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Featured Article

Synergism of antihypertensives and cholinesterase inhibitors in Alzheimer's disease

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AbstractIntroduction: We investigated the effect of antihypertensive (aHTN) medications and cholines-
terase inhibitors (ChEIs) on the cognitive decline in patients with Alzheimer's disease (AD) and
analyzed synergism by chemogenomics systems pharmacology mapping.
Methods: We compared the effect of aHTN drugs on Mini-Mental State Examination scores in 617

AD patients with hypertension, and studied the synergistic effects. **Results:** The combination of diuretics, calcium channel blockers, and renin-angiotensin-aldosterone

system blockers showed slower cognitive decline compared with other aHTN groups ($\Delta\beta = +1.46$, P < .0001). aHTN medications slow down cognitive decline in ChEI users ($\Delta\beta = +0.56$, P = .006), but not in non-ChEI users ($\Delta\beta = -0.31$, P = .53). **Discussion:** aHTN and ChEI drugs showed synergistic effects. A combination of diuretics, renin-

angiotensin-aldosterone system blockers, and calcium channel blockers had the slowest cognitive decline. The chemogenomics systems pharmacology–identified molecular targets provide system pharmacology interpretation of the synergism of the drugs in clinics. The results suggest that improving vascular health is essential for AD treatment and provide a novel direction for AD drug development. © 2018 The Authors. Published by Elsevier Inc. on behalf of the Alzheimer's Association. This is an open access article under the CC BY-NC-ND license (http://creativecommons.org/licenses/by-nc-nd/4.0/).

Keywords: Alzheimer's disease; Cognitive decline; Cholinesterase inhibitors; Antihypertensive medications; Combination therapy; Systems pharmacology; Clinical data mining

1. Introduction

Alzheimer's disease (AD) has loomed as a major health challenge worldwide. To date, there is no cure for AD, and

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the currently available treatments, such as cholinesterase inhibitors (ChEIs), achieve very limited therapeutic advantage [1]. Because of the complexity in the pathology and etiology of AD, it has been proposed that combination therapies may be more advantageous compared with monotherapies, and there is a great need for studies on combination therapies in AD [2].

Hypertension is one of the most prevalent coexisting diseases in patients with AD, comprising 42% of the AD population [3]. In addition to the fact that both diseases are age-related, it has also been proposed that vascular abnormalities can etiologically contribute to the onset and

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progression of AD [4]. Therefore, targeting the vascular system is a potential strategy for treating AD.

Numerous epidemiologic studies have investigated the relationship among hypertension, antihypertensive (aHTN) medications, and AD, but the results were mixed [5]. Several longitudinal studies have consistently reported that mid-life hypertension is an important risk factor for developing AD and dementia in late age [6]. In older subjects, however, the role of blood pressure in relation to AD seems more obscure and intricate, whereas high incidences of AD or dementia have been reported to have an association with low blood pressure in elderly patients [7,8]; other studies also reported no association [9,10], association with high blood pressure [11], or association with both high and low blood pressure [12]. On the other hand, aHTN drugs have been associated with reduced risk of dementia in observational studies [13,14] and randomized controlled trials (RCTs) [15,16]. Particularly, the use of diuretics, reninangiotensin-aldosterone system (RAAS) inhibitors, and β-adrenergic blockers was related to a slower rate of cognitive decline and lower risk of dementia in elderly patients [17]. Importantly, however, the study population of all the aforementioned studies was elderly individuals rather than patients already diagnosed with AD. Therefore, these results do not necessarily translate to a therapeutic effect in patients with AD. In one study, the use of diuretics was associated with a slower decline in cognitive function among patients diagnosed with dementia [18]. However, this study did not differentiate between AD and other types of dementia including vascular dementia, which is pathologically distinct from AD and may interact with hypertension via different pathways. Therefore, direct evidence of the therapeutic effect of aHTN medications on patients with AD is still lacking. Moreover, none of the studies done in patients diagnosed with AD have examined the effect of combinations of different classes of aHTN drugs, or the combination of aHTN drugs and currently available treatments for AD.

In this observational study, we applied clinical datamining analyses to investigate the effect of aHTN medications on the cognitive decline in patients with clinically diagnosed AD. We then assessed the effect of the combined use of multiple hypertensive drug classes and the combination of aHTN drugs with ChEIs. Furthermore, we applied systems pharmacology drug-target network analyses to better understand, from the molecular level, the effect of anti-AD and aHTN medications on AD and their synergism in clinical contexts. Systems pharmacology applies the principles of systems biology to study pharmacology, and it seeks to understand how medicines work on various molecular targets from complex systems of the body. Then, the drug-target network studies can serve as predictors on new indication(s) for approved drugs and to guide combinational therapy in clinics. Xie and Wang have developed chemogenomics systems pharmacology target (CSP-target) mapping technique [19] by computational analyses on interaction networks of multiple drugs and multiple targets, from a systems pharma-

perspective, using high-throughput cology docking (HTDocking) [20] and TargetHunter algorithm [21] based on the AD [22] and cardiovascular disease (CVD) [23] domain-specific chemogenomics knowledgebases. Such integrated computational methodologies enable the classification of drugs according to their chemical structures and to which proteins they bind, and the CSP-target analyses make predictions about the therapeutic effects of drugs for complex diseases and possible off-target effects. The outcomes from such integrated CSP-target analyses of AD and HTN medications used in patients led to new understanding of drug action synergy by correlating molecular pharmacology with clinical observation for complex diseases. By investigating the molecular targets involved in HTN and AD, the CSP-target map analysis will provide a mechanistic insight into the effect of aHTN medications on AD in clinical contexts. We also correlated such analyses with the reported AD pathology and the mechanism of drug actions in the literature and clinical trials for AD treatment. The outcomes of such studies could be used to guide rationale clinical therapy and even to new drug design and discovery for AD.

2. Methods

2.1. Participants

The subjects of this study were patients examined at the University of Pittsburgh Alzheimer's Disease Research Center from April 1983 to March 2015. The protocols for patient diagnosis and information collection were published previously [24]. Briefly, the patients received a series of clinical examinations evaluating their physical, cognitive, and neurologic status, and a diagnosis was made by a neurologist and a psychiatrist and then reviewed by a committee. Follow-up surveys and cognitive evaluations using the Mini-Mental State Examination (MMSE) were conducted at annual clinic visits and semiannual phone interviews regarding their current status and history of disease and medication.

