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# A novel BACE inhibitor isolated from *Eisenia bicyclis* exhibits neuroprotective activity against $\beta$ -amyloid toxicity

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## Abstract

Alzheimer's disease (AD) is a disturbing and advanced neurodegenerative disease and is characterized pathologically by the accumulation of amyloid beta ( $A\beta$ ) and the hyperphosphorylation of tau proteins in the brain. The deposition of  $A\beta$  aggregates triggers synaptic dysfunction, and neurodegeneration, which lead to cognitive disorders. Here, we found that FF isolated from an eatable perennial brown seaweed *E.bicyclis* protect against  $A\beta$ -induced neurotoxicity in neuroblastoma cells stably transfected with two amyloid precursor protein (APP) constructs: the APP695 cDNA (SH-SY5Y-APP695swe). The FF demonstrated strong inhibitory activity for  $\beta$ -secretase ( $IC_{50}$  16.1  $\mu$ M) and its inhibition pattern was investigated using Lineweaver-Burk and Dixon plots, and found to be non-competitive. Then, we tested whether FF could inhibit production of  $A\beta$  in SH-SY5Y-APP695swe. FF inhibited the production of  $A\beta$  and soluble-APP, residue of APP from cleaved APP by  $\beta$ -secretase. Our data show that FF can inhibit the production of  $A\beta$  and soluble-APP $\beta$  via inhibition of  $\beta$ -secretase activity. Taken together these results suggest that FF may be worthy of future study as an anti-AD treatment.

**Keywords:**  $\beta$ -Secretase inhibitory activity, *Eisenia bicyclis* ethanol extract, Fucofuroeckol-b,  $\beta$ -Amyloid, Amyloid precursor protein, Alzheimer's disease

## Background

Alzheimer's disease (AD) is a neurodegenerative disease, and its prevalence is increasing rapidly in the elderly population. Dementia is the most prevalent symptom of AD, which is also characterized by progressive cognitive decline, memory impairment, vast neuronal loss, decline in cholinergic transmission, and the presence of senile plaques. AD is pathologically characterized by the presence of extracellular deposition of plaques comprised of amyloid- $\beta$  ( $A\beta$ ) peptide and neurofibrillary tangles (NFTs) comprised of hyperphosphorylated-tau protein, accumulating evidence suggests that these abnormal protein deposits are unlikely the causative events in AD as  $A\beta$  plaque or NFT volume poorly correlate with the severity of dementia (Šimić et al. 2016). Moreover, the degree of dementia in postmortem brains of AD patients correlates more closely with the level of soluble

oligomers of  $A\beta$  species, especially in hippocampal and cortex regions associated with learning and memory function.  $A\beta$  peptides are proteolytic products of the amyloid precursor protein (APP) and are sequentially cleaved by  $\beta$ - and  $\gamma$ -secretases (Murphy and LeVine 2010). Although  $A\beta$  peptides of varying length are produced,  $A\beta$ -42 is considered to be comparatively more amyloidogenic and readily assembles into soluble oligomers and consequent fibril deposits. Instead,  $A\beta$  oligomers termed as  $A\beta$ -derived diffusible ligands, are thought to induce synaptic loss and progressive cognitive decline in AD (Masters and Selkoe 2012).

Currently, marine plants are intriguing materials as nutraceuticals and pharmaceuticals, given their numerous biological and phytochemical benefits. Marine algae can be classified into three classes based on their pigmentation, namely brown, red, and green algae, which are referred to as *Phaeophyceae*, *Rhodophyceae*, and *Chlorophyceae*, respectively (Khan et al. 2010). Since the 1940s, production of algal polysaccharides has attained commercial significance through their application as thickening and gelling agents for various food and

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industrial applications (Burtin 2003). Researchers have revealed that marine algae-originated compounds exhibit various biological activities, including anticoagulant (Athukorala et al. 2007), anti-viral (Artan et al. 2008), antioxidant (Heo et al. 2005), anti-allergic (Li et al. 2008), anti-cancer (Kong et al. 2009), and anti-inflammatory (Kim et al. 2009) activities, among others. Furthermore, several scientific studies have provided insight into the putative neuroprotective properties of marine algae (Zarros 2009). *Eisenia bicyclis* (Kjellman) is a perennial brown alga, of the family *Laminariaceae*. It is extensively distributed in Korea and Japan, and is especially abundant in Ulleung and on Dok Island in Korea. It is frequently used as a dietary alternative for *Laminaria japonica*, along with *Ecklonia stolonifera* and *Ecklonia cava*. This species contains several bioactive components, including phlorotannins, polysaccharides, pyropheophytin, sterol, lipids, tripeptides, and oxylipins (Noda et al. 1989; Kurata et al. 1990; Kousaka et al. 2003; Okada et al. 2004). In particular, phlorotannins produced by the polymerization of phloroglucinol (1,3,5-trihydroxybenzene) (Ragan and Glombitza 1986), have been reported to possess various biological activities, such as antioxidant (Kang et al. 2003), anti-inflammatory (Shin et al. 2006), and  $\beta$ -secretase1 (BACE1) inhibitory (Jung et al. 2010) activities. Although several reports on the cholinesterase and  $\beta$ -secretase inhibitory activities of brown algae and their phlorotannins have been reported, the neuroprotective effects of *E. bicyclis* and its phlorotannins on A $\beta$ -induced neurotoxicity have not yet been studied. Natural products from brown algae effectively modulate the type of neurodegenerative changes seen in AD, suggesting therapeutic potential. The present work focuses on the chemistry of polyphenols and their role in modulating APP processing. We also provide new hypotheses on how these therapeutic molecules may modulate APP processing, prevent A $\beta$  aggregation.

Therefore, the aims of this study were to study the potential of *E. bicyclis* and its isolated  $\beta$ -secretase inhibitor to protect against A $\beta$ -induced neurotoxicity, and to reduce A $\beta$  in neuroblastoma cells.

