Quantitative Measurement of Cerebrospinal Fluid Amyloid-β Species by Mass Spectrometry

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Abstract.

Background: High sensitivity liquid chromatography mass spectrometry (LC-MS/MS) was recently introduced to measure amyloid- β (A β) species, allowing for a simultaneous assay that is superior to ELISA, which requires more assay steps with multiple antibodies.

Objective: We validated the $A\beta_{1-38}$, $A\beta_{1-40}$, $A\beta_{1-42}$, and $A\beta_{1-43}$ assay by LC-MS/MS and compared it with ELISA using cerebrospinal fluid (CSF) samples to investigate its feasibility for clinical application.

Methods: CSF samples from 120 subjects [8 Alzheimer's disease (AD) with dementia (ADD), 2 mild cognitive dementia due to Alzheimer's disease (ADMCI), 14 cognitively unimpaired (CU), and 96 neurological disease subjects] were analyzed. Aβ species were separated using the Shimadzu Nexera X2 system and quantitated using a Qtrap 5500 LC-MS/MS system. $A\beta_{1-40}$ and $A\beta_{1-42}$ levels were validated using ELISA.

Results: CSF levels in CU were $666 \pm 249 \text{ pmol/L}$ in $A\beta_{1-38}$, $2199 \pm 725 \text{ pmol/L}$ in $A\beta_{1-40}$, $153.7 \pm 79.7 \text{ pmol/L}$ in $A\beta_{1-42}$, and $9.78 \pm 4.58 \text{ pmol/L}$ in $A\beta_{1-43}$. The ratio of the amounts of $A\beta_{1-38}$, $A\beta_{1-40}$, $A\beta_{1-42}$, and $A\beta_{1-43}$ was approximately 68:225:16:1. Linear regression analyses showed correlations among the respective $A\beta$ species. Both $A\beta_{1-40}$ and $A\beta_{1-42}$ values were strongly correlated with ELISA measurements. No significant differences were observed in $A\beta_{1-38}$ or $A\beta_{1-40}$ levels between AD and CU. $A\beta_{1-42}$ and $A\beta_{1-43}$ levels were significantly lower, whereas the $A\beta_{1-38/1-42}$, $A\beta_{1-38/1-43}$, and $A\beta_{1-40/}A\beta_{1-43}$ ratios were significantly higher in AD than in CU. The basic assay profiles of the respective $A\beta$ species were adequate for clinical usage.

Conclusion: A quantitative LC-MS/MS assay of CSF $A\beta$ species is as reliable as specific ELISA for clinical evaluation of CSF biomarkers for AD.

Keywords: $A\beta_{1-38}$, $A\beta_{1-40}$, $A\beta_{1-42}$, $A\beta_{1-43}$, cerebrospinal fluid, ELISA, liquid chromatography mass spectrometry

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INTRODUCTION

Cerebrospinal fluid (CSF) amyloid- β (A β)₁₋₄₂ is a definite biomarker of Alzheimer's disease (AD). A decrease in the A β ₁₋₄₂ level, which begins 25 years earlier, allows the prediction of the onset and

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progression of A β amyloidosis in the AD brain. CSF A β_{1-42} is the earliest and most reliable biomarker for the prediction and diagnosis of AD [1–3]. Low CSF A β_{1-42} and cortical amyloid PET ligand binding have been adopted as biomarkers of A β plaques (labeled "A") in the ATN classification system of the NIA-AA research framework and have been proposed as diagnostic criteria based on the biological definition of AD for observational and interventional research [4].