Probable AD cases with concomitant hypertension (n = 617) were selected from a total of 4364 participants in clinics. These patients will be referred to as set 1. Among these AD cases, 399 had records of aHTN drug use. The aHTN drugs were categorized into four drug classes: diuretics; calcium channel blockers (CCBs); RAAS inhibitors; and others (β -adrenergic blockers, α -adrenergic blockers, arterial vasodilators, and miscellaneous aHTN agents). On the basis of their use of aHTN drugs or their combinations, the set 1 patients were divided into nine groups (Table 1). Later, we contrasted the particular treatment group with slowest cognitive decline (combination the of diuretics + CCB + RAAS) against all other groups. Their baseline characteristics are shown in Table 2. Apolipoprotein E (APOE) genotyping was performed on isolated DNA from blood as described previously or by using Taq-Man genotyping assays [25].

Table 1 Baseline characteristics for set 1 patients (number of subjects under different aHTN drug classes in set 1)

Group	No. of years since the first record of aHTN drug use						
	0	1	2	3	4	5	
No aHTN drug	218	182	99	53	36	34	
Diuretics only	29	27	18	11	6	5	
CCB only	25	22	13	12	7	1	
RAAS only	79	72	47	29	17	8	
Diuretics + RAAS	44	38	28	16	10	8	
CCB + RAAS	31	29	23	17	11	6	
Diuretics + CCB	7	5	4	3	4	2	
Diuretics + CCB + RAAS	24	20	17	13	12	9	
Other groups	160	142	108	63	42	30	
Total	617	537	357	217	145	103	

Abbreviations: aHTN, antihypertensive; CCB, calcium channel blocker; RAAS, renin-angiotensin-aldosterone system.

To investigate the potential synergistic effect of ChEI and aHTN drugs on the cognitive decline during the first 2 years after being diagnosed with probable AD, we selected the patients who did not switch treatment (i.e., from ChEI user to nonuser and vice versa) from set 1. These patients will be later referred to as set 2 (N = 419). On the basis of their usage of ChEI and aHTN drugs, the patients in set 2 were divided into four groups, and their baseline characteristics are shown in Table 3.

2.2. Statistical analysis

The cognitive function, as measured by MMSE, was the primary outcome of interest in this study. Because of skewness, the MMSE scores were power transformed before conducting the mixed-effect linear model analysis. Student *t* test, analysis of variance, or χ^2 tests compared the baseline characteristics of the two sets of patients.

The study data were maintained and managed using SPSS for Windows (v12–v15); the analyses were carried out using SPSS and SAS (SAS Institute, Cary, NC). $\alpha = 0.05$ was used as the threshold for statistical significance for all analyses.

2.3. Effect of aHTN drugs on cognitive decline

To test the effects of different classes and combinations of aHTN drugs on the rate of cognitive decline in AD patients

Table 2 Baseline characteristics for set 1 (diuretics + CCB + RAAS vs. all other groups)

Group	Diuretics + CCB + RAAS	Other groups	χ^2/t	P value
N	24	593		
Age	77.3 (7.2)	76.3 (-7.6)	-0.54	0.54
Education	13.7 (3.2)	13.2 (3.3)	-0.7	0.48
Baseline MMSE (SD)	21.9 (4.7)	20.4 (5.2)	-1.39	0.16
Gender: female (%)	20 (83%)	397 (67%)	2.83	0.09
APOEE4 carrier (%)	11 (46%)	280 (47%)	0.02	0.89

Abbreviations: CCB, calcium channel blocker; MMSE, Mini-Mental State Examination; RAAS, renin-angiotensin-aldosterone system.

with hypertension (set 1), we implemented mixed-effect regression analysis with random intercept and trend. The nine aHTN treatment groups (see Table 1) and time after first record of aHTN drug use were used as the primary predictors for the power-transformed MMSE score. We also assessed the interaction term between the treatment group and time to test whether the treatment groups differed in their rates of cognitive decline. The analysis was conducted using an autoregressive variance-covariance matrix, and controlled for covariates including age at baseline, sex, years of education, and $APOE\varepsilon4$ carrier status.

2.4. Synergistic effects of aHTN drugs and ChEIs on cognitive decline

To test for the possible synergistic effects of aHTN drugs and ChEIs on the rate of cognitive decline, we divided the set 2 patients into two layers: ChEI users and nonusers. We first compared the time trends of MMSE decline in these two layers, and then investigated the effect of aHTN drug use on the MMSE decline in both layers using a mixed-effect regression model. Baseline characteristics including age, sex, years of education, $APOE\varepsilon4$ genotype, and MMSE score were adjusted for in the analyses, and an autoregressive variance-covariance matrix was used in the analyses.

2.5. Data collection for CVD and AD

The information on approved drugs, drugs in clinical trials, and protein targets associated with AD and CVD were gathered from various databases, including the Drug-Bank, ClinicalTrials.gov, BindingDB, AlzGene, PubChem, ChEMBL, Therapeutic Target Database, and SciFinder. The information from different sources was standardized with the same format, including protein full name, gene name, UniProt Entry ID, and Entry name. All the information was double checked by a second person, according to our data collection protocols [22]. To be noticed, CVDs included coronary artery diseases such as angina and myocardial infarction, stroke, hypertension, rheumatic heart disease, cardiomyopathy, heart arrhythmia, congenital heart disease, valvular heart disease, carditis, aortic aneurysms, peripheral artery disease, and venous thrombosis. The tables listing the disease-specific targets can be found on our website (http://cbligand.org/AD/target_list.php for AD and http://cbligand.org/CVD/target_list.php for CVD) [20,21].