## Material and methods

### Sample preparation

Brown alga (*E. bicyclis*) was collected from the coast of Ullengdo, Ullengdo Island, South Korea, in July 2013. The samples were washed three times with tap water to remove salt, epiphytes, and sand attached to the surface and then rinsed with fresh water, dried under shade, and homogenized with a grinder.

### Preparation of ethanolic extract and fractions

The algal powder (1 kg) was extracted with 3 L of 80% ethanol at 37 °C for 24 h and then filtered. After filtration, the ethanol extracts were dried using a rotary

evaporator under vacuum. Each extract was partitioned with organic solvents to yield n-hexane, ethyl acetate, and H<sub>2</sub>O fractions (Fig. 1a).

### Isolation of compounds

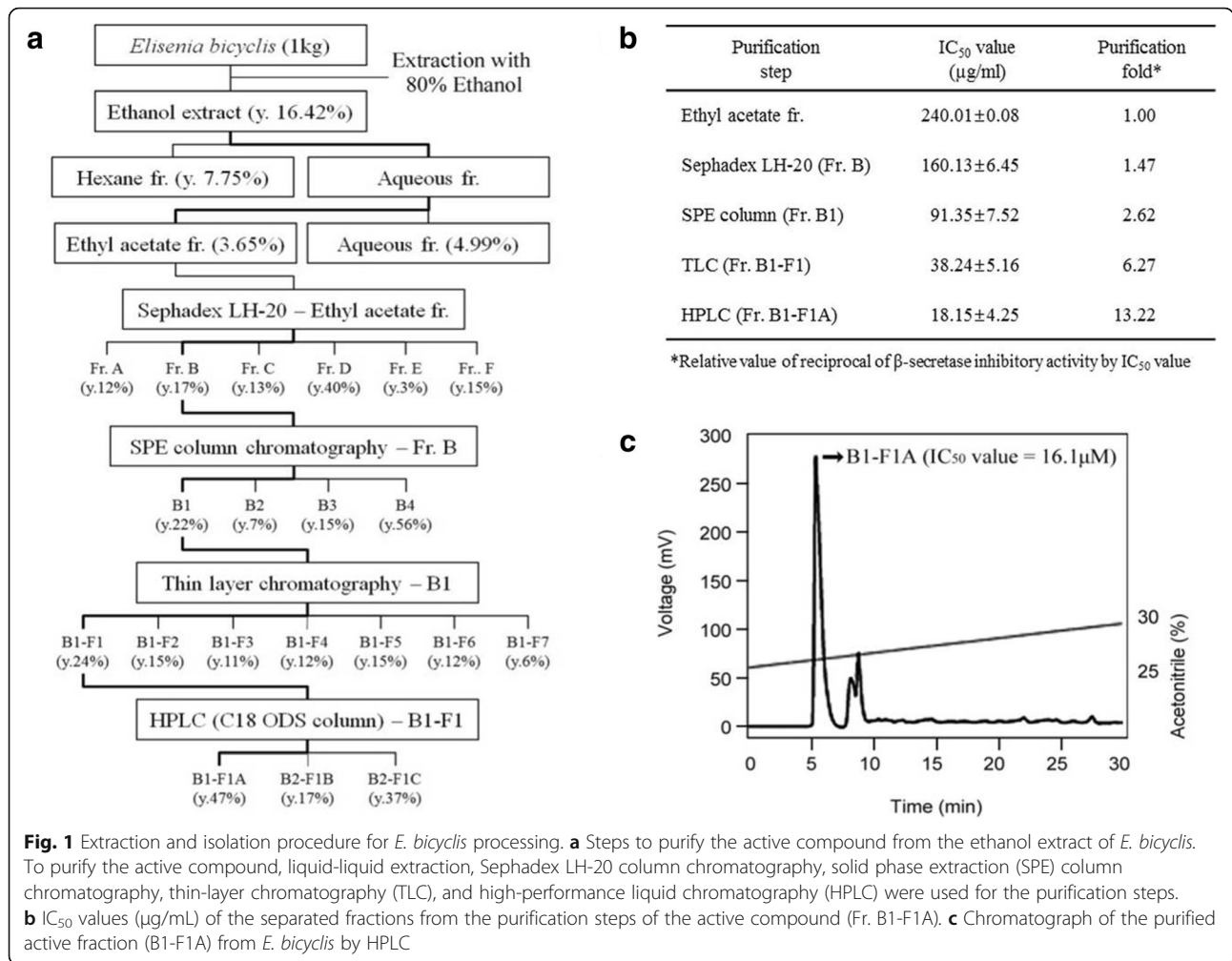
The active fraction was purified using a silica gel-filled Sephadex LH-20 (Sigma-Aldrich, MO, USA) chromatography column, and the column was eluted with methanol at a flow rate of 1.5 mL/min. The isolated fraction from the Sephadex LH-20 column was isolated using a solid phase extraction (SPE) column (Hi-Load 18, 5 g, 20 mL, Chrom Tech, Inc.) and a thin-layer chromatography (TLC) (0.25 mm Silica gel 60 F254, Merck, Co.) preparative separation. The active fraction was separated by high-performance liquid chromatography (HPLC) (Agilent 1100 series, USA) on a C18 ODS column (4.6 × 250 mm, 5  $\mu$ m, Shodex, Tokyo, Japan) at a flow rate of 1.0 mL/min at 215 nm.

### NMR analysis

All non-aqueous reactions were performed in flame-dried glassware under a nitrogen atmosphere with magnetic stirring. NMR spectra were obtained on a JOEL Lambda 300 spectrometer and recorded at 300 MHz for <sup>1</sup>H (75 MHz for <sup>13</sup>C) with CDCl<sub>3</sub> as a solvent and (CH<sub>3</sub>)<sub>4</sub>Si (<sup>1</sup>H) or CDCl<sub>3</sub> (<sup>13</sup>C, 77.0 ppm) as internal standards, unless otherwise stated. All <sup>31</sup>P NMR chemical shifts are reported in ppm relative to 85% H<sub>3</sub>PO<sub>4</sub> (the external standard). FT-IR spectra were recorded on a JASCO FR-IR 460 series unit. High-resolution FAB mass spectra were attained from a hybrid LC-quadrupole-TOF tandem mass spectrometer at Gangneung-Wonju National University.