γ-Secretase cleaves the β-carboxyl terminal fragment (β -CTF) of the amyloid- β protein precursor (A β PP) at the ε -site into long A β s (A β ₁₋₄₉ and $A\beta_{1-48}$), which are ultimately converted from $A\beta_{1-43}$ to $A\beta_{1-40}$ and from $A\beta_{1-42}$ to $A\beta_{1-38}$, respectively, by the successive cleavage of tripeptides and tetrapeptides [5, 6]. The half-life of $A\beta$ in the brain is approximately 9 hours. However, the faster turnover of A β_{1-42} than A β_{1-38} and A β_{1-40} in CSF has been reported with aging and AD [7, 8]. γ -Secretase activity in AD is enhanced by the conversion of $A\beta_{1-43}$ to $A\beta_{1-40}$ and $A\beta_{1-42}$ to $A\beta_{1-38}$, leading to low CSF levels of A β_{1-42} and A β_{1-43} [9]. A β_{1-38} and A β_{1-40} have been shown to attenuate $A\beta_{1-42}$ toxicity [10]. $A\beta_{1-43}$ is known to be enriched in the AD brain [11-13] and promotes the rates and extent of protofibril aggregation [14]. CSF A β_{1-43} is of similar diagnostic accuracy to $A\beta_{1-42}$ [15, 16]. The quantification of $A\beta_{1-38}$, $A\beta_{1-40}$, and $A\beta_{1-42}$ in CSF using mass spectrometry and ELISA has already revealed that the $A\beta_{1-42/1-40}$ and $A\beta_{1-42/1-38}$ ratios are superior to CSF A_{β1-42} alone for the detection of amyloid deposition in AD and the differentiation of AD dementia (ADD) from non-AD dementias [17-19]. However, $A\beta_{1-40}$ and $A\beta_{1-38}$ have lower discriminative powers [20-23]. Collectively, these findings imply that the precise quantitation of these AB species, including A β_{1-38} , A β_{1-40} , A β_{1-42} , and A β_{1-43} , is relevant to biomarkers for observational and interventional research.

However, antibody-based immunoassays, such as ELISA, have several disadvantages [24, 25]. Currently available ELISA systems using antibodies against different epitopes or the conformation of A β species measure the levels of various forms of A β species. The specific characteristics of A β with self-aggregation, binding to tubes, post-translational modifications, and cross-reactions with endogenous components of CSF decrease assay sensitivity and lead to variable results depending on the ELISA systems. Therefore, discrepant findings have been reported across each assay procedure and laboratory. This variability represents a major issue for global clinical observational and interventional studies. Commercial ELISA kits and systems are expensive, time-consuming (often include overnight incubations), and indicate the variation of measured values due to the different lots of the ELISA kits.

An approach based on liquid chromatography mass spectrometry (LC-MS/MS) has been developed to overcome these methodological issues [26–29]. The direct confirmation of A β_{1-38} , A β_{1-40} , A β_{1-42} , and $A\beta_{1-43}$ is possible, and detailed comparisons can be made with advanced LC-MS/MS equipment [8, 17–19]. However, the lower sensitivity and intraassay variability of this system compared to immunological methods have yet to be reliably improved [26-29]. Therefore, we herein developed and validated a novel LC-MS/MS method to assay CSF A β_{1-38} , A β_{1-40} , A β_{1-42} , and A β_{1-43} , and further confirm their contribution as biomarkers for AD pathological changes by comparing the results obtained by the conventional ELISA system for $A\beta_{1-40}$ and $A\beta_{1-42}$.

MATERIALS AND METHODS

Subjects

One hundred and twenty CSF samples from 106 patients with neurological diseases and 14 cognitively unimpaired subjects (CU) were examined. The AD group consisted of 4 cases of autosomal inherited ADD with PSEN-1 mutations, 4 of ADD, and 2 of AD due to mild cognitive impairment (ADMCI) based on the criteria of NIA-AA [4, 30, 31]. They were all defined by positive A+T+N+biomarkerchanges. Clinically diagnosed neurological diseases by the respective recent criteria included 17 cases of amyotrophic lateral sclerosis [32], 9 of corticobasal degeneration [33], 6 of encephalopathy, 3 of frontotemporal dementia [34, 35], 14 of multiple sclerosis and related disorders, 6 of multiple system atrophy [36], 11 of Parkinson's disease [37], 4 of peripheral neuropathy, 11 of progressive supranuclear palsy [38], and 15 of spinocerebellar degeneration. Basic profiles are summarized in Table 1 and no significant differences were observed in age among the disease groups. CSF samples were taken in polypropylene vials by lumber puncture in the morning. Cell numbers and protein levels were examined in the first 1 mL of CSF. Each 1 mL of remaining CSF sample was transferred into polypropylene vials after centrifugation at $1,500 \times g$ for 10 min and stored at -80°C until analysis.

