was sequenced (Fig. 1B). The library was enriched in three sequences: PYRWQLWWHNWS, TLAHPYH, and SSSPSKH. Since the 12-mer peptide library was used for screening against $A\beta_{1-10}$, TLAHPYH and SSSPSKH were regarded as contaminants of the other sequences selected in our experiment. The target clone possessed a peptide, PYRWQLWWHNWS, which was then used to synthesize the special peptide.



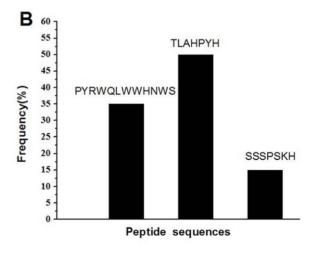


Figure 1. The specific affinity and amino acid sequences of the selected phages. (A) Specific affinity between $A\beta_{1-10}$ and the phages screened for three rounds. Real dark line: binding specificity test of the screened phages; Dash line: competitive inhibition test of the small peptide; Horizon line: the baseline of reaction unit; (B) Amino acid sequence of peptides displayed by the 20 selected phage clones. Frequency of peptides was displayed by the phage clones. Three sequences were highly represented. doi:10.1371/journal.pone.0027649.g001

Morphological changes in $A\beta_{1-42}$ aggregation under various conditions

Some obvious plaques were observed for $A\beta_{1-42}$ or $A\beta_{1-10}$ alone. However, after the addition of the selected phages or the special synthetic peptide, the plaques were instead of bundles of short fibrils. This indicates that the selected phages and special synthetic peptide can inhibit the aggregation of $A\beta_{1-42}$ into plaques (Fig. 2A).

Effects of the special peptide on $A\beta_{1-42}$ -mediated neurotoxicity *in vitro*

As shown in Fig. 2B, the viability of cells treated with $A\beta_{1-42}$ only at concentrations >0.01 μ M was significantly decreased in a concentration-dependent manner (Ps<0.05). Based on these results, a final concentration of 20 μ M $A\beta_{1-42}$ was selected as the optimal concentration for subsequent experiments because cell viability was about 60% at that concentration. In addition, cell death with 20 μ M $A\beta_{1-42}$ appeared to be mainly due to apoptosis (Fig. 2D).

To investigate the effect of the special peptide on $A\beta_{1-42}$ -medidated neurotoxicity, PC12 cells were incubated in the presence of 20 μ M $A\beta_{1-42}$ together with several concentrations of the special peptide. Compared with the control, cell viability was significantly decreased with 20 μ M $A\beta_{1-42}$ treatment (P<0.01). In the treatment groups, the pattern of cell viability exhibited an inverted "U" shape. The intermediate concentrations (i.e., 0.004, 0.02, 0.1, and 0.5 μ g/ μ l) of the special peptide significantly increased cell viability (P_3 <0.05), while smaller (0.0004 μ g/ μ l) and larger doses (2.5 μ g/ μ l) were ineffective (P_3 >0.05) (Fig. 2C). This neuroprotective effect of the special peptide on $A\beta_{1-42}$ -induced cell death resulted completely from the net action of the special peptide on $A\beta_{1-42}$ aggregation, because the viabilities of PC12 cells incubated with different concentrations of the special peptide alone were similar to those of the control group (data not shown).

PI staining revealed that the percentage of apoptotic cells among groups exhibited a "U" shape; i.e., the percentage of apoptotic cells was significantly higher in the presence of $20~\mu M$