### Measurement of $\beta$ -secretase inhibitory activity

$\beta$ -Secretase inhibitory activity was measured following Johnston's method (Johnston et al. 2008), using a commercially available fluorogenic substrate, MCA-EVKMD AEFK-(DNP)-NH<sub>2</sub>. This substrate is analogous to the wild-type APP sequence and is derivatized at its N-terminus with a fluorescent 7-methoxycoumarin-4-yl acetyl (MCA) group and at its C-terminal lysine residue with a 2,4-dinitrophenyl (DNP) group. In the whole peptide, the fluorescence of the MCA group is eliminated by internal quenching from the DNP group. Once cleaved by  $\beta$ -secretase activity, the MCA fluorescence can be detected. Tests were carried out in 96-well black plates using a spectrofluorometer (Tecan, Switzerland).  $\beta$ -Secretase and  $\beta$ -secretase substrate I were incubated in a final volume of 200  $\mu$ L of assay buffer (50 mM sodium acetate, pH 4.5). The hydrolysis of  $\beta$ -secretase substrate I was followed at 37 °C for 30 min by measuring the associated increase in fluorescence. Readings (excitation 325 nm, emission 393 nm) were obtained every 60 s. The inhibition ratio was calculated using the following equation: inhibition (%) = [1 - {(S



**Fig. 1** Extraction and isolation procedure for *E. bicyclis* processing. **a** Steps to purify the active compound from the ethanol extract of *E. bicyclis*. To purify the active compound, liquid-liquid extraction, Sephadex LH-20 column chromatography, solid phase extraction (SPE) column chromatography, thin-layer chromatography (TLC), and high-performance liquid chromatography (HPLC) were used for the purification steps. **b** IC<sub>50</sub> values (μg/mL) of the separated fractions from the purification steps of the active compound (Fr. B1-F1A). **c** Chromatogram of the purified active fraction (B1-F1A) from *E. bicyclis* by HPLC

$-S_0)/(C-C_0) \times 100]$ , where *C* is the fluorescence of a control (enzyme, assay buffer, and substrate) after 60 min of incubation, *C*<sub>0</sub> is the fluorescence of the control at zero time, *S* is the fluorescence of test samples (enzyme, sample solution, and substrate) after 60 min of incubation, and *S*<sub>0</sub> is the fluorescence of the relevant test sample at zero time. All data are the mean of triplicate experiments.

**Kinetic analysis**

To investigate the mode of β-secretase inhibition by the purified inhibitor, the purified inhibitor was added to each reaction mixture. The inhibition constants (*K<sub>i</sub>*) of β-secretase inhibitors were calculated using Dixon plots. To determine the kinetics of purified compounds, inhibition rates were measured at four concentrations of the substrate (500, 750, 1000, and 1500 nM). For the Dixon plots, data were acquired as mean values of 1/*V* (reaction velocity), the inverse of the intensification of fluorescence intensity per min (min/relative fluorescence units, RFU) of three independent tests with different concentrations

(0, 2.5, 5.0, and 10 μM) of inhibitor. The assay was carried out in the presence of purified inhibitor.

**Cell culture**

Human neuroblastoma SH-SY5Y cells were maintained in Dulbecco’s modified Eagle medium nutrient mix F12 medium (GIBCO, Carlsbad, CA, USA) containing L-glutamine and 15 mM HEPES, supplemented with 10% fetal bovine serum (GIBCO, Carlsbad, CA, USA) and 100 U/mL penicillin-streptomycin at 37 °C, under 5% CO<sub>2</sub>. SH-SY5Y-APP695swe cells stably transfected with the human Swedish mutation were generated as previously described (Jämsä et al. 2011). The cDNA encoding human APP695swe was inserted into the expression vector pIRESHyg (Clontech) before being stably transfected into SH-SY5Y cells. DNA (30 μg) was introduced into the cells by electroporation in 4-mm cuvettes with a pulse of 250 V and 1650 μF using an ECM630 electroporator (BTX Harvard Apparatus, Holliston, MA). Selection for cells containing the required construct was performed in

normal growth medium with 0.15 mg/mL hygromycin B (Invitrogen, Paisley, UK). Cells were grown to 90–100% confluency, washed twice with phosphate-buffered saline solution (pH 7.4, GIBCO, Carlsbad, CA, USA), and incubated in 10 mL of serum-free Opti-MEM (Invitrogen, ON, USA) for 24 h. The conditioned medium was harvested, and 5 mL was concentrated to 200  $\mu$ L using 10-kDa cutoff Vivaspin filtration columns (Millipore, Billerica, MA). The remaining 5-mL conditioned media sample from the SH-SY5Y cells was used for analysis of A $\beta$ 42 using an ELISA kit (Invitrogen, ON, USA).

#### A $\beta$ -induced toxicity on SH-SY5Y cells

On the experiment day, the culture media were removed, and cells were incubated at 37 °C in fresh culture media and exposed to either A $\beta$ 42 for 24 h, in the presence or absence of a  $\beta$ -secretase inhibitor. A $\beta$ 42 was dissolved by briefly vortexing in distilled water containing ammonium hydroxide 0.02% (*v/v*) and stored at –20 °C before use. Cells were also treated with the inactive reverse sequence peptide A $\beta$ 42 to confirm the specificity of the toxic effects of the fragment A $\beta$ 42. Cell viability was assessed 24 h later using the 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyl tetrazolium bromide (MTT) colorimetric assay.