Table 1

Subject profiles. Subjects consisted of 10 cases of Alzheimer's disease with dementia (ADD) and AD due to MCI (ADMCI), 17 of amyotrophic lateral sclerosis (ALS), and 9 of corticobasal degeneration (CBD). Six cases of encephalopathy (ENC) consisted of meningoencephalitis, Creutzfeldt-Jacob disease, leukoencephalopathy, superficial siderosis, or cerebral angiitis. Three cases of frontotemporal dementia (FTD), 14 cognitively unimpaired healthy controls (CU), 14 cases of multiple sclerosis (MS) and related disorders including neuromyelitis optica in the multiple sclerosis group, 6 of multiple system atrophy (MSA), 11 of Parkinson's disease (PD), 4 of peripheral neuropathy (PN), 11 of progressive supranuclear palsy (PSP), and 15 of spinocrebellar degeneration (SCD) were analyzed. No significant differences were observed in age among the CU and other groups

| Disease | | Numb | er | Mean age | Age range | |
|-----------|-------|-------|---------|----------|-----------|--|
| | Total | Males | Females | | | |
| ADD/ADMCI | 10 | 5 | 5 | 59 | 41-80 | |
| ALS | 17 | 13 | 4 | 67 | 53-83 | |
| CBD | 9 | 4 | 5 | 72 | 67-78 | |
| ENC | 6 | 1 | 5 | 60 | 41-78 | |
| FTD | 3 | 2 | 1 | 73 | 70-75 | |
| CU | 14 | 5 | 9 | 61 | 22-80 | |
| MS | 14 | 3 | 11 | 40 | 19-75 | |
| MSA | 6 | 2 | 4 | 66 | 58-81 | |
| PD | 11 | 6 | 5 | 65 | 49-79 | |
| PN | 4 | 2 | 2 | 51 | 30-62 | |
| PSP | 11 | 6 | 5 | 74 | 67-82 | |
| SCD | 15 | 7 | 8 | 64 | 39-80 | |
| Total | 120 | 56 | 64 | 62 | 19–83 | |

LC-MS/MS chemicals and reagents

Synthetic human $A\beta_{1-38}$, $A\beta_{1-40}$, $A\beta_{1-42}$, and $A\beta_{1-43}$ were purchased from the Peptide Institute (Osaka, Japan). Uniformly stable isotope-labeled peptides manufactured by rPeptide (Watkinsville, GA, USA), corresponding to each peptide ($A\beta_{1-38}$, $A\beta_{1-40}$, $A\beta_{1-42}$, and $A\beta_{1-43}$), were used as internal standards (ISs). Artificial CSF was purchased from Tocris Bioscience (Bristol, UK), and rat plasma was purchased from Charles River Laboratories Japan (Yokohama, Japan). Other materials and reagents were purchased from Kanto Chemical (Tokyo, Japan), Kokusan Chemical (Tokyo, Japan), and Sigma-Aldrich (St. Louis, MO, USA). Ultrapure water was prepared using Milli-Q Integral 10 (Merck, Darmstadt, Germany).

LC-MS/MS sample preparation by solid-phase extraction

Human CSF samples $(300 \,\mu\text{L})$ and IS solution were mixed in 1500 μL of water/propionic acid (PA)/ trifluoroacetic acid (TFA) (950:50:1) and loaded onto an Oasis HLB 96-well plate (Waters Corporation, MA, USA). After loading, the plate was washed with 1000 μ L of water/Methanol (MeOH)/PA/TFA (900:50:50:1) and then analytes and ISs were eluted with 100 μ L of EtOH/2,2,2-trifluoroethanol (TF-EtOH)/PA/Ethylene glycol (1:1:1:0.1) twice. Eluates from individual samples were evaporated to dryness at approximately 40°C under nitrogen. The resulting residue was re-dissolved with 100 μ L of water/Me OH/TF-EtOH/PA (85:5:5:5). Calibration standards were also prepared by spiking standard solutions to 300 μ L of artificial CSF containing 0.25% rat plasma, with concentration ranges of 50–5000 pmol/L for A β_{1-38} , 100–10000 pmol/L for A β_{1-42} , and 2–200 pmol/L for A β_{1-43} .

LC-MS/MS analysis

Chromatographic separation was performed using the Shimadzu Nexera X2 system (Shimadzu Corporation, Kyoto, Japan) with a column of Acquity UPLC BEH C8 (1.7 μ m, 2.1 × 100 mm, Waters Corporation, MA, USA) set at 50°C. The mobile phases of (A) water/acetic acid (AA) (8:2) and (B) MeOH/ AA (8:2) were used with a linear gradient program as follows (Time - B conc.): 0 min-25%, 3.5 min-55%, 4.0 min-100%, 6.0 min-100%, 6.1 min-25%, and 8.0 min-25%. The flow rate was set at 0.4 mL/min. Mass spectra were obtained by a Qtrap 5500 system (AB Sciex, CA, USA) with the positive electrospray ionization and multiple reaction monitoring (MRM) mode. MRM transitions are shown in Table 2. MS/MS conditions were as follows: ion spray voltage at 5500 V, temperature at 600°C, curtain gas at 10 psi, ion source gas 1 at 90 psi, ion source gas 2 at 90 psi, and collision gas at level 11. Quantification was performed using Analyst 1.6.2 (AB Sciex, Redwood City, CA, USA). Calibration curves were constructed by analyzing calibration standards. The peak area ratios of analytes to the corresponding ISs