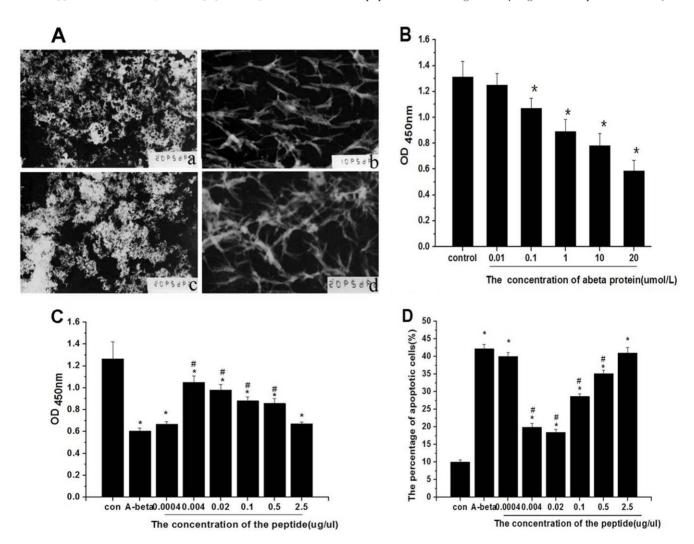


Figure 2. The special peptide alleviates Aβ aggregation and toxic effects *in vitro.* (A) Electron microscopic analysis of morphological change showing the effect of the special peptide or selected phage on A β_{1-42} aggregation. A β_{1-42} (a) or A β_{1-10} (c) alone was incubated for 7 days; A β_{1-42} with special peptide (b) or the selected phage (d) was incubated for 7 days; (B) Different concentrations of aggregated A β_{1-42} were tested to assess its effect on neuronal cell viability by CCK-8 assay. *P<0.05 when compared with the control group; (C) Cell viability was assessed in the presence of 20 μM A β_{1-42} and different concentrations of the special peptide by CCK-8 assay. *P<0.05 when compared with the control group. *P<0.05 when compared with the group treated with 20 μM A β_{1-42} only (note: 0.04 μg/μl peptide equal to 20 μM); (D) Detection of apoptosis in PC12 cells by Pl staining. *P<0.01 when compared with the control group. *P<0.01 when compared with 120 μM A β_{1-42} only. doi:10.1371/journal.pone.0027649.q002

 $A\beta_{1-42}$ than that of the control (P<0.05). Furthermore, the increase was markedly attenuated by combined treatment with several concentrations of the special peptide (i.e., 0.004, 0.02, 0.1, and 0.5 μ g/ μ l) (Ps<0.05), but it was not significantly different from these doses (Ps>0.05) (Fig. 2D).

Performance in the MWM test

Fig. 33 shows the performance of rats in finding the hidden platform in the MWM task. The latency progressively decreased with days for all rats combined ($F_{(9,225)}=91.49$, P<0.01; Fig. 3A), suggesting that the rats were able to learn the task. The repeated-measures ANOVA indicates that the effect of treatment on the latency was significant ($F_{(4,25)}=3.121$, P=0.0324). Post-hoc analysis shows that the latency in the AD model rats was longer than that in the control rats (P=0.041). However, among the three special peptide-treated groups, only the latency of the 0.522 µg/µl group was significantly shorter than that of the AD model group (P=0.047), and was similar to that of the control group.

The cumulative distance also declined daily ($F_{(9,225)} = 78.09$, P<0.01; Fig. 3B). The repeated-measures ANOVA indicates that treatment had significant effect on the cumulative distance ($F_{(4,25)} = 4.941$, P=0.005). The AD model rats swam significantly longer distances than the control rats (P=0.021); only the $0.522~\mu g/\mu l$

special peptide-treated group had a shorter latency than the AD model group (P=0.029). The results of the latency and distance suggest that repeated daily infusion of the special peptide at 0.522 μ g/ μ l significantly ameliorates the impairment of performance caused by A β ₁₋₄₂. Swimming speed was similar among groups (F_(4,25) = 0.897, P=0.467; Fig. 3C), indicating that the cognitive performance observed under the various treatments is not due to potential alterations of locomotor function.

In the probe trial test, the percentage of time in the target quadrant showed significant effect of treatment ($F_{(4,25)} = 7.984$, P < 0.01; Fig. 3D). This indicates that the AD model rats searched the target quadrant for a significantly less time than the control rats (P < 0.001); only the 0.522 µg/µl special peptide-treated rats spent significantly more time than the AD model rats (P = 0.025). The percentages of cumulative distance in each group of rats corresponded with the percentages of escape latency (data not shown).