#### MTT assay

To evaluate the toxicity of the active sample, cells were seeded in 96-well plates ( $5 \times 10^4$  cells/mL) and incubated with varying sample concentrations (6.5, 15, 25, and 50  $\mu$ M) for 24 h. The cell culture media were changed immediately before the MTT assay to avoid interaction between MTT and the sample, and 10  $\mu$ L MTT (5 mg/mL, Sigma-Aldrich, MO, USA) solution was added, followed by incubation for 4 h. The supernatant was removed, and 100  $\mu$ L DMSO was added to dissolve the dye crystals. Absorbance was read at 490 nm, and the percentage of viable cells was calculated.

#### Western blot

Cells were harvested, and proteins were prepared by direct extraction on ice in Tricine sample buffer (Bio-Rad Laboratories, Hercules, CA, USA) containing a protease inhibitor cocktail (Roche, Indianapolis, IN, USA), followed by centrifugation at 4 °C for 5 min. The supernatant protein concentration was measured using a BCA Protein Assay Kit (Thermo Scientific, Waltham, MA, USA). The protein samples were aliquoted and then frozen at –80 °C. Approximately 40  $\mu$ g of total protein was loaded into the wells in 8–12% sodium dodecyl sulfate (SDS)-polyacrylamide gels for determination of full-length APP or in 16% SDS–Tricine polyacrylamide gels for sAPP $\beta$  and A $\beta$  blots. Afterward, the proteins were transferred onto polyvinylidene fluoride (PVDF) membranes (Millipore, Billerica, MA, USA). After protein transfer, the PVDF membranes

were blocked with 5% dry nonfat milk in Tris-buffered saline plus 0.05% Tween-20 (TBST) for 1 h at room temperature and then incubated with primary antibodies against the C-terminus of APP (1:2000, A8717), sAPP $\beta$  (3  $\mu$ g/mL, SIG-39138), A $\beta$  (1:1000, B-4, Santacruz Biotech), and  $\beta$ -actin (1:2000, Sigma-Aldrich, MO, USA) in TBST containing 5% dry nonfat milk overnight at 4 °C and then for 30 min at room temperature. The membranes were washed with TBST and incubated with horseradish peroxidase-conjugated secondary antibodies for 2 h at room temperature. The immunoreactive proteins were detected with ECL Advance Western Blotting Detection Reagents (Amersham Biosciences, Piscataway, NJ, USA). All bands were standardized to  $\beta$ -actin and then compared and quantified with ImageJ software (National Institutes of Health, Bethesda, MD, USA).

#### Statistical analyses

Data were analyzed for statistical significance using analysis of variance (ANOVA) followed by Dunnett's multiple comparison test with SPSS software (version 14). All values obtained from three different experiments were expressed as the mean value  $\pm$  SEM.

## Results and discussion

#### Isolation of a $\beta$ -secretase inhibitor using ethanol extraction and partitioning into solvent fractions

A novel  $\beta$ -secretase inhibitor from *E. bicyclis* was purified using multiple purification methods include organic solvent extraction, LH-20 Sephadex chromatography, thin layer chromatography (TLC), solid phase extraction (SPE) and high performance chromatography (Fig. 1a). Among the fractions separated, the active fractions were selected through  $\beta$ -secretase inhibitory activity assay.

As shown in Table 1, ethanol extract or its fractions isolated by hexane, ethyl acetate and H<sub>2</sub>O fraction exhibited  $\beta$ -secretase inhibitory activity with an IC<sub>50</sub> value of 0.70, 1.02, 0.24 and 2.12 mg/mL, respectively. Among the fractions, the ethyl acetate fraction had the highest  $\beta$ -secretase inhibitory activity.

The ethyl acetate fraction was separated on a silica gel-filled Sephadex LH-20 column into six fractions (A–F). Among the separated fractions, fraction B had the highest  $\beta$ -secretase inhibitory activity, with an IC<sub>50</sub> value

**Table 1** IC<sub>50</sub> values of  $\beta$ -secretase inhibitory activity of the ethanol extract and its individual fractions from *E. bicyclis*

Sample	IC <sub>50</sub> value (mg/mL)*
Ethanol extract	0.70 $\pm$ 0.21 <sup>b</sup>
Hexane fraction	1.02 $\pm$ 0.11 <sup>c</sup>
Ethyl acetate fraction	0.24 $\pm$ 0.08 <sup>a</sup>
H <sub>2</sub> O fraction	2.12 $\pm$ 0.27 <sup>d</sup>