2.6. Construction of disease-target network for AD and hypertension

To find the shared multiple drug targets in the overlapping pathways between AD and hypertension, which may point to the combinational treatment for these two coexisting medical conditions, we used our established AD (http://www. cbligand.org/AD) and CVD (http://www.cbligand.org/ CVD) databases for the analysis. From the CVD database, we only extracted the drug targets specifically for

Table 3 Baseline characteristics for set 2 patients by ChEI and aHTN drug use

aHTN drug use	No ChEI		ChEI			
	No	Yes	No	Yes	$\chi^2 F$	P value
N	40	46	169	174		
Education (SD)	13.2 (2.8)	13.1 (2.6)	14.8 (3.1)	13.6 (3.0)	21.93	<.0001
Age (SD)	75.7 (9.8)	77.6 (6.8)	72.7 (9.4)	75.9 (8.0)	18.25	<.0001
Baseline MMSE (SD)	19.4 (4.2)	22.2 (3.4)	21.5 (4.3)	21.2 (4.1)	10.59	<.0001
Gender: female (%)	30 (75.0)	36 (78.7)	90 (53.3)	108 (61.9)	2.926	.09
APOEε4 carrier* (%)	23 (57.5)	23 (50.0)	105 (62.1)	100 (57.5)	1.546	.238

Abbreviations: aHTN, antihypertensive; ChEI, cholinesterase inhibitor; MMSE, Mini-Mental State Examination.

*Missing data for APOEɛ4 genotype: n = 2, both in the ChEI-/aHTN+ group.

hypertension treatment to simplify the analysis. Most of the targets we included were the proteins with one or more drugs in the market or under investigation for the treatment of hypertension. We mainly focused on targets with confirmed associations with the disease validated by at least two different sources. Then we combined the target information from two disease-specific databases. The UniProt Entry name and the corresponding disease (AD or hypertension) were further used to map out a disease-target network (DTN) by using Cytoscape 3.1.2, an open-source program for visualizing complex networks.

2.7. Mechanism study of diuretics + CCB + RAAS combination using CSP-target mapping

To understand the molecular mechanism of the synergistic effect among the diuretics + CCB + RAAS combination on the cognitive decline in patients with AD, we carefully examined drug (most frequently used) from each of the aHTN drug classes. The representative drugs are hydrochlorothiazide (HCTZ) for diuretics, amlodipine for CCB, and losartan for RAAS inhibitor. Then, we predicted the ADrelated protein targets that each of the drugs potentially acts on by docking the drug molecule against the AlzPlatform target library [22] using our HTDocking program, a web-based software package for high-throughput docking (http://www.cbligand.org/HTDocking/). A docking score of >6 was used as a threshold for potential interactions. Then, we ranked the potential protein targets for each drug based on the docking scores, and searched in the literature for experimental validation results for the top 10 interactions. To visualize the results, a CSP-target mapping was constructed using SpiderPLot to construct DTN. The docking poses and detailed interactions were visualized using Py-MOL.

3. Results

3.1. Effect of aHTN medications on cognitive decline

Among the 617 patients who had been diagnosed with AD and hypertension (set 1), 399 (64.7%) patients had at least one record for using an aHTN agent. The number of

patients under different drug classes over time is shown in Table 1. Time zero is defined as the time for the first record of aHTN drug use in the database. For patients without hypertension and hypertensive patients who did not have any record of aHTN drug use, time zero is defined as the date of the first record in the database. The rate of cognitive decline in probable AD patients without HTN (n = 627) was not significantly different from that of those who had probable AD and HTN but were not taking aHTN medications (P = .14, see Fig. S1). A significant difference (P = .02) was found among different treatment groups in their trajectories of cognitive decline (see Fig. 1). The patient group under the combination of CCB, diuretics, and RAAS showed the lowest rate of cognitive decline, with virtually no decrease in the first 3 years.

To further confirm this observation, we compared the MMSE trajectories of patients with the CCB + diuretics + RAAS combination group against patients in all other

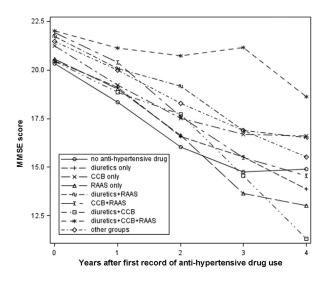


Fig. 1. The cognitive decline in set 1 patients by different aHTN treatment groups was analyzed using a mixed-effect regression model. Combination of diuretics, calcium channel blockers, and renin-angiotensin-aldosterone system inhibitors (diuretics + CCB + RAAS) was associated with the slowest rate of cognitive decline among all groups tested ($\Delta\beta = +1.46$, P < .0001). This effect was most prominent in the first 3 years of aHTN treatment. Abbreviations: aHTN, antihypertensive; MMSE, Mini-Mental State Examination.

groups in the first 3 years using the mixed-effect regression model with random intercept and slope. There was no significant difference between the CCB + diuretics + RAAS group and other groups in terms of the baseline characteristics (Table 2). However, the estimated slope of cognitive decline of all other groups was -2.15, whereas the slope of the CCB + diuretics + RAAS group was -0.69($\Delta\beta = +1.46$, P < .0001), indicating a significantly slower rate of decline.

3.2. Synergistic effect of aHTN medications and ChEI on cognitive decline

Table 3 shows the baseline characteristics of set 2 patients. In general, ChEI users received more education than nonusers. aHTN drug users were older compared with nonusers. Those who never used either ChEI or aHTN drugs also had the lowest MMSE score at baseline.

We first compared the rate of cognitive decline over the 24-month follow-up between the ChEI group and the non-ChEI group, controlling for baseline characteristics including age, sex, years of education, *APOE*e4 genotype, and MMSE score. The result showed that ChEI use had a trend of slowing MMSE decline (P = .068). Next, we compared the effect of aHTN medications in the ChEI group and non-ChEI group in set 2, respectively (see Fig. 2). In the non-ChEI stratum, the ChEI-/aHTN- patients had a slope of -2.09 (P < .0001), and the ChEI-/aHTN+ patients also had a similar trajectory ($\Delta\beta = -0.31$, P = .53). On the other hand, ChEI+/aHTN- patients had a similar slope to that of ChEI-/aHTN- patients ($\beta = -2.13$, P < .0001). Interest-

ingly, however, ChEI+/aHTN+ patients had a slope for cognitive decline of -1.57, indicating a significantly reduced rate of cognitive decline compared with the ChEI+/aHTN- patients (slope difference $\Delta\beta = +0.56$, P < .01).