Discussion

Using phage display technology, we identified a peptide that binds specifically to $A\beta_{1-10}$. Meanwhile, the special peptide inhibited the aggregation of $A\beta_{1-42}$ into plaques, and reduced

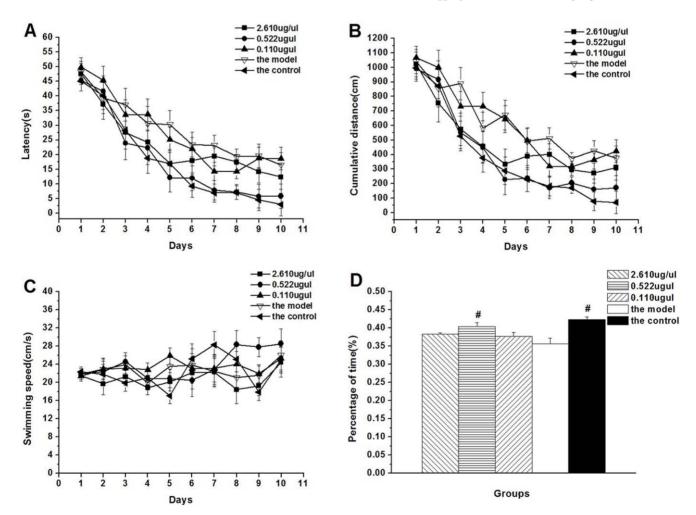


Figure 3. Performance of the rats in the Morris water maze. (A) latency, (B) distance and (C) swimming speed in the place learning test; (D) the percentage of time spent in the target quadrant in the probe trial. $^{\#}P$ <0.05 when compared with the model group. (note: the 0.522 μ g/ μ l special peptide equimolar to A β_{1-42}). doi:10.1371/journal.pone.0027649.q003

neuronal damage induced by $A\beta_{1-42}$ in PC12 cells at appropriate concentrations. Moreover, the special peptide exhibited a protective effect against $A\beta_{1-42}$ -induced learning and memory deficits in the MWM task.

Based on previous studies, we reasoned that a peptide could be selected by phage display technology using $A\beta_{1-10}$ as the target. After three rounds of screening, the phages were selected. The affinity between the selected phages and $A\beta_{1-10}$ was assessed by BIA; the results show that the affinity between them was highly specific. Meanwhile, we found that this method is suitable for identifying positive correlations between selected phages and peptides; it is superior to the orthodox ELISA because it maintains the natural properties of biomolecules without the need for fluorescent and isotope labeling, therefore making the results more reliable. Three amino acid sequences were identified; the sequence of one was interesting, while the others were considered to be contaminants. We chose to use this target sequence to synthesize the special peptide.

The presence of the selected phages or special peptide elicited dramatic morphological features in that some observed plaques were instead of bundles of fibrils. This suggests that specific phages and the special peptide could inhibit A β aggregation into plaques. Although this method led to fibril formation, which is considered to be an aggregation pathway originating from a high entropic barrier and a thermodynamically unfavorable event, a recent study indicates that the equilibrium between toxic and non-toxic A β intermediates exhibits a dynamic nature [48]. Furthermore, the non-aggregated and aggregated states of $A\beta$ are in equilibrium, and soluble forms are more accessible to clearance and degradation than insoluble forms [49]. In this study, the special peptide was able to disaggregate plaques due to steady-state equilibrium between $A\beta$ in plaques and in the monomeric form [50]. This ultimately led to increased amounts of monomeric A β , which is more easily cleared from the brain [51] and can play a neuroprotective role in the brain [52].