Table 2

MRM transitions. Mass spectra were obtained by the Qtrap 5500 system with the positive electrospray ionization (ESI) and multiple reaction monitoring (MRM) mode. MRM transitions are shown, respectively

| Analyte and IS | Precursor ion $[M+5H]^{5+}(m/z)$ | Product ion (<i>m/z</i>) |
|-----------------------------------|----------------------------------|-------------------------------|
| Αβ1-38 | 827.3 | 801.1 |
| $[^{15}N_{51}]$ -A β_{1-38} | 837.2 | 810.6 |
| Αβ ₁₋₄₀ | 867.0 | 843.8 |
| $[^{15}N_{53}]$ -A β_{1-40} | 877.2 | 853.6 |
| Αβ ₁₋₄₂ | 903.7 | 863.2 |
| $[^{15}N_{55}]$ -A β_{1-42} | 914.7 | 873.7 |
| Αβ ₁₋₄₃ | 924.1 | 1000.9 |
| $[^{15}N_{56}]$ -A β_{1-43} | 935.1 | 1012.8 |

were plotted against analyte concentrations, and the curves were fit by a linear least-squares regression method (a weighting factor of 1/X2).

ELISA

Sandwich ELISA was used to quantify $A\beta_{1-40}$ and A β_{1-42} levels using Human β Amyloid (1-40) ELISA Kit Wako II and a Human/Rat β Amyloid (1-42) ELISA Kit Wako, High-Sensitive according to the manufacturer's instructions (Wako Pure Chemical Industries, Ltd., Osaka, Japan). Microplates were precoated with monoclonal BAN-50 (IgG, anti-AB1-16) and sequentially incubated with 25 µL of samples followed by horseradish-peroxidase-conjugated BA 27 (anti-A β_{1-40} (Fab')2, specific for A β_{40}) or BC05 (anti-AB35-43 (Fab')2, specific for $A\beta_{42/43}$) [39]. Sensitivities were 0.019 pmol/L (assay range 1.0-100 pmol/L) in the A β_{1-40} assay and 0.06 pmol/L (assay range 0.01–20.0 pmol/L) in the A β_{1-42} assay. Intra- and inter-assay coefficients of variation were less than 10% in both assay systems.

Data analysis

Simple linear regression models were used in statistical analyses. The linear regression line was drawn by the method of least squares. Comparisons among the groups of assay values were performed using the Kruskal-Wallis test with multiple comparison tests for non-parametric data. Two-tailed *p*-values less than 0.05 were considered to be significant. These analyses were performed with IBM SPSS Statistics, version 24 (IBM Japan, Tokyo) and GraphPad Prism, version 8 (GraphPad Software, San Diego, CA, USA).

RESULTS

The minimal to maximal LC-MS/MS assay ranges of A β_{1-38} , A β_{1-40} , A β_{1-42} , and A β_{1-43} were 193.231~1,224.771, 391.624~3,560.087, 29.326~ 328.450, and 0.714~20.624 pmol/L, respectively (Table 3). CSF levels as mean ± standard deviation values of CU were 666 ± 249 pmol/L in A β_{1-38} , 2199 ± 725 pmol/L in A β_{1-40} , 153.7 ± 79.66 ± 79.7 pmol/L in A β_{1-42} , and 9.78 ± 4.58 pmol/L in A β_{1-43} . The ratios of A $\beta_{1-38/1-42}$, A $\beta_{1-38/1-43}$, A $\beta_{1-40/1-42}$, A $\beta_{1-40/1-43}$, and A $\beta_{(1-40/1-43)}/A\beta_{(1-38/1-42)}$ were 4.66 ± 0.94, 71.7 ± 15.8, 15.7 ± 3.7, 241 ± 58, and 53 ± 13, respectively. The ratio of the amounts of A β_{1-38} , A β_{1-40} , A β_{1-42} , and A β_{1-43} was approximately 68:225:16:1 (Table 4).

