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journal or	Foods
publication title	
volume	9
number	6
year	2020
URL	http://hdl.handle.net/10258/00010265

doi: info:doi/10.3390/foods9060825



Article

Evaluation of Amyloid β₄₂ Aggregation Inhibitory Activity of Commercial Dressings by A Microliter-Scale High-Throughput Screening System Using Quantum-Dot Nanoprobes

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Received: 30 April 2020; Accepted: 22 June 2020; Published: 24 June 2020



Abstract: The aggregation and accumulation of amyloid β (A β) in the brain is a trigger of pathogenesis for Alzheimer's disease. Previously, we developed a microliter-scale high-throughput screening (MSHTS) system for A β_{42} aggregation inhibitors using quantum-dot nanoprobes. The MSHTS system is seldom influenced by contaminants in samples and is able to directly evaluate A β_{42} aggregation inhibitory activity of samples containing various compounds. In this study, to elucidate whether the MSHTS system could be applied to the evaluation of processed foods, we examined A β_{42} aggregation inhibitory activity of salad dressings, including soy sauces. We estimated the 50% effective concentration (EC₅₀) from serial diluted dressings. Interestingly, all 19 commercial dressings tested showed A β_{42} aggregation inhibitory activity. It was suggested that EC₅₀ differed by as much as 100 times between the dressings with the most (0.065 ± 0.020 v/v%) and least (6.737 ± 5.054 v/v%) inhibitory activity. The highest activity sample is traditional Japanese dressing, soy sauce. It is known that soy sauce is roughly classified into a heat-treated variety and a non-heat-treated variety. We demonstrated that non-heat-treated raw soy sauce exhibited higher A β_{42} aggregation inhibitory activity than heat-treated soy sauce. Herein, we propose that MSHTS system can be applied to processed foods.

Keywords: Alzheimer's disease; Amyloid β ; amyloid β aggregation inhibitor; quantum dot; soy sauce

1. Introduction

One of the problems facing an aging society is the increase of patients with dementia. While various diseases are known to cause dementia, Alzheimer's disease (AD) in particular, accounts for the majority of cases [1–3]. Four AD drugs approved in Japan, donepezil, galantamine, rivastigmine, and memantine, only function by delaying the progression of pathological conditions by temporarily enhancing neurotransmission, and are not fundamental therapeutic agents [4]. The amyloid cascade hypothesis notes that AD is caused by the aggregation and accumulation of 38 to 43 residues of the amyloid β (A β) peptide excised from amyloid precursor protein in the brain [5–8]. Recently, Biogen and Eisai reported that a patient's cognitive decline had been blunted in clinical trials using antibodies, aducanumab, that bind specifically to A β aggregates [9]. However, in March of 2019, a phase III clinical trial of aducanumab was halted because of insufficient evidence to support its effect in AD [10]. In October of 2019, both companies announced that they would apply for a new drug application of aducanumab to the U.S. Food and Drug Administration in 2020, as the effect was confirmed in some



patients who received the drug at a high dose. However, these events remind us of the difficulties in developing AD therapeutics. Therefore, attention is now focused on AD prevention and treatment schemes that target the aggregation and accumulation of A β . There is currently a global search for candidate substances that can inhibit A β aggregation. Since the aggregation and accumulation of A β begins several decades before the expression of AD [11], long-term prevention with functional foods may be more effective than treatment with therapeutic medicine.