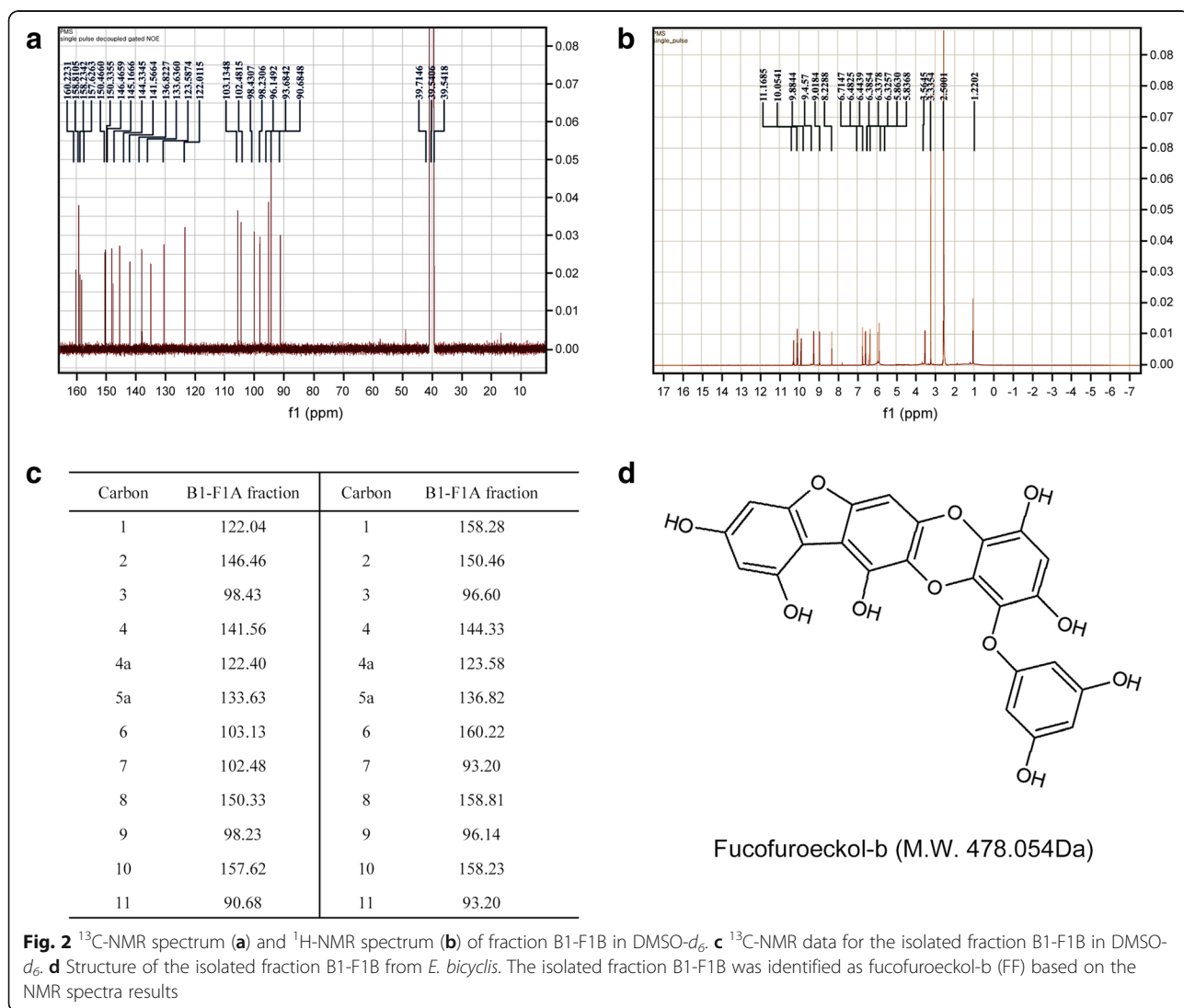
\*Values (mean  $\pm$  S.E.M. of three replications) not sharing a common superscript letter are significantly different ( $p < 0.05$ )

of 0.16 mg/mL (Fig. 1b). Fraction B from the Sephadex LH-20 column was then purified using an SPE column. In addition, that purified fraction was further isolated using a TLC preparative separation. The TLC purification was carried out in developing solvent (n-Hexane-ethyl-acetate-acetone (6:5:1, v/v)) and scanned under wavelength UN light (365 nm). Finally, an active fraction, B1-F1A, was purified from the B1-F1 fraction from TLC separation by HPLC. The  $\beta$ -secretase inhibitory activity of B1-F1A was purified over 13.2-fold from the ethyl acetate fraction using a purification of five steps (Fig. 1b). The purified fraction possessed  $\beta$ -secretase inhibitory activity with an  $IC_{50}$  value of 16.1  $\mu$ M (Fig. 1c) and exhibited a molecular weight of 478.054 Da according to Q-TOF-MS (data not shown). The structure of the purified fraction B1-F1A was identified by comparing its NMR spectral data with those in the existing literature.

**Identification of compounds isolated from *E. bicyclis***

Figure 2 shows the spectroscopic properties of the purified fraction, B1-F1A:  $C_{24}H_{14}O_{11}$ ;  $^1H$ -NMR (300 MHz,  $DMSO-d_6$ )  $\delta$  11.58 (1H, s, 14-OH), 10.05 (1H, s, 4-OH), 9.88 (1H, s, 10-OH), 9.45 (1H, s, 2-OH), 9.11 (2H, s, 3' 5'-OH), 8.22 (1H, s, 8-OH), 6.71 (1H, s, H-13), 6.48 (1H, d,  $J = 1.1$  Hz, H-11), 6.34 (1H, s, H-3), 6.38 (1H, d,  $J = 1.5$  Hz, H-9), 5.86 (1H, s, H-4'), 5.83 (2H, d,  $J = 1.5$  Hz, H-2, 6');  $^{13}C$ -NMR (300 MHz,  $DMSO-d_6$ )  $\delta$  160.22 (C-1'), 158.81 (C-3, 5'), 158.23 (C-11a), 157.62 (C-10), 150.46 (C-12a), 150.33 (C-8), 146.46 (C-2), 144.33 (C-14), 141.56 (C-4), 136.82 (C-15a), 133.63 (C-5a), 123.58 (C-14a), 122.4 (C-4a), 122.04 (C-1), 103.13 (C-6), 102.48 (C-7), 98.43 (C-3), 98.23 (C-9), 96.6 (C-4'), 96.14 (C-13), 93.2, (C-2',6') 90.68 (C-11).

The molecular structure of the purified compound  $C_{24}H_{14}O_{11}$  based on the NMR spectra data is shown in Fig. 2a, b. The proton NMR ( $^1H$ -NMR) spectrum



**Fig. 2**  $^{13}C$ -NMR spectrum (a) and  $^1H$ -NMR spectrum (b) of fraction B1-F1B in  $DMSO-d_6$ . c  $^{13}C$ -NMR data for the isolated fraction B1-F1B in  $DMSO-d_6$ . d Structure of the isolated fraction B1-F1B from *E. bicyclis*. The isolated fraction B1-F1B was identified as fucofuroeckol-b (FF) based on the NMR spectra results

contained signals characteristic of active aromatic protons. AB2 system had signals at  $\delta$ -6.71 (1H, s) and 6.47 (1H, d,  $J$  = 1.10 Hz),  $\delta$ 6.25 (1H, d,  $J$ -1.46 Hz) and 5.76 (2H, d,  $J$  = 1.46 Hz), and two singlets at  $\delta$ 6.29 (1H, s) and 5.83 (1H, s), as well as eight phenolic hydroxyl signals at  $\delta$ 10.05, 9.88, 9.76, and 8.22. The carbon-13 nuclear magnetic resonance ( $^{13}\text{C}$ -NMR) spectrum of the purified compound indicated the presence of 7 methanes, 15 O-bearing aromatic carbons, and 2 quaternary carbons ( $\delta$ 103.1, 102.4). The compound is composed of four benzene rings that are joined together via a furanoid, a 1,4 dioxin structure, and a phenyl ether bridge. The chemical shifts of C-5a, C-13, C-14, and C-14a were very close to those of the subsequent signals of eckol, and those of the two other carbons [C-6 ( $\delta$ 103.1) and C-12a ( $\delta$ 150.5)] were different from the subsequent signals.