3.3. Mechanism study using CSP-target mapping

Using our AD and CVD chemogenomics-guided CSPtarget mapping, we identified 128 molecular targets for the signaling pathways related to hypertension and 108 molecular targets related to AD (Fig. 3). Twenty-eight targets were found in the intersection between the two diseases, including several targets for RAAS inhibitors (angiotensin-converting enzyme [ACE], angiotensin receptor 1 [AGTR1], and AGTR2) and CCBs (CACNA1A, CACNB2, CACNA2D1, and CAC-NA2D4). We also identified the drugs targeting two important molecular targets that are related to aHTN treatment, lipoprotein lipase, and ACE (Table S1). These drugs have already been approved by the Food and Drug Administration or are in clinical trials for treating AD.

We docked the three representative drugs (HCTZ, amlodipine, and losartan) from the three aHTN drug classes against the AD-specific target library. From the docking results, we identified 43 potential targets for HCTZ, 38 potential targets for amlodipine, and 46 potential targets for losartan, whose docking scores with the corresponding aHTN drugs were ≥ 6 . We searched in the literature for the reported drug-target interactions among the top 10 targets

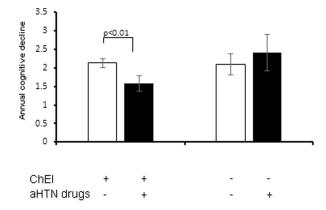


Fig. 2. Annual MMSE decline in AD cases with hypertension (set 2) by ChEI and aHTN drug use. Trajectories for cognitive decline among different groups in set 2 were analyzed using a linear mixed-effect regression model. Bars and error bars represent the coefficient β and standard error of the mean for the time variable in each group, respectively. In ChEI+ group, concomitant use of aHTN drugs was associated with significantly slower cognitive decline (slope difference $\Delta\beta = +0.56$, P < .01). The use of aHTN drug was not associated with a significantly different rate of cognitive decline in the ChEI- group ($\Delta\beta = -0.31$, P = .53). Note that "+" denotes drug users and "-" denotes nonusers. For example, ChEI+/aHTN- represents patients who used ChEIs but not aHTN drugs. Abbreviations: AD, Alzheimer's disease; aHTN, antihypertensive; ChEI, cholinesterase inhibitor; MMSE, Mini-Mental State Examination.

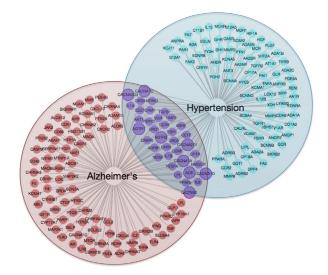


Fig. 3. Overlapping protein targets for Alzheimer's disease (red) and hypertension (cyan). Each node represents a protein target name that is associated with the disease(s) linked to it by a straight line. Targets highlighted in solid line circles indicate the drug classes that are shown to have potential synergetic effects by our data-mining analysis in clinical and molecular levels. Abbreviations: ACE, angiotensin-converting enzyme; AGTR1, angiotensin receptor 1; CACNA1A, calcium voltage-gated channel auxiliary subunit alpha1 A; CACNA2D1, calcium voltage-gated channel auxiliary subunit alpha2delta1; CACNA2D4, calcium voltage-gated channel auxiliary subunit alpha2delta4; CACNB2, calcium voltage-gated channel auxiliary subunit beta1; PPARγ, peroxisome proliferator–activated receptor gamma.

for each aHTN drug and constructed a CSP-target mapping analysis as shown in Fig. 4. For each aHTN compound, we selected two predicted protein targets (one experimentally validated and one nonvalidated) as examples; their docking poses and detailed DTN interactions are shown in Fig. 5. The CCB drug amlodipine has been predicted to form hydrogen bonds with its validated target, acetylcholinesterase (AChE) [26], in residues Gln71, Tyr72, Tyr124, Ser125, and Tyr137 (Fig. 5A). Amlodipine has also been predicted to form two hydrogen bonds with its nonvalidated target, the mitogenactivated protein kinase (MAPK), in residues Ala111 and Asp168 (Fig. 5B). Meanwhile, the diuretic drug HCTZ was predicted to form four hydrogen bonds with its validated target phosphodiesterase 4B (PDE4B) [27] in residues Tyr233, Tyr403, Met411, and Met431 (Fig. 5C). Three hydrogen bonds were predicted to form between HCTZ and its nonvalidated target, cyclooxygenase 2 (COX-2) in residues Phe381, Met522, and Ser530 (Fig. 5D). Finally, the RAAS drug losartan was predicted to form four hydrogen bonds with its validated target, the peroxisome proliferator-activated receptor gamma (PPARy) [28] in residues Cys285, Gln286, Tyr327, and Tyr473, respectively (Fig. 5E). With the nonvalidated target monoamine oxidase B (MAOB), Losartan was predicted to form four hydrogen bonds in residues Ile198, Gln206, Lys296, and Tyr 435 (Fig. 5F). The distances of the hydrogen bonds are all within 2.5 to 4.0 Å range. These results showed that these three representative aHTN medications may potentially act on a number of AD-related protein targets in addition to their primary targets, and a number of these interactions have been experimentally confirmed in the literature.

4. Discussion

We applied clinical data-mining analyses to investigate patients with AD in three decades of clinical observations and evaluated whether/which medication(s) used for HTN, a common coexisting disease of AD, is related to cognitive decline in patients diagnosed with AD. As shown previously, the use of aHTN medications was associated with a reduced rate of cognitive decline only in those patients who also used ChEI. This suggests that aHTN medications and ChEIs may produce a synergistic effect against cognitive decline in AD patients with hypertension. Moreover, patients under one specific combination of three classes of aHTN drugs, namely diuretics, CCB, and RAAS, were associated with the most significant reduction in the rate of cognitive decline compared with all other groups.