Rosmarinic acid (RA) is a polyphenol found in abundance in plants of the Lamiaceae such as rosemary, perilla, and lemon balm. RA is a known inhibitor of A β aggregation [12,13]. Its A β aggregation inhibitory activity was examined using AD model mice and its safety was confirmed in human studies using lemon balm extract [14,15]. Among many other polyphenols, curcumin, which is found in turmeric, is also a famous Aβ aggregation inhibitor [12]. Further, it was reported that importance of functional foods on AD. The extract obtained from miso, a traditional fermented dressing in Japan, suppresses Aβ-induced neuronal damage [16]. Hsu et al., reported that nattokinase degraded amyloid fibrils [17]. Thus, the use of functional foods has attracted attention as a possible AD countermeasure. However, it is technically very difficult to evaluate plant extracts and processed foods as these include various impurities. In general, the Thioflavin T (ThT) method has been used to evaluate A β aggregation inhibitory activity of various substances [18]. ThT emits fluorescence when bound to amyloid fibrils. In this method, the level of A β aggregation is measured from the fluorescence intensity of ThT. However, the excitation and emission wavelengths of ThT are 455 and 490 nm, respectively, so they compete with the absorption wavelengths of many natural substances. Therefore, the ThT method is unsuitable to evaluate food samples that contain various contaminants. A method of directly observing A β aggregates with a transmission electron microscope (TEM) is widely used. Because it is necessary to dry the A β aggregates sample when preparing, the observation under physiological conditions is difficult. Further, the amount of aggregates is biased depending on the field of view even in the same sample, suggesting that there is a problem in quantitative. In addition, the ThT and TEM method generally require several steps for sample preparation and observation, and it is difficult to analyze a large amount of the sample at one time. In other words, previous conventional method could not perform accurate and quick high throughput quantitative analysis.

Previously, we succeeded in real-time imaging of the $A\beta_{42}$ aggregation process with a fluorescence microscope using a quantum dot (QD) nanoprobe and developed a microliter-scale high-throughput screening (MSHTS) system for A β_{42} aggregation inhibitors by applying this imaging method [19,20]. The MSHTS system has some advantages: (1) only a small sample volume of 5 μ L is required, (2) high-throughput analysis uses a 1536-well plate, and (3) filter effects due to contaminants in the sample are avoided because the amount of $A\beta_{42}$ aggregates is quantified from standard deviation (SD) value estimated from the variation in fluorescence intensity of each pixel of obtained images and the emission wavelength of QD605 does not overlap with the absorption of almost natural products [20,21]. Thus, the MSHTS system can evaluate the magnitude of inhibitory activity for A β_{42} aggregation as EC_{50} values. Before, we evaluated the A β_{42} aggregation inhibitory activity of 52 spices using this method and demonstrated that the herb-based spices of the Lamiaceae family exhibited high $A\beta_{42}$ aggregation inhibitory activity [20]. Then, we found that the activity of boiling water extracts of 11 seaweeds was higher than that of ethanolic extracts and revealed that $A\beta_{42}$ aggregates morphology was affected with seaweed-derived polysaccharide including in boiling water extracts [22]. Further, we recently developed an automated MSHTS system to evaluate larger numbers of samples at once [21]. Screening 504 plant extracts collected in Hokkaido, Japan, we found that Geraniales and Myrtales within Rosids showed high A β_{42} aggregation inhibitory activity. Thus, MSHTS system is useful for quantitative evaluation of $A\beta_{42}$ aggregation inhibition ability of various natural products. However, it is unclear whether MSHTS system can evaluate $A\beta_{42}$ aggregation inhibitory activity in foods including various natural substances with many impurities. In this study, to elucidate whether the MSHTS system is applied to processed foods such as salad dressings, including soy sauces, we evaluated $A\beta_{42}$ aggregation inhibitory activity of dressings using the MSHTS system. We found that all tested

commercial dressings showed $A\beta_{42}$ aggregation inhibitory activity despite there were differences in their activities. Especially raw soy sauce showed the highest inhibitory activity among the tested samples. These results suggest that the MSHTS system is a powerful and useful tool that is expected to be applied and developed in various processed foods.

2. Materials and Methods

2.1. Materials

Human A β_{42} (4349-v, Peptide Institute Inc., Osaka, Japan) and Cys-conjugated A β_{40} (23519, Anaspec Inc., Fremont, CA, USA) kits were purchased commercially. Twenty different commercially available salad dressings and soy source brands were purchased from Japanese companies (Kewpie, Sameura Foods, Sanyo Coffee Foods, Shiranukacho Shinko Kosha, Shinshu Shizen Okoku, Seijo Ishii, Taiyo Sangyo, Tsukiboshi Foods, Nihon Syoyu Kogyo, Big Chef, Pure Foods Toya, Yamada Bee Farm, Riken Vitamin, H+B Life Science) using catalog shopping in June 2016.