Extracellular A β peptides are highly cytotoxic to neuronal cells, and the underlying mechanisms may include free radical damage, oxidative stress, and mitochondrial dysfunction of neurons, which ultimately induce apoptosis [2]. Through interactions with unidentified targets on the cell surface, AB initiates a cascade of intracellular events that culminate in neuronal death. Recent studies show that the dying cells exhibit apoptotic characteristics in AD brains and cultures of primary neurons and neuronal cell lines exposed to A β [53,54]. Furthermore, apoptosis in neurons may be responsible for neuronal death in AD [55]. Applying Aβ to cultured cells even at micromolar concentrations can cause apoptosis [56]. PC12 cells were used in this study as a neuronal model because they biochemically and morphologically resemble neurons after differentiation and are particularly sensitive to Aβ peptides [57]. Moreover, they are relatively easy to culture and survive longer than primary cultured neurons [58]. We found that Aβ-induced cell death was attenuated by the addition of the special peptide at 0.004, 0.02, 0.1, and 0.5 µg/µl. Meanwhile, the percentage of apoptotic cells was markedly reduced by combined treatment with the special peptide (i.e., 0.004, 0.02, 0.1, and 0.5 µg/ml). The positive effect of the special peptide was significant at a molar ratio of peptide: AB ranging from 0.1 to 12.5. Interestingly, the relationship between changes in concentration and efficacy of the special peptide in these experiments met the pattern of inverted U-shaped dose-effect curve (IUSDEC). The IUSDEC is widely described and poorly understood phenomenon in the pharmacological field. The basic concept is that, the effects of increasing dosages of a given compound appear to increase up to a maximum, and then the effects decrease [59]. A recent

comprehensive review has exhibited many examples of such curves in pharmacology, such as cardiac glycosides, anti-tumor drugs and drugs of central nervous system, chemoprevention of stroke, traumatic brain injuries [60]. Similarly, it is difficult to elucidate the mechanisms underlying the IUSDEC in our study. We speculated that the lower concentration of the special peptide might be too low to interfere sufficiently the interaction between the special peptide and AB, so that the protective effect could not be produced. On the other hand, the reduced effect at the higher concentration of the special peptide might be due to that it stimulates the body to generate endogenous AB or enhances the toxic process of AB, or produces self-cytotoxicity. For the latter, however, our unpublished data had suggested that the special peptide had no cytotoxicity for PC12 cells, even up to a concentration of 2.5 µg/µl.

An animal AD model induced by intracerebroventricular AB infusion is particularly attractive for evaluating drugs for AD. Ample experimental evidence indicates that single or chronic intracerebroventricular administration of several kinds of AB peptides (i.e., $A\beta_{25-35}$, $A\beta_{1-40}$, and $A\beta_{1-42}$) can induce cognitive impairment [61,62,63,64]. Although the effects of Aβ on learning and memory have been extensively studied, the mechanism by which AB causes cognitive deficits is not clearly understood. A recent study suggests that the spatial learning and memory deficits induced by AB peptides in rodents may not be entirely related to Aβ-induced neuronal damage such as the activation of glial cells, and neuroinflammatory and oxidative responses [65]. And it is also indicated that $A\beta_{1-42}$ -induced mitochondrial mislocalization contributes to late-onset behavioral deficits in a transgenic Drosophila model [66]. According to the validated in vitro concentrations of the special peptide, we chose three in vivo doses (i.e., 0.110, 0.552, and 2.610 μ g/ μ l); the corresponding molar ratios of peptide: A β were 1, 5, and 25, respectively. The results showed that $A\beta_{1-49}$ injection into the lateral ventricle induced spatial learning and memory impairment, and infusing the special peptide alone at 0.552 µg/µl significantly ameliorated impairment in the MWM. The beneficial effect of the special peptide on learning and memory may be related to its activity as a direct binder to $A\beta_{1-10}$. Further studies are needed to fully understand the mechanisms of the protective activity of the peptide against the $A\beta_{1-42}$ -induced learning and memory impairment. Note should be made that a relatively small difference occurred among groups in the MWM, which might be attributable to the AD model based on intracerebroventricular infusion of AB or/and the inadequate sample size in this study. Further, the effect of the special peptide on cognitive function should be confirmed using AD transgenic mice, such as the 5xFADAD model mice [21].

In summary, by using phage display technology, we successfully identified a 12-amino acid peptide that can specifically bind to $A\beta_{1-10}$. Such a peptide may not only serve to inhibit the aggregation of AB into plaques, but also inhibit the toxic effects of Aβ. Therefore, the special peptide may be a potential drug for the treatment of AD.

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Author Contributions

Conceived and designed the experiments: G-HC Y-PC. Performed the experiments: Fang Wang. Analyzed the data: Q-GY. Contributed reagents/materials/analysis tools: W-HX X-LZ. Wrote the paper: Fang Wang. Contributed to obtaining permission for use of SD rats and interpreting the data in the manuscript: Fei Wang.

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