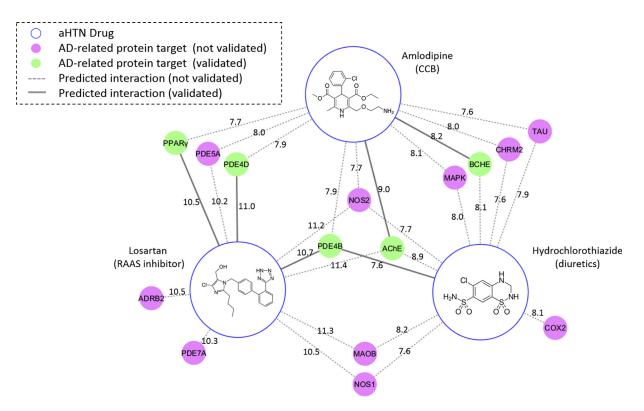


Fig. 4. Chemogenomics systems pharmacology (CSP) target mapping analysis for the molecular mechanism of HCTZ, amlodipine, and losartan for AD treatment. Each blue circle represents an aHTN drug, and each of the other nodes represents a predicted protein target either with experimentally validated binding affinities (green) or without experimental validation (magenta). Each edge connecting an aHTN drug and a protein target represents either an unconfirmed (dashed line) or a confirmed (solid line) drug-target interaction predicted by the HTDocking algorithm. The numbers on the edges represent the docking scores (predicted log K_i 's) of the drug-target interaction. Abbreviations: AD, Alzheimer's disease; AChE, acetylcholinesterase; ADRB2, β -2 adrenergic receptor; aHTN, antihypertensive; BChE, butyrylcholinesterase; CHRM2, cholinergic receptor muscarinic 2; COX, cyclooxygenase; MAOB, monoamine oxidase B; MAPK, mitogen-activated protein kinase; NOS, nitric oxide synthase; PDE, phosphodiesterase; PPAR γ , peroxisome proliferator–activated receptor gamma.

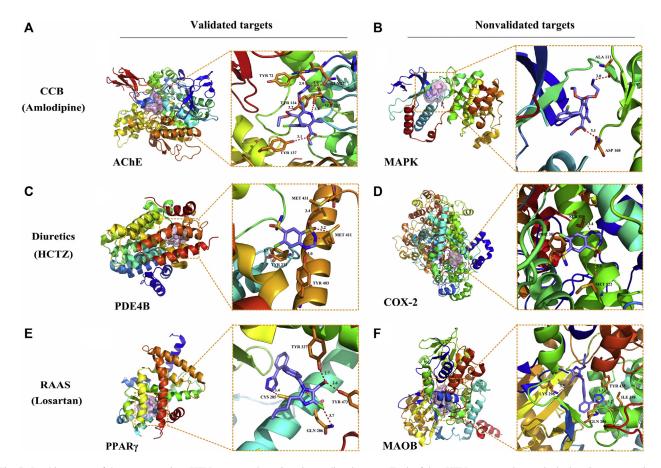


Fig. 5. Docking poses of the representative aHTN compounds against the predicted targets. Each of the aHTN compounds was docked against an experimentally validated target (left two columns—A, C, and E) and a nonvalidated target (right two columns—B, D, and F) predicted in our CSP-target mapping study. The first and third columns show the relative positions of the binding pockets (pink surfaces) in the proteins. The second and fourth columns show the detailed interactions between the aHTN compounds and the adjacent residues in the binding pocket of the protein targets. The aHTN compounds were shown as blue sticks, and the interacting residues were shown as orange sticks with residue numbers labeled. The polar interactions between the aHTN compounds and their bond lengths (Å) were labeled. Abbreviations: AChE, acetylcholinesterase; aHTN, antihypertensive; CCB, calcium channel blockers; COX, cyclooxygenase; CSP, chemogenomics systems pharmacology; HCTZ, hydrochlorothiazide; MAOB, monoamine oxidase B; MAPK, mitogen-activated protein kinase; PDE, phosphodiesterase; PPAR γ , peroxisome proliferator–activated receptor gamma; RAAS, renin-angiotensinaldosterone system.

Our study provides new and enriched data on the protective effect of aHTN medications against cognitive decline in patients with AD, and the results are congruent with the previous reports of decreased risk of cognitive decline and developing AD in hypertension-medicated patients and elderly people [13–18]. Our results are also in line with the most recent findings from the National Institutes of Health Systolic Blood Pressure Intervention Trial-Memory and Cognition IN Decreased Hypertension (SPRINT-MIND) [29] presented at the Alzheimer's Association International Conference 2018, which suggested that aggressive blood pressure control reduced the risk of mild cognitive impairment and dementia. Our study provides new evidence on the protective effect of aHTN medications against cognitive decline in patients with AD. This finding bears clinical importance because it suggests that aHTN medications are not only a prophylaxis against developing AD and cognitive decline, but can also be used as an add-on treatment.

Moreover, our study is the first to discover an optimal combination of different aHTN drug classes on the cognitive decline in AD patients with hypertension. According to a recent review article on the protective effect of aHTN medications against AD and cognitive decline [30], most studies have supported a preference for RAAS inhibitors [31–33] and CCBs [15,34] in protecting against cognitive decline, whereas some studies also reported no difference among the aHTN classes [35]. It is worth noting that although many studies also involved patients under combinations of aHTN drug classes, most of them tested the effect of different drug classes separately [16,17]. Few studies have reported different combination therapies that had a more pronounced protective effect against cognitive decline [13,15,32,36,37], but their patient populations are generally elderly patients or elderly patients identified with high-risk comorbidities, rather than those already diagnosed with AD. According to our results, there seems to be no difference

in cognitive decline when each of the drug classes were used alone, but the combination of RAAS + CCB + diuretics was significantly better. This observation does not contradict either side of the previous studies, but further points out a possibility that the preference for RAAS inhibitors and CCBs could be explained by the fact that they can produce a stronger protective effect when used in the specific combination. Our result also highlights the importance of considering specific drug combinations as a distinct group instead of testing each of its components separately.