2.2. Preparation of QDAβ Nanoprobe

The QDA β nanoprobe was prepared using QD-PEG-NH₂ (QdotTM 605 ITKTM Amino (PEG) Quantum dot; Q21501MP, Thermo Fisher Scientific, Waltham, MA USA) according to our previous reports [19–22]. The QDA β nanoprobe was prepared by first reacting 10 μ M QD-PEG-NH₂ with 1 mM sulfo-EMCS (22307, Thermo Fisher Scientific, Waltham, MA, USA) in PBS (phosphate-buffered saline) for 1 h at room temperature. QDA β concentration was determined by comparing absorbance at 350 nm to that of unlabeled QD-PEG-NH₂.

2.3. Estimation of EC_{50} by the MSHTS System

The EC₅₀ values of various dressings were determined by a modified MSHTS system, as was described in our previous reports [20–22]. More specifically, various concentrations of each dressing, 30 nM QDA β , and 30 μ M A β_{42} in PBS containing 5% EtOH and 3% DMSO were incubated in a 1536-well plate (782096, Greiner, Kremsmünster, Austria) at 37 °C for 24 h. The QDA β -A β_{42} aggregates that formed in each well were observed by an inverted fluorescence microscope (TE2000, Nikon, Tokyo, Japan). Standard deviation (SD) values of fluorescence intensities of 40,000 pixels (200 × 200 pixels) around the central region of each well were measured by ImageJ software Ver 1.53b (NIH). The SD values, which were approximately proportional to the amount of aggregates [20–22], were plotted against the concentrations of added salad dressings to establish an inhibition curve.

2.4. Fluorescence Microscopy

Aggregates in the 1536-well plate were observed by an inverted fluorescence microscope (TE2000-S, Nikon) using a 4× objective lens (Plan Fluor 4×/0.13 PhL DL, Nikon) equipped with a color CCD camera (DP72, Olympus, Tokyo, Japan).

2.5. Transmission Electron Microscopy

Samples were deposited in 10 μ L aliquots onto 200-mesh copper grids and negatively stained with 1% phosphotungstic acid at room temperature. Specimens were examined under an H-7600 TEM (Hitachi, Tokyo, Japan) at 60 kV.

2.6. ThT Assay

The ThT assay was conducted according to the method of Levine modified in our laboratory [18,21]. Statistical analyses between +ThT and –ThT samples were performed with EZR (Saitama Medical Center, Jichi Medical University, Saitama, Japan), which is a graphical user interface for R (The R Foundation for Statistical Computing) [23]. More precisely, it is a modified version of R commander designed to add statistical functions frequently used in biostatistics.

2.7. SDS-Polyacrylamide Gel Electrophoresis

SDS-polyacrylamide gel electrophoresis (SDS-PAGE) was performed using standard techniques. Soy sauce and raw soy sauce were heated by block incubator at 80 °C for 60 min before electrophorese. For dialysis protocol, soy sauce and raw soy sauce were dialyzed against distilled water. Distilled were changed three times for overnight. Then, the gel was silver-stained by staining kit (2D-SLVER STAIN II, COSMO BIO)

3. Results and Discussion

3.1. Evaluation of $A\beta_{42}$ Aggregation Inhibitory Activity of Commercial Dressings by ThT Method

First, in order to assess whether 19 liquid salad dressings could be evaluated for activity by the ThT method, their absorbance spectra were measured using a Nanodrop 2000c (Thermo Fisher Scientific) (Figure 1). The excitation and emission wavelengths of ThT are 450 nm and 490 nm, respectively. As shown in Figure 1, only 3 of the 19 dressings (samples L, N, M) showed no absorbance of 1 or more at each wavelength. Most of the samples contained soy sauce as a raw material, and the color was black or brown depending on the content of soy sauce. In the 15 samples that showed absorption peaks at the ThT excitation and emission wavelengths, the absorption peak shifted to the right in proportion as the color became darker. This indicates that the evaluation of $A\beta_{42}$ aggregation inhibitory activity of the liquid dressings using the ThT method was difficult because the absorption wavelength of almost samples overlapped with the excitation and emission wavelengths of ThT.



Figure 1. Absorbance of 19 commercial liquid dressings. Absorbance of the 19 commercial liquid dressings shown in Table 1 was measured. As for the absorption wavelength of dressings, three samples (L, M, N) do not show an overlap with the excitation (450 nm) and emission (490 nm) wavelengths of ThT. Two samples (H, G) overlap with the ThT excitation wavelength. The remaining 14 samples had an overlap with both excitation and emission wavelengths.