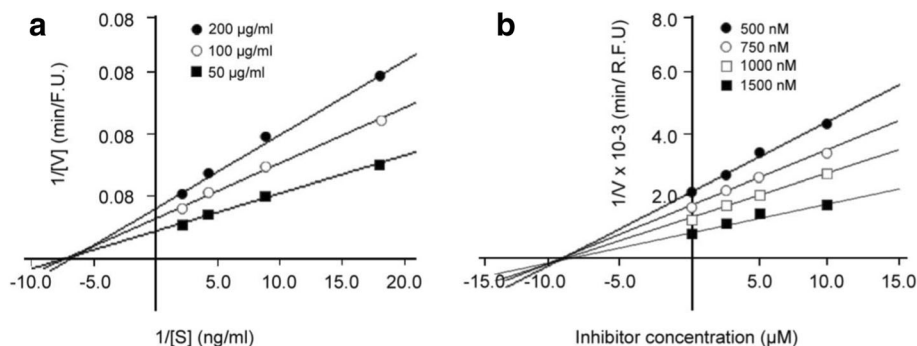
Thus, the purified compound had to be an eckol switched with phloroglucinol via a biaryl bond at C-9. The removal of one molecule of water from the hydroxyl groups at C-2 of the phloroglucinol and C-8 of the eckol unit must then have caused the creation of the furan ring. These data indicate that this compound contains an eckol moiety and that C-12a or C-13 can be altered or can exist close to the variation. Therefore, the purified compound was identified as fucofuroeckol-b (FF) based on the NMR spectrum results.

FF was first isolated as its free form from natural sources from *E. bicyclis*, although the hepta-acetate of FF has been previously isolated from *Eisenia arborea* (Glombitza and Gerstberger 1985). The  $\beta$ -secretase inhibition pattern of FF from *E. bicyclis* was investigated using Lineweaver-Burk plots and Dixon plots, and it was found to be non-competitive (Fig. 3a). Thus, it could bind with both the enzyme and the enzyme-substrate complex and interact independently of the substrate. In addition, the inhibitor did not change the substrate-enzyme binding affinity. With increasing concentrations, a sequence of lines was observed that had a shared intercept on the  $1/[S]$  axis

but different slopes. To determine the manner of  $\beta$ -secretase inhibition, kinetic analysis was performed using different substrate (500, 750, 1000, and 1500 nM) and inhibitor concentrations. The inhibition constants ( $K_i$ ) were obtained by analysis of the Dixon plots, in which the value of the  $x$ -axis denoted  $K_i$ . The Dixon plot is a graphical approach [plot of  $1/\text{enzyme velocity}$  ( $1/V$ ) against inhibitor concentration] to determine the type of enzyme inhibition, and it can be used to determine the dissociation or  $K_i$  for the enzyme-inhibitor complex (Cornish-Bowden and Eisenthal 1974). In the case of competitive inhibition, the  $x$ -axis indicates  $K_i$  when  $1/V = 1/V_{\text{max}}$ . In non-competitive inhibition, the  $x$ -axis indicates  $K_i$  when  $1/V = 0$ . As shown in Fig. 3b, FF ( $K_i = 10.1$ ) showed a non-competitive inhibition pattern in the Dixon plots. Typically, the lower the  $K_i$  value, the stronger the binding with the enzyme, and the more efficient an inhibitor is; thus, the results indicated that FF may be a strong candidate for a  $\beta$ -secretase inhibitor. Jung et al. (2010) reported that various phlorotannins demonstrated significant and non-competitive inhibition against  $\beta$ -secretase: dioxinodehydroeckol ( $K_i = 8.0$ ), eckol ( $K_i = 13.9$ ); phlorofurofuoceckol-A ( $K_i = 1.3$ ), triphloroethol-A ( $K_i = 12.1$ ), and 7-phloroethol ( $K_i = 7.2$ ). The  $K_i$  value of FF was higher than that of eckol and triphloroethol but lower than that of phlorofurofuoceckol-A. However, they all exhibited a non-competitive inhibition pattern. The *E. bicyclis* extract and the phlorotannins contained in it likely have value for the development of therapeutic and preventive agents for AD.

#### Cell cytotoxicity of FF in neuronal cells

We tested the toxicity of compound FF on SH-SY5Y cells overexpressing APP695swe. Cells were treated with increasing concentrations of FF (15, 25, 50, and 100  $\mu\text{g}/\text{mL}$ ) for 24 h, and cell viability was assessed by MTT assay. Cells treated with concentrations of the compound up to 15  $\mu\text{g}/\text{mL}$  did not exhibit reduced viability,



**Fig. 3** **a** Lineweaver-Burk plots for the inhibition of  $\beta$ -secretase in the presence of FF. Concentrations of FF for the curves were 50, 100, and 200  $\mu\text{g}/\text{mL}$ , using the substrate as the control. **b** Dixon plots for the inhibition of  $\beta$ -secretase by FF in the presence of different concentrations of the substrate: 500 nM (black circle), 750 nM (white circle), 1000 nM (white square), and 1500 nM (black square)