In addition, our results uncovered that aHTN medications may have a synergistic effect with ChEIs on decreasing the rate of cognitive decline in AD patients with hypertension. First, we analyzed the effect of ChEI on the cognitive decline in AD patients with hypertension over a 24-month followup. The result showed that ChEI users had a marginally significant slower rate of cognitive decline (P = .068) compared with the non-ChEI users. We then compared the effect of aHTN medications in the ChEI users and nonusers, respectively (Fig. 2). The rate of cognitive decline in the ChEI monotherapy group and the ChEI-/aHTN+ group was not significantly different from that in the ChEI-/aHTNgroup, whereas the cognitive decline rate of ChEI+/aHTN+ group was significantly slower. To the best of our knowledge, our study is the first to discover the interaction between ChEIs and aHTN medications. As the first-line treatment for AD, the ChEIs have been shown by numerous RCTs and observational studies to have a significant effect on slowing cognitive decline in different AD populations [38-40]. However, it is important to note that most of these RCTs typically have a short time span of 24 to 26 weeks. Studies on the long-term effect (≥24 months) of the ChEIs on cognitive decline are few and reveal little to no clinically significant benefits on MMSE measures [41], whereas some studies did find a significant improvement using other end points such as nursing home admission [42]. The beneficial effect of ChEI monotherapy is also reported to have considerable interindividual variability [43]. A recent study [44] has searched for clinical factors that can predict ChEI response, yet it did not find any correlations between the pattern of ChEI response and any of the factors investigated, except for initial response in the first few months. Their scope of search included ChEI dose, APOE genotype, and CYP2D6 polymorphisms but did not include the common comorbidities and coadministered medications. In our study, we were the first to discover that the therapeutic effect of ChEIs may be dependent on concomitant use of aHTN medications in AD patients with coexisting hypertension. This synergistic effect suggests that aHTN drugs may serve as an add-on therapy for delaying cognitive decline in these patients.

To understand the molecular mechanisms behind the synergistic effects between AD and HTN medications observed in clinical settings, we applied our developed AD and CVD database-guided CSP-target mapping methodology techniques to map out DTNs for AD and HTN [20–23]. As shown in Fig. 3, we identified certain protein targets associated with two diseases, which indicated that aHTN drug(s) targeting these proteins could also have a direct effect on AD pathologic pathways. Such systems pharmacology DTN mapping analyses also suggested a molecular level synergism in accordance with the clinical level synergistic treatment of patients with AD with combinations of ChEI and aHTN drugs.

There have been some studies investigating the relationship between aHTN medication use and cognitive improvement. However, these articles only reported a reduced risk of AD in the population with hypertension treatment [15,45,46]. Because many previous studies have shown that elevated blood pressure is one of the major risk factors for AD, researchers make an assumption that aHTN medications may reduce the incidence of AD by controlling the blood pressure. Nevertheless, others have suggested that the aHTN drugs belonging to different drug classes may have specific protective effects in reducing AD risk [47]. In addition, some reports also found that controlling changes in blood pressure did not significantly alter the risk of AD dementia [46]. Thus, it is suggested that the aHTN drugs have a beneficial role in reducing the incidence of AD that is in addition to or independent from their benefit on blood pressure control.

The mechanism for the protective effect of diuretics against AD has not been widely studied. Although diuretics are a general class of aHTN medications with different mechanisms of action, further analysis of the medication history of the patients in diuretics + CCB + RAAS group in set 1 patients indicated that potassium-sparing diuretics and thiazides are the prevalent diuretics used in this combination. Some studies indicated that potassium-sparing diuretics had a potential to decrease AD risk because of a protective role of high potassium levels related to reduced vasoconstriction and chronic inflammation [45-47], presumably inhibiting their primary therapeutic via target mineralocorticoid receptor (NR3C2) [48]. However, some other studies found no significant differences between potassium-sparing diuretics and other nonsparing diuretics in decreasing AD risks [46]. On the other hand, the thiazide diuretics have been reported to inhibit carbonic anhydrases (CA1, CA2, and CA4) [49] in addition to their primary target SLC12A3 [50]. Although there has not been any study showing a connection between SLC12A3 and AD risk, inhibition of carbonic anhydrases has been reported to lead to a decreased release of cytochrome c from mitochondria to the cytoplasm, and hence reduce the amyloid beta $(A\beta)$ -induced neurotoxicity [51], which could be a potential mechanism for the protective effect of thiazide diuretics against AD.

The effect of CCBs in reducing AD incidence is controversial. Some epidemiologic studies showed that the use of CCB is related to a reduced risk of dementia [15,52]. Some others found no significant improvement in primary outcome measures [53–55]. Many CCBs were tested in clinical trial for AD treatment. Nimodipine and nilvadipine were shown to prevent cognitive decline in some trials, whereas other drugs within the same family failed [55]. Calcium homeostasis has been implicated in a role in AD. A β neurotoxicity results in an intracellular calcium influx via CACNA1C channels, which further leads to hyperphosphorylated tau and autophagy dysfunction [53,56]. In addition, L-type voltage-gated calcium channel (CAC-NA1C, CACNA1D, CACNA1S, and CACNA1F) blockers prevent neurotoxicity with the potential to reduce A β formation and maintain calcium homeostasis [53].

The aHTN therapies targeting RAAS [57], including ACE inhibitors (ACEI), angiotensin II receptor blocker (ARB) [58], and renin inhibitor, have been indicated to play a complicated role in AD pathogenesis. The beneficial effect of RAAS drugs to improve brain function was implicated in many studies. It was thought that the main mechanism of this improvement is to increase cerebral blood flow (CBF) by reducing vasoconstriction [47,59].

As for ACEIs, the effect is conflicted. On one hand, ACEIs have been reported to slow down the cognitive decline and dementia process [59,60]. On the other hand, ACE may be involved in the degradation of $A\beta$, thus ACEIs might contribute to $A\beta$ pathology and induce both the incidence [45] as well as the mortality [61] of AD. In addition, ACEIs augment levels of substance P, a substance degraded by ACE, which leads to an increased activity of another $A\beta$ degrading enzyme, neprilysin, thus having indirect beneficial effects for $A\beta$ clearance [59,62]. It makes the effect of ACEIs on AD more complicated [59].

ARBs also block angiotensin II signaling by acting on AGTR1 and AGTR2, and hence reduce vasoconstriction. This can result in increased CBF and improved cognitive function. Yet, recent reports suggested that the level of angiotensin II in the brain had a more important role in cognitive function. If ARB blocks the interaction between angiotensin II and the angiotensin II receptor 1, more angiotensin II will be converted to angiotensin III and then to angiotensin IV [47,63]. Angiotensin IV acts on c-Met receptor, which is associated with increases in longterm potentiation, synaptic plasticity, and CBF [64]. Angiotensin IV can also inhibit receptor insulin-regulated aminopeptidase, thus reducing the catalytic activity on vasopressin and oxytocin, both of which are related to memory consolidation [47,65,66]. The interaction between angiotensin IV and insulin-regulated aminopeptidase has also been reported to increase the uptake of glucose, further supporting its cognitive enhancing effects [47,66]. Another aspect of angiotensin II that is related to AD is the regulation of glycogen synthase kinase 3β (GSK3β). Angiotensin II can enhance the GSK3β level, which is thought to contribute to tau phosphorylation [67], inhibition of acetylcholine (ACh) release [68], and induced oxidative stress [58,59]. ARBs can reduce angiotensin II-mediated GSK3ß elevation, and its contribution to these cognitive-impairing factors, thus enhancing brain function [59].