To confirm whether the evaluation using ThT method was performed correctly, the fluorescence intensity of five samples at a high concentration (40 v/v%) was measured in three conditions; +A β_{42} and +ThT, -A β_{42} and +ThT, -A β_{42} and -ThT (Figure 2). Soy sauce (sample A), soy sauce containing perilla (sample B), Japanese style dressing (sample C), oil dressing (sample D), and Chinese dressing (sample E) were selected and evaluated. Soy sauce (sample A) was used as the control for the other four samples. At 40 v/v% sample concentration, the fluorescence intensity of the -A β_{42} solution, -A β_{42} and -ThT solution were not significantly different from that of the solution containing A β_{42} and ThT. We confirmed that samples A and B showed higher fluorescence intensity than only A β_{42}

and ThT sample in all conditions (negative control, black line). The fluorescence intensity of samples C, D, and E did not show a higher value than the negative control. The color and some components of the evaluated samples may affect the ThT method by absorbing the excitation or emission of ThT. If the sample solution without $A\beta_{42}$ exhibits a higher fluorescence intensity than the negative control, it is difficult to determine whether the sample solution affects ThT. Therefore, the ThT method might not accurately evaluate the $A\beta_{42}$ aggregation inhibitory activity of a commercial dressing.



Figure 2. Effect of sample solution on ThT fluorescence intensity. $A\beta_{42}$ (+), ThT (+): $A\beta_{42}$ solution and ThT solution are mixed with sample; $A\beta_{42}$ (-), ThT (+): only ThT solution is mixed with sample solution; $A\beta_{42}$ (-), ThT (-): neither $A\beta_{42}$ solution nor ThT solution are mixed and only the sample solution is used. As a negative control, the sample used was an assay buffer (10% EtOH, 1 × PBS) under the conditions of $A\beta_{42}$ (+) and ThT (+) (black line), and its average value of absorbance was 5.28 (A.U.). There is no significant difference between +ThT (+/– $A\beta_{42}$) and ThT (–) condition in all samples (One way ANOVA, p > 0.05).

3.2. Evaluation of $A\beta_{42}$ Aggregation Inhibitory Activity of Commercial Dressings Using MSHTS Sysytem

Previously, we reported the real-time imaging of the $A\beta_{42}$ aggregation process with a fluorescence microscope using QD nanoprobes (Figure 3A) and developed MSHTS system (Figure 3B) for $A\beta_{42}$ aggregation inhibitors by applying this imaging method [19,20]. In the MSHTS system, QD-labeled Aß co-aggregated with intact $A\beta_{42}$, so that amyloid aggregates were observed by fluorescence microscopy. Since the emission wavelength of QD605 does not overlap with the absorption of almost dressings, it is less susceptible to substances that exhibit an inner filter effect. These aggregates (complex of $A\beta_{42}$ and QD A β) caused an inhomogeneous distribution of fluorescence intensity in images, resulting in an increased standard deviation (SD) value estimated from the variation in fluorescence intensity of each pixel in the images. Therefore, we could estimate the effects of certain aggregation inhibitors by detecting changes in the SD value. Here, we evaluated the $A\beta_{42}$ aggregation inhibitory activity of 19 commercial dressings using the MSHTS system (Figure 3C), and estimated the EC_{50} values from the SD value of each image (Table 1). In Table 1, EC_{50} values are sorted in ascending order of aggregation inhibitory activity, then each dressing was assigned a letter from A to S. As shown in Figure 3C, all commercial dressings almost completely inhibited A β_{42} aggregation at a concentration of 40 v/v%. Samples A to J completely inhibited aggregation even at 4 v/v%, whereas samples K to S formed small aggregates. At a concentration of 0.4 v/v%, a slight change in the shape of aggregates was observed in samples A to E. The EC₅₀ value of the sample with the highest activity was 0.065 v/v° , and that with the lowest activity was 6.737 v/v%. There was an about 100-fold difference in activity between these 2 samples. Among the 19 dressings, only three samples (D, K, and N) contained plant oil while the other 16 samples were non-oil type dressings. We compared the mean value of EC_{50} value of non-oil and oil type (non-oil type: $1.330 \pm 1.692 v/v\%$, oil type: $1.019 \pm 0.946 v/v\%$) and