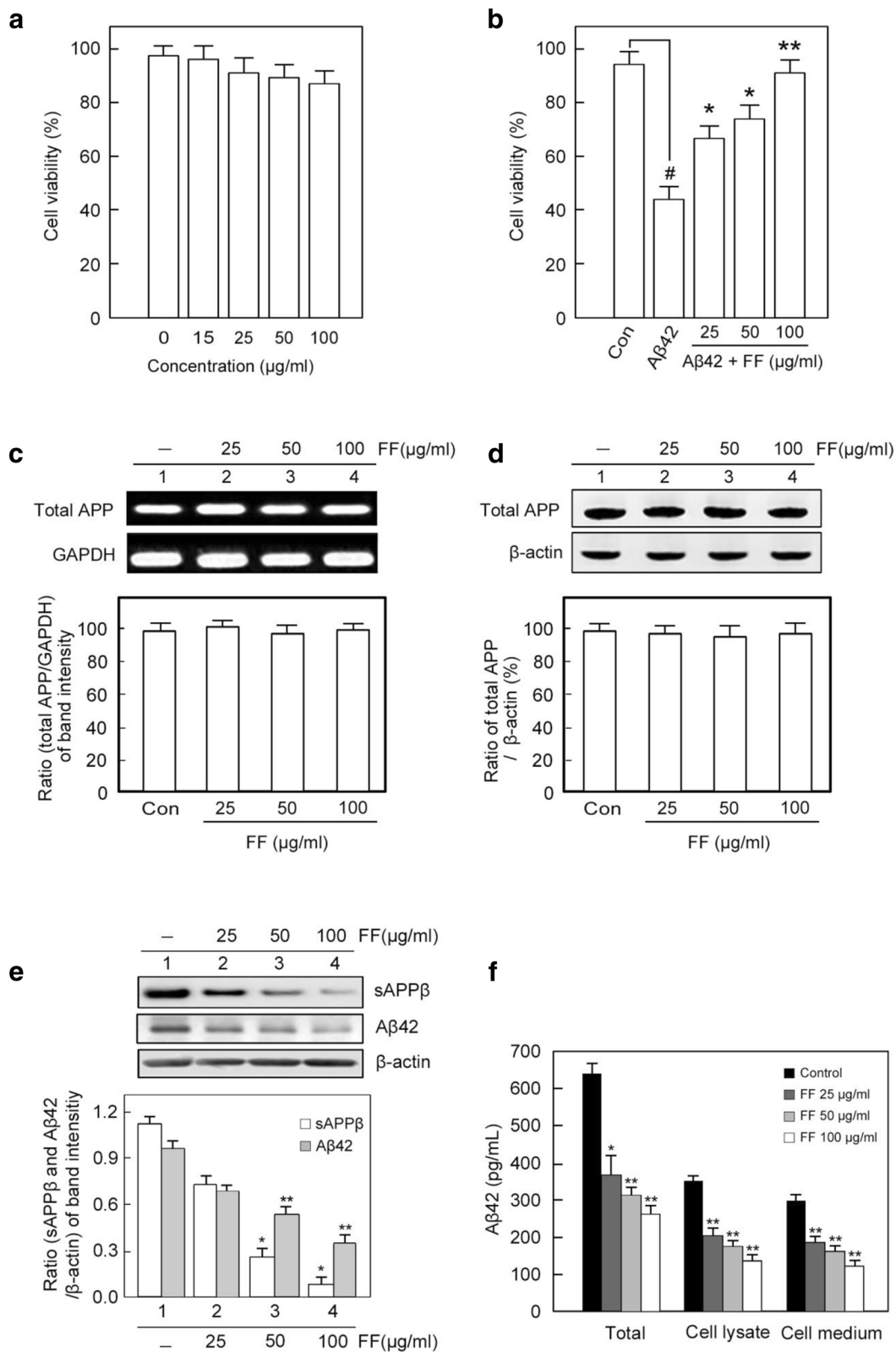


Fig. 4 (See legend on next page.)

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**Fig. 4** Cytotoxicity of **a** SH-SY5Y cells overexpressing APP695swe that were treated with FF (0, 15, 25, 50, and 100 µg/mL) for 24 h. **b** Effects of purified FF on Aβ42-induced cytotoxicity in SH-SY5Y cells. Cells were exposed to various concentrations of FF (25, 50, and 100 µg/mL) for 24 h. After pretreatment, cells were treated with Aβ42 (5 µM) for 24 h, and cell viability was determined by MTT assay. \**P* < 0.05; \*\**P* < 0.01 versus Aβ42-treated cells. #*P* < 0.01 versus Aβ42 non-treated cells. **c** FF treatment did not alter APP expression levels. SH-SY5Y cells overexpressing APP695swe were treated with increasing concentrations of FF (25, 50, and 100 µg/mL) for 24 h. The expression of full-length APP was determined by both RT-PCR and western blotting analysis. The levels of APP mRNA (**c**) and protein (**d**) did not significantly differ between control and FF-treated cells. Results are shown as the mean ± SEM of experiments performed in triplicate (*n* = 3). **e** Expression levels of sAPPβ and Aβ42 in SH-SY5Y-APP695swe cells treated with Leu-Asn peptide measured by western blot analysis. **f** FF reduced Aβ42 levels in both cell culture medium supernatants and cell lysates. SH-SY5Y cells overexpressing APP695swe were treated with increasing concentrations of FF (25, 50, and 100 µg/mL) for 24 h. Aβ42 levels were determined using Aβ40- and Aβ42-specific sandwich ELISAs. Total Aβ42 was the sum of Aβ42 in the supernatant and lysate. Results are expressed in picogram per milliliter, and all experiments were performed in triplicate (*n* = 3). \**P* < 0.05, \*\**P* < 0.01 compared with the control

nor was cell toxicity observed at concentrations of 50 and 100 µg/mL (Fig. 4a). As illustrated in Fig. 4b, Aβ42 (5 µM for 24 h) treatment decreased SH-SY5Y cell viability to  $47.5 \pm 0.5\%$ , while FF (25, 50, and 100 µg/mL) treatment prevented Aβ42-induced damage, restoring cell survival to  $66.9 \pm 1.6\%$ ,  $80.7 \pm 2.5\%$ , and  $98.0 \pm 1.3\%$  (mean ± S.E.M., *n* = 3), respectively.