In addition to the aforementioned pharmacologic effects, our CSP-target map (Fig. 4) also suggests that the diuretics + CCB + RAAS combination may directly act on several AD-related protein targets, with significant overlap. These targets include important proteins from several distinct pathologic pathways of AD, such as the enzymes involved in the clearance of neurotransmitters (AChE and butyrylcholinesterase [BChE]), enzymes related to oxidative stress (MAOB, NOS1, and NOS2), enzymes mediating cellular signaling pathways (PDE4B, PDE4D, PDE5A, PDE7A, MAPK, and COX-2), neurotransmitter receptors (cholinergic receptor muscarinic 2 [CHRM2] and β-2 adrenergic receptor [ADRB2]), a microtubule-associated protein (TAU), and a nuclear receptor related to neuroinflammation (PPAR γ). Subsequent literature study has indicated that a number of these predicted interactions have already been reported previously. Fig. 5 demonstrates the binding poses and detailed interactions between the three aHTN drugs and some of their predicted targets. These results suggest that the synergistic effect of the three aHTN medications on cognitive decline may be a net result of all these drugtarget interactions. The role of each predicted target in AD as well as its therapeutic effects is addressed subsequently.

The AChE has been predicted to interact with all three representative drugs from the three aHTN medication classes. The AChE catalyzes the breakdown of the neurotransmitter ACh at the synaptic clefts, and the inhibitors of AChE (ChEIs) are the first-line treatment for AD because of their ability to reverse the deficit of ACh in patients with AD. Among the three aHTN drugs, amlodipine has been reported to bind to AChE with a K_i of 0.19 μ M [26]. The same study also confirmed that amlodipine may interact with the BChE with a K_i of 0.11 μ M. BChE is another sub-type of cholinesterase inhibitor, which shares ~65% amino acid sequence identity with that of AChE and has also played an important role in AD [69].

Our CSP-target map analysis has also predicted that the MAOB, one of the important proteins involved in the oxidative stress mechanism [70], is a potential target for both losartan and HTCZ, which is congruent with the experiments reported in the literature. Oxidative stress is characterized by an imbalance in the redox state of the cells, which occurs as a result of the overproduction of reactive oxygen species or the defects in the antioxidant system [71]. Oxidative stress has been considered to be a major part of the pathophysiology of AD, causing significant tissue damage and neuronal death [72]. Localized in the outer mitochondrial membrane, MAOB catalyzes the oxidative deamination of multiple neurotransmitters including dopamine, serotonin, and norepinephrine, as well as exogenous amine species [73], during which it produces hydrogen peroxide (H2O2) as a side product, which is a potential source of oxidative stress [74]. It is possible that these two drugs together alleviate the cognitive symptoms of AD via reducing the oxidative stress generated by MAOB. The two isoforms of nitric oxide synthase, NOS1 and NOS2, also known as neuronal NOS and inducible NOS, have been reported to contribute to neuronal oxidative stress by producing nitric oxide (NO) [75], which is a free radical itself and can also form peroxynitrite when combined with superoxide anions [76]. In our CSP-target mapping analysis, all three aHTN drugs were predicted to target on NOS2, whereas losartan and HCTZ were predicted to act on NOS1, which may also partially explain the synergistic effect of the three aHTN medications against the cognitive decline in AD.

In addition, several isoforms of the phosphodiesterases (PDE4B, PDE4D, PDE5A, and PDE7A) have also been predicted to be the targets of the aHTN medications, and the inhibition of different PDE isoforms by losartan and HCTZ has been experimentally validated [27,77]. It has been previously shown in many AD animal models that specific PDE inhibitors improved memory performances by elevating the levels of cyclic adenosine monophosphate and/or cyclic guanosine monophosphate, and hence promoting the gene expression regulated by cyclic adenosine monophosphate response element-binding [78,79]. These genes are crucial for long-term memory formation and potentiation [80]. Therefore, the inhibition of PDEs provides another possible mechanism underlying the cognitive benefits of the diuretics, CCB, and RAAS inhibitor combination. The p38 MAPK is another cell signaling modulator that has been proposed as a target for treating AD [81]. Activation of the p38MAPK leads to the phosphorylation of serine and threonine residues in various kinases and transcription factors, and upregulates the inflammatory response to cellular stress [82]. Previous studies suggested that elevated MAPK activity is a significant contributing factor for the AD-related neuroinflammation in the brain, particularly in the microglia and astrocytes [83,84]. Furthermore, the p38MAPK has also been shown to have a direct role in the hyperphosphorylation of the tau protein, which is one of the hallmarks of AD pathology [85]. Therefore, the predicted interaction between MAPK and HCTZ and amlodipine may implicate a role of neuroinflammation in the mechanism underlying the protective effect of the diuretics + CCB + RAAS combination against cognitive decline in patients with AD. Finally, our CSP-target map analysis has also indicated a potential interaction between HCTZ and the COX-2 enzyme. The COX enzyme, also known as prostaglandin H synthase, catalyzes the synthesis of prostanoids such as prostaglandins, prostacyclin, and thromboxane, each of which plays an important role in the inflammation pathway. Because of its role in inflammatory reactions, the COX enzymes have been the primary targets for the nonsteroidal anti-inflammatory drugs [86,87]. Two selective COX-2 inhibitors, rofecoxib and naproxen, have been accessed in a phase 3 clinical trial for their potential to dampen the cognitive decline in patients with mild to moderate AD, but did not show significant benefit [88]. Given the importance of neuroinflammation in the pathology of AD, however, the author also commented in the article that additional trials using other nonsteroidal anti-inflammatory drugs may still be warranted.