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performed a statistical analysis. There was no significant difference between non-oil type and oil type (Student's *t*-test, p > 0.05). The activity did not depend on the presence of oil, suggesting that oil in the dressing did not affect the A β_{42} aggregation inhibitory activity. Next, to confirm whether A β_{42} aggregation was inhibited by the effect of the dressing, we observed the A β_{42} aggregates at a sample concentration of 0.04 v/v% using TEM (Figure 4). Sample A, which had high A β_{42} aggregation inhibitory activity, showed a significant decrease in A β_{42} aggregates compared to samples J and S. Sample J also significantly decreased aggregates than sample S. These results were consistent with the A β_{42} aggregation inhibitory activity calculated by the MSHTS system. Sample A, which showed the highest A β_{42} aggregation inhibitory activity, was soy sauce, a traditional Japanese liquid dressing. The remaining 17 samples, except for sample D, contained soy sauce. In other words, soy sauce evidently exhibited high A β_{42} aggregation inhibitory activity. Natto, a traditional Japanese food made of fermented soybeans, has antibacterial, as well as a soybean peptide with a neuroprotective effect [24]. It is possible that soybean-derived proteins found in soy sauce, and/or various metabolic products caused by soybean fermentation, may have a positive effect on AD.



Figure 3. Evaluation of A β_{42} aggregation inhibitory activity using the MSHTS system. (**A**) Real-time imaging of A β_{42} aggregation using a quantum-dot nanoprobe using fluorescence microscopy. A β_{42} and QDA β were mixed and incubated for 24 h at 37 °C. Co-aggregates of A β_{42} and QDA β formed. (**B**) A scheme of the MSHTS system of A β_{42} aggregation inhibitors. (**C**) Fluorescence microscope image of concentration-dependent inhibition of A β_{42} aggregation of dressings observed by the MSHTS system. At a sample concentration of 4 v/v% or more, the brightness of the image was uniform, indicating that no A β_{42} aggregates formed. At a sample concentration of 0.04 v/v% or less, A β_{42} aggregates were observed in all samples. All images were captured using a conventional fluorescence microscope.



Figure 4. Electron microscopy images of A β_{42} aggregates. The A β_{42} solution was mixed with each sample (0.04 v/v%) and was incubated for 24 h at 37 °C. The images of A β_{42} aggregates were captured by TEM at 3000× magnification. Bars: 500 nm.

EC ₅₀ (v/v%)	Oil Type
$0.122 \pm 0.034 \; (w/v\%)$	-
0.065 ± 0.020	Non-oil
0.094 ± 0.017	Non-oil
0.133 ± 0.021	Non-oil
0.227 ± 0.026	Oil
0.230 ± 0.026	Non-oil
0.334 ± 0.075	Non-oil
0.395 ± 0.130	Non-oil
0.413 ± 0.084	Non-oil
0.480 ± 0.101	Non-oil
0.508 ± 0.025	Non-oil
0.763 ± 0.607	Oil
1.350 ± 0.247	Non-oil
1.360 ± 0.590	Non-oil
2.067 ± 0.728	Oil
2.132 ± 1.473	Non-oil
2.150 ± 0.887	Non-oil
2.313 ± 0.490	Non-oil
2.580 ± 0.173	Non-oil
6.737 ± 5.054	Non-oil
	$\begin{array}{c} {\rm EC_{50}} \ (v/v\%) \\ \hline 0.122 \pm 0.034 \ (w/v\%) \\ 0.065 \pm 0.020 \\ 0.094 \pm 0.017 \\ 0.133 \pm 0.021 \\ 0.227 \pm 0.026 \\ 0.230 \pm 0.026 \\ 0.334 \pm 0.075 \\ 0.395 \pm 0.130 \\ 0.413 \pm 0.084 \\ 0.480 \pm 0.101 \\ 0.508 \pm 0.025 \\ 0.763 \pm 0.607 \\ 1.350 \pm 0.247 \\ 1.360 \pm 0.590 \\ 2.067 \pm 0.728 \\ 2.132 \pm 1.473 \\ 2.150 \pm 0.887 \\ 2.313 \pm 0.490 \\ 2.580 \pm 0.173 \\ 6.737 \pm 5.054 \\ \end{array}$

Table 1. EC₅₀ values of 19 commercial dressing samples using MSHTS system.