Diverse pathologies and etiologies are involved in neurodegenerative disorders, including AD. A growing body of data suggests free radical-induced toxicity, oxidative impairment, mitochondrial dysfunction, and inflammatory mechanisms may represent common aspects in the pathogenesis of neurodegeneration (Crouch et al. 2008). Among the brain's cells, microglia have been reported to play important roles in neuropathological conditions. In response to a variety of neurotoxic stimuli, microglia become active and secrete a number of pro-inflammatory cytokines and cytotoxic molecules, including ROS and reactive nitrogen species (RNS) (Durrenberger et al. 2012). Conversely, vitamin E and minocycline have been shown to exhibit neuroprotective roles by inhibiting microglial activation (Ryu et al. 2004). Therefore, in this study, we examined the protective effect of FF, a major constituent of brown algae polyphenolics, against Aβ-induced oxidative cell death in SH-SY5Y neuroblastoma cells. We initially examined Aβ-induced cytotoxicity in SH-SY5Y cells with an MTT reduction assay. Aβ decreased cell viability in SH-SY5Y cells, though this damage was effectively attenuated in the presence of FF. The molecular mechanisms of Aβ-mediated neurotoxicity are not completely understood, but this finding suggested that β-secretase inhibitor, FF rescues cell death induced by Aβ may rely on inhibition β-secretase-mediated downregulation of Aβ.

#### Inhibition of amyloidogenic APP processing in SH-SY5Y-APP695swe cells

Human neuroblastoma SH-SY5Y-APP695swe cells overexpressing APP were treated with 25, 50, and 100 µg/mL FF, and APP expression was determined via RT-PCR and western blot analysis (Fig. 4c and d). As shown in Fig. 4c and d, treatment with 25, 50, and

100 µg/mL FF did not induce significant changes in APP levels compared to those in the control. We then examined whether FF affected Aβ production, we measured soluble-APPβ (sAPPβ) and Aβ42 levels produced in SH-SY5Y-APP695swe cells after administration of FF at different concentrations for 24 h. The western blot results indicated that FF treatment decreased sAPPβ and Aβ42 expression in a dose-dependent manner in SH-SY5Y-APP695swe cells (Fig. 4e).

Both the culture medium supernatants and cell lysates were collected, and Aβ concentrations were determined by ELISA. Total Aβ was the sum of Aβ in the supernatant and lysate. FF significantly decreased Aβ in both the lysate of SH-SY5Y-APP695swe cells and the cell culture supernatant in a dose-dependent manner (Fig. 4f). Aβ production is likely a primary event in the pathogenesis of AD; therefore, understanding the manner in which APP and the secretases are brought together could provide needed data for designing therapeutic strategies for AD. Previous studies have shown that synthesized peptides such as OM99-2 and KMI 420 bind to β-secretase and prevent its access to APP, thereby reducing Aβ generation. However, the molecular mechanisms by which APP and β-secretase come in close proximity to each other in transient or stable complexes are essentially unknown (Ghosh et al. 2012). In this study, we demonstrated that FF strongly reduced β-secretase cleavage of APP and Aβ generation in a transgenic SH-SY5Y cell line.

#### Conclusions

We have attempted purification of a novel β-secretase inhibitor, FF from marine brown algae, *E. bicyclis* extracts for the first time. FF exhibited the strongly inhibitory activity against β-secretase and Aβ production on in-vitro. We have characterized this β-secretase inhibitor to aid the development of a novel anti-AD treatment for the drug or food supplement industries.

#### Abbreviations

AD: Alzheimer's disease; ANOVA: Analysis of variance; APP: Amyloid precursor protein; Aβ: β-Amyloid peptide; BACE1: The β-site APP-cleaving enzyme; BCA: Bicinchoninic acid; DMSO: Dimethyl sulfoxide; DNA: Deoxyribonucleic



acid; DNP: 2,4-Dinitrophenyl; ECL: Enhanced chemiluminescence; ELISA: Enzyme-linked immunosorbent assay; FAB: Fast atom bombardment; FF: Fucufuroeckol-b; FT-IR: Fourier-transform infrared; HPLC: High-performance liquid chromatography; IRES: Internal ribosome entry site; LC: Liquid chromatography; MCA: 7-Methoxycoumarin-4-yl acetyl; MEM: Modified Eagle's medium; MTT: 3-(4,5-Dimethylthiazol-2-yl)-2,5-diphenyl tetrazolium bromide; NMR: Nuclear magnetic resonance; ODS: Octadecylsilane; PVDF: Polyvinylidene fluoride; Q-TOF-MS: Quadrupole time-of-flight mass spectrometry; RFU: Relative fluorescence units; RNS: Reactive nitrogen species; ROS: Reactive oxygen species; RT-PCR: Reverse transcription polymerase chain reaction; sAPP $\beta$ : Soluble-APP $\beta$ ; SDS: Sodium dodecyl sulfate; SEM: Scanning electron microscope; SPE: Solid phase extraction; SPSS: Statistical package for the social sciences; TBST: Tris-buffered saline with Tween20; TLC: Thin layer chromatography; TOF: Time-of-flight; UV: Ultraviolet visible

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#### Availability of data and materials

All datasets generated during and/or analyzed during the current study are available from the corresponding author on reasonable request.

#### Authors' contributions

HGB and LJK conceived and designed the study and helped to draft the manuscript and revised the manuscript. LJK performed the experiments, analyzed the data, and drafted the manuscript. All authors read and approved the final manuscript.

#### Ethics approval and consent to participate

Not applicable.

#### Consent for publication

Not applicable.

#### Competing interests

The authors declare that they have no competing interests.

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