Furthermore, the CHRM2 has been predicted to be the target of both amlodipine and HCTZ by the HTDocking program. The choline deficit is a major symptom of AD, and elevating the choline level by the use of ChEIs has long been the mainstream treatment strategy for AD. The CHRM2 receptor has been shown to regulate the release of ACh from cholinergic neurons, and CHRM2 selective inhibitors have been proposed as a promising strategy to treat AD by elevating the ACh levels [89]. On the other hand, the ADRB2 has been predicted to be a potential target of losartan. Activation of the ADRB2 receptor has been reported to enhance the activity of γ -secretase, and hence accelerate the production of amyloid plaque [90]. Therefore, as a potential inhibitor of ADRB2, losartan may also slow down the AD progression by modulating A β production.

The tau phosphorylation pathway is another promising target for the treatment of AD [91,92]. As a microtubuleassociated protein, tau participates in the assembly of tubulin into microtubules in the brain [93]. The hyperphosphorylation of tau protein has been found to be the major cause of the breakdown of microtubules and the formation of neurofibrillary tangles, which is known to be one of the hallmarks of AD [94]. Current strategies for blocking tau hyperphosphorylation mainly include targeting the upstream enzymes of tau phosphorylation, such as (1) inhibiting the kinases catalyzing tau phosphorylation, such as GSK3β, cyclindependent kinase 5 (CDK-5), and other kinases; (2) enhancing the activity of tau phosphatases (PP2A); and (3) enhancing the glycosylation of tau by the β -N-acetylglucosamine (O-GlcNAcylation) group [91]. Direct inhibition of tau protein has not yet been reported. In our CSP-target mapping analysis, HCTZ and amlodipine were both predicted to bind to the tau protein, which may potentially block its phosphorylation, and the consequent formation of neurofibrillary tangles.

Importantly, in our docking study, both losartan and amlodipine have been predicted to target the PPAR γ , and it has been found that a metabolite of losartan, EXP3179, has a partial agonist activity on PPAR γ [28]. PPAR γ is another protein target that has been extensively studied for its implications in AD. The PPARs are a family of three nuclear receptors (α , γ , and δ), each of which regulates a set of genes involved in lipid and energy metabolisms [95]. Because of the ability of PPAR γ to regulate both lipid and carbohydrate metabolisms, particularly the serum glucose levels and insulin sensitivity, the PPAR γ agonists (pioglitazone and rosiglitazone) have been developed into medications for treating type II diabetes mellitus [96]. Apart from that, the PPAR γ agonists have also been found to have a therapeutic potential for AD by targeting multiple aspects of AD pathology, including AB homeostasis, neuroinflammation, insulin sensitivity, energy metabolism, and lipid metabolism [97]. As a matter of fact, the PPAR γ agonist pioglitazone has entered phase III clinical trials for its effect in slowing the cognitive decline in patients with AD. Our CSP-target mapping revealed that these results may also explain the effect of the diuretics + CCB + RAAS combination against ADrelated cognitive decline.

AD is a multifactorial disease. The limitations of current treatments are that they target specific downstream neurochemical pathology whereas an upstream underlying mechanism remains to be unveiled. A combination medication that acts on a number of molecular and cellular pathologic pathways in AD, including A β accumulation, tau phosphorylation, chronic inflammation, oxidative stress, and impaired CBF, might have more beneficial effects compared with a single medication to treat AD [46]. Using these drugs in combination may help to achieve such effects.

We also point out that the present study has limitations. First, the patient number in set 2 was not large enough to allow further analysis for different aHTN drug classes. Therefore, interpretation of the molecular mechanisms underlying the possible synergism between hypertensive drugs and ChEIs is difficult. Second, we did not measure the blood pressure levels, which might have been an intermediate factor for the effect of aHTN drugs on cognitive decline. Although some studies reported that high blood pressure is a contributing factor to cognitive decline [98,99], other studies also reported a U-shaped relationship between blood pressure and cognitive impairment [100,101], suggesting that mildly elevated blood pressure may be beneficial in patients with AD, particularly in APOEe4 carriers [35]. Without blood pressure measurements, our study does not reveal the role of blood pressure in the therapeutic effect of aHTN medications on cognitive decline. Third, one recent study has pointed out that the cognitive benefits of ACEIs in patients with AD are dependent on rs1800764 and rs4291 genotypes [102]. Because we did not have the information regarding the rs1800764 and rs4291 variants, the genetic influence on the effectiveness of the combination therapy remains to be tested. Fourth, one potential factor that may contribute to the better cognitive outcome in patients treated with the three aHTN medications is that these patients might have visited their doctors more frequently, and consequently, they have better control of hypertension and other disorders that we did not examine here. Finally, the prediction of drug-target interactions using the CSP-target mapping analysis was carried out in silico using the HTDocking platform. The actual binding affinities for these interactions need to be experimentally validated in future studies.

5. Conclusions

Our clinical outcome analyses supported the protective effect of aHTN medications against cognitive decline in patients who have already been diagnosed with AD. The combined use of aHTN medications and ChEIs was associated with a significantly slower rate of cognitive decline compared with each of the drugs alone in patients with AD, suggesting a potential synergistic effect. These findings indicate that improving the vascular health of patients with AD can produce a cognitive benefit, and also suggest that the mechanisms of actions of aHTN medications may provide a novel approach for developing therapies against AD. Well-controlled clinical trials that test the effect of the combined use of ChEIs and different classes of aHTN drugs, especially the combination of diuretics, CCBs, and RAAS inhibitors, could be valuable for determining the specific mechanisms of the synergism.

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Supplementary data

Supplementary data to this article can be found online at https://doi.org/10.1016/j.trci.2018.09.001.

RESEARCH IN CONTEXT

- 1. Systematic review: Hypertension is an important risk factor for developing Alzheimer's disease (AD) in later stages of life; hypertension also etiologically contributes to the pathologies of AD.
- 2. Interpretation: Some antihypertensive (aHTN) drug classes had been shown to prevent or slow cognitive decline in elderly patients. In this study, we confirmed the effect of aHTN medications on slowing cognitive decline in AD cases and examined combinations of aHTN drugs that achieved synergistic effects against cognitive decline. This study also provides a molecular level interpretation for the synergism using a chemogenomics systems pharmacology approach and identified the protein targets of the aHTN drugs that may explain their anti-AD activities.
- 3. Future directions: The effect of aHTN medications on slowing cognitive decline needs further confirmation by randomized controlled trials. Further studies could also investigate whether aHTN medications have an indirect effect on cognitive decline through lowering blood pressure.

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