3.3. Effect of Salt Concentration on $A\beta_{42}$ Aggregation

In general, it is well known that the soy sauce contains a large amount of NaCl and that high salt concentration affects protein aggregation. In order to determine whether the A β_{42} aggregation inhibitory activity of soy sauce was due to NaCl, we examined the effect of NaCl concentration on the shape of $A\beta_{42}$ aggregates (Figure 5A). We prepared a 4000 mM NaCl solution. Then, the solution was gradually diluted to six concentrations with five-fold dilutions. These 7 concentrations of NaCl solution was mixed with the 50 μ M A β_{42} solution and observed using the MSHTS system. The formation of A β_{42} aggregates was slightly affected by 200 mM NaCl solution. At 1000 mM of NaCl, AB42 aggregates were fragmented, and at 2000 mM, abnormal aggregates such as large clumps were observed. The SD values obtained from images were gradually decreased from 2000 mM to 200 mM (Figure 5B). However, 0.32-40 mM NaCl did not affect the SD value, suggesting that $A\beta_{42}$ aggregates were formed. The NaCl concentration in sample A soy sauce is 16.2% (2.77 M). Because the EC₅₀ of sample A is $0.065 \pm 0.020 v/v\%$, it was indicated that 1.8 mM NaCl is included. The NaCl concentration of the 0.4 v/v% and 0.04 v/v% solution samples is 11 mM and 1.1 mM, respectively. Therefore, the NaCl in 0.065 v/v° (EC₅₀ value) soy sauce solution might not affect the formation of A β_{42} aggregates. In fact, as shown in Figure 3C, aggregation was inhibited in samples A to J when NaCl was included at 0.4 v/v%. Especially, 0.04 v/v% of sample A inhibited the aggregation formation. These results suggest that the A β_{42} aggregation inhibitory activity by soy sauce was due to not NaCl but other components.



Figure 5. Influence of salt concentration on $A\beta_{42}$ aggregation. (**A**) NaCl solution adjusted to each concentration and the $A\beta_{42}$ solution were mixed then incubated for 24 h at 37 °C. Using the MSHTS system, the influence of NaCl concentration on the formation of $A\beta_{42}$ aggregates was examined. At 2000 mM NaCl, $A\beta_{42}$ and QDA β were salted out. From their morphology, it is believed that these solids were not $A\beta_{42}$ aggregates. At 1000 mM, aggregates started to form. At 40 mM or less, no significant effect was observed on the aggregates. From 40 to 0.32 mM, normal aggregates were formed. Bars: 100 μ m. (**B**) Ratio of SD value at each NaCl concentration. At 40 mM or less, the SD value was not affected by NaCl concentration.

3.4. Influence of Heating and Dialysis Treatment on $A\beta_{42}$ Aggregation Inhibitory Activity

As shown in Figure 3, among the 19 dressings, soy sauce showed the highest activity when the MSHTS system was used. In fact, there are several types of soy sauce, which can be classified according to the sterilization method, the composition of the raw materials, the color, and salt concentration of the product. Among them, we focused on raw soy sauce that was not heat-sterilized. Since raw soy sauce is sterilized by filtration, it has a feature that compounds produced during the fermentation process and enzymes derived from microorganisms are not inactivated. Here, we evaluated A β_{42} aggregation inhibitory activity of raw soy sauce purchased from NIHON SYOYU KOGYO using the MSHTS system. The A β_{42} aggregation inhibitory activity of raw soy sauce was 0.0045 ± 0.0015 v/v% (Data not shown). This activity is about 15 times higher than that of soy sauce which showed the highest activity among the 19 dressings. Therefore, it is likely that the difference in the aggregation inhibitory activity of A β_{42} between soy sauce and raw soy sauce is caused by the raw material-derived protein and the microorganism-derived enzyme, which are lost by heating.

To examine whether proteins and/or low molecular weight compounds in soy sauce and raw soy sauce affect $A\beta_{42}$ aggregation inhibitory activity, each sample was subjected to SDS-PAGE after heat treatment (80 °C, 60 min) and/or dialysis treatment and their band patterns were compared using silver staining (Figure 6A). As revealed by an SDS-PAGE gel (lanes 1 and 5), the property and amount of proteins in soy sauce and raw soy sauce differed. Bands were detected at 15 and 25 kDa in both samples while 30 and 100–200 kDa bands were detected only in raw soy sauce. Heat treatment did not change the banding pattern of soy sauce, but reduced the intensity of the100–200 kDa band of raw soy sauce (Figure 6A, from lanes 2 and 6). Dialysis treatment reduced the 10–15 and 25 kDa bands in both samples (Figure 6A, lanes 3 and 7). Heat treatment after dialysis reduced the band at about 25 kDa in

А

Soy sauce

soy sauce less than the dialyzed sample, and reduced the bands at about 15, 25 and 100–200 kDa in raw soy sauce (Figure 6A, lanes 4 and 8).

1.76

0.962

6.28

3.71

100

10

1

С

Raw soy sauce

G Control o Heat -1 Dialys EC50 (v/v%) 8 MW 0.1 0.0412 kDa 0.0210 200 100 0.01 0.001 * Dialysis Dialysis Control × Oiai Heat Soy sauce Raw soy sauce В Concentration of raw soy sauce (v/v%) Concentration of soy sauce (v/v%) 0.001 0.01 0.1 10 0.001 0.01 0.1 1 10 1 control Heated Dialyzed Heated Dialyzed

Figure 6. Influence of heat and dialysis treatment on $A\beta_{42}$ aggregation inhibitory activity of soy sauce and raw soy sauce. (A) SDS-PAGE analysis demonstrated that heating and dialysis treatments changed the band pattern of soy sauce and raw soy sauce. Lanes 1–4: soy sauce; lanes 5–8: raw soy sauce. Lanes 1 and 5: control; lanes 2 and 6: heat treatment; lanes 3 and 7: dialysis treatment; lanes 4 and 8: dialysis and heat treatment. MW: molecular weight marker. (B) Fluorescence images of $A\beta_{42}$ aggregates in each condition. Effect of heating and dialysis treatment on $A\beta_{42}$ aggregation inhibition of soy sauce and raw soy sauce. All images were captured using a conventional fluorescence microscope. Bars: 100 μ m. (C) A β_{42} aggregation inhibitory activity (EC₅₀) of soy sauce and raw soy sauce after heating and dialysis calculated by the MSHTS system. Whereas the activity of soy sauce was not changed by heating, the EC_{50} of heated raw soy sauce was about 30 times higher than that of the control sample, suggesting that the activity was greatly reduced. In both soy sauce and raw soy sauce, EC_{50} was approximately double after dialysis.

Next, using the MSHTS system, we assessed whether heating and dialysis treatment would affect the A β_{42} aggregation inhibitory activity of soy sauce and raw soy sauce (Figure 6B). Furthermore, the dialyzed soy sauce and raw soy sauce were also heated and analyzed. As shown in Figure 6C, the A β_{42} aggregation inhibitory activity of soy sauce was not changed by heating. This result was consistent with the SDS-PAGE banding pattern (Figure 6A, lanes 1 and 2). In addition, the activity of soy sauce decreased even after dialysis treatment, even more so when heat treatment followed dialysis. The A β_{42} aggregation inhibitory activity of raw soy sauce was reduced by about 30 times or by about half after heat treatment and dialysis, respectively. Dialysis treatment did not affect the 100-200 kDa band that was reduced by heat treatment, i.e., the 100-200 kDa protein did not contribute to the $A\beta_{42}$ aggregation inhibitory activity of raw soy sauce. The heat treatment after dialysis decreased

9 29

0.630

A β_{42} aggregation inhibitory activity by about 440 times, which was consistent with the SDS-PAGE result in which many bands were reduced (Figure 6A, lane 8). These results suggest that the A β_{42} aggregation inhibitor found in soy sauce and raw soy sauce is a low molecular weight compound that is removed by dialysis and a protein that is thermally denatured by heat treatment at 80 °C for 60 min. Since the A β_{42} aggregation inhibitory activity of raw soy sauce was greatly reduced after heat treatment, the proteins found in raw soy sauce are considered to be particularly important for the inhibition of A β_{42} aggregation.

In general, proteins and peptides are denatured by heating, then lose their physiological activity. It is possible that the presence or absence of heat treatment in the fermentation process may be involved in the physiological activity of dressings such as soy sauce, which contains A β_{42} aggregation inhibitory activity. The main raw materials of soy sauce, soy, and wheat, are decomposed into amino acids, peptides and saccharides in the manufacturing process. It is known that soybeans have many physiologically active ingredients such as soy protein and isoflavone [25–27], so these physiologically active ingredients are also present in soy sauce. Soy is widely applied to fermented foods such as miso and natto, Japanese traditional foods. Actually, physiological activity has also been reported for these foods. Miso extract suppresses $A\beta$ -induced neuronal damage [16]. Genistein, one of isoflavone, mitigated A β deposition and neuroinflammation in mice [28]. It was reported that natto peptide exhibited antimicrobials effects and that nattokinase has amino residues playing a intramolecular chaperone [24,29] Further, vitamin K2 (menaquinone-7), which is abundant in natto, is an important factor in the synthesis of sphingolipids present in brain cell membranes that support cell signaling function and structure formation [30–32]. We are currently investigating the active compound in raw soy sauce using this analytical method. Soy sauce is used in many traditional dishes in Japan, and its effectiveness against AD would be significant for the prevention of this disease.

4. Conclusions

In this work, we evaluated A β_{42} aggregation inhibitory activity of 19 commercial liquid dressings using the MSHTS system. All tested dressings exhibited $A\beta_{42}$ aggregation inhibitory activity, suggesting that the MSHTS system can be applied to processed food containing various impurities. Japanese traditional liquid dressings, soy sauce, exhibited the highest inhibitory activity. However, these findings are limited to in vitro conditions. The physiological activity of the dressings should be clarified through animal experiments, taking into account dynamics such as intestinal absorption and metabolism, particularly the permeability of the blood-brain barrier. Although there is a strong demand for functional food products that help maintain and improve brain function, it is not realistic to subject all food products to animal testing. We are confident that we can progress quickly to a second screening stage such as animal testing by using the MSHTS system as a first screening tool to discover food materials with high A β_{42} aggregation inhibitory activity. Recently, we confirmed that aggregation of various amyloid proteins A β_{42} , tau, and α -synuclein could be visualized using nonlabelled QD and succeeded in the evaluation of aggregation inhibitory activity of RA [33]. MSHTS system using nonspecific binding of QD to amyloid proteins might bring speed and simplification of the screening of various foods using various amyloids. Furthermore, we expected that the combination of our previously developed automated-MSHTS system [21] and non-specific MSHTS system allows enormous, comprehensive screening of foods, thereby creating the potential for new approaches to overcome AD.

Author Contributions: Conceptualization: K.T.; Methodology: K.T.; Software: M.K., W.Y., and K.S.; Formal analysis: M.K., W.Y., and K.S.; Data curation: M.K., W.Y., and K.S.; Writing—original draft preparation: M.K.; Writing—review and editing: X.L., K.S., K.U., and K.T.; Supervision: K.T. All authors have read and agree to the published version of the manuscript.

Funding: This work was supported by JSPS KAKENHI Grant Number JP16H03288 to K.T. and the Matching Planner Program from Japan Science and Technology Agency (MP28116808300 to K.T.).

Acknowledgments: Part of this work was conducted at the Chitose Institute of Science and Technology, and supported by the Nanotechnology Platform Program (Synthesis of Molecules and Materials) of the Ministry of Education, Culture, Sports, Science and Technology (MEXT), Japan.

Conflicts of Interest: Authors have no conflicts of interest.

Abbreviations

Αβ	amyloid β
MSHTS system	microliter-scale high-throughput screening system
PBS	phosphate buffered saline
QD	quantum dot
RA	rosmarinic acid
SD value	standard deviation value
SDS-PAGE	SDS-polyacrylamide gel electrophoresis
TEM	transmission electron microscope
ThT	thioflavin T

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