First order regression analysis showed a correlation between A β_{1-38} and A β_{1-40} ($Y=0.3035^*X+8.409$; $r^2=0.8395$; p<0.0001, Fig. 1a), between A β_{1-38} and A β_{1-42} ($Y=2.686^*X+247.8$; $r^2=0.547$; p<0.0001, Fig. 1b), between A β_{1-38} and A β_{1-43} ($Y=36.4^*X+317.9$; $r^2=0.424$; p<0.0001, Fig. 1c), between A β_{1-40} and A β_{1-42} ($Y=8.204^*X+860.3$; $r^2=0.5601$; p<0.0001, Fig. 1d), between A β_{1-40} and A β_{1-43} ($Y=112.8^*X+1064$; $r^2=0.4469$; p<0.0001, Fig. 1e), and between A β_{1-42} and A β_{1-43} ($Y=13.4^*X+27.09$; $r^2=0.7576$; p<0.0001, Fig. 1f).

 $A\beta_{1-40}$ and $A\beta_{1-42}$ values measured by LC-MS/ MS were strongly correlated with those obtained by ELISA. The first order regression analysis was Y = 0.9726 * X + 165.9 (r² = 0.8127, p < 0.0001) for AB1-40 values measured by LC-MS/MS and ELISA (Fig. 2a). Regarding $A\beta_{1-42}$ values, the regression curve was $Y = 0.8584 \times X + 2.545$ (r² = 0.6207, p < 0.0001) between LC-MS/MS and ELISA (Fig. 2b). Thus, these quantitative assays of A β_{1-40} and A β_{1-42} . by LC-MS/MS were equivalent to those by specific ELISA. Comparisons between $A\beta_{1-42} + A\beta_{1-43}$ levels by LC-MS/MS and AB1-42 values by ELISA showed the same relationship as those between LC-MS/MS A β_{1-42} levels and A β_{1-42} by ELISA (Y= 0.9100*X + 2.389, $r^2 = 0.6245$, p < 0.0001; Fig. 2c). These results show that conventional ELISA using BCO5 for A β 42/43 measures both A β ₁₋₄₂ and $A\beta_{1-43}$ in CSF.

The regression curve between the $A\beta_{1-38/1-42}$ ratio and $A\beta_{1-40/}A\beta_{1-43}$ ratio measured by LC-MS/MS

| Table 3 |
|---------|
|---------|

Basic profile of the LC-MS/MS analysis. The minimal to maximal assay ranges, median, mean, and standard deviations of $A\beta_{1-38}$, $A\beta_{1-40}$, $A\beta_{1-42}$, and $A\beta_{1-43}$ values measured by LC-MS/MS assay and ELISA

| Item | Αβ ₁₋₃₈ | Αβ ₁₋₄₀ | | Αβ ₁₋₄₃ | | |
|--------|--------------------|--------------------|----------|--------------------|---------|----------|
| | LC-MS/MS | LC-MS/MS | ELISA | LC-MS/MS | ELISA | LC-MS/MS |
| Max | 1224.771 | 3560.087 | 3401.273 | 328.450 | 306.233 | 20.624 |
| Min | 193.231 | 391.624 | 634.258 | 29.326 | 41.120 | 0.714 |
| Median | 506.495 | 1739.419 | 1555.430 | 100.789 | 122.838 | 5.798 |
| Mean | 546.038 | 1771.370 | 1619.538 | 111.056 | 126.683 | 6.267 |
| SD | 198.920 | 600.511 | 561.217 | 54.778 | 50.425 | 3.559 |

Assay data obtained by LC-MS/MS. Mean \pm standard values (pmol/L) of A β_{1-38} , A β_{1-40} , A β_{1-42} , and A β_{1-43} , and the ratios of A $\beta_{1-38/1-42}$, A $\beta_{1-38/1-42}$, A $\beta_{1-40/1-42}$, A $\beta_{1-40/1-43}$, and A $\beta_{(1-40/1-43)}/A\beta_{(1-38/1-42)}$ in respective disease groups including with ADD and ADMCI, ALS, CBD, 6 ENC, 3 FTD, MS, MSA, PD, PN, PSP, and SCD

Table 4

| Disease | mean level \pm standard deviation pmol/L | | | ratio±standard deviation | | | | | | | |
|---------------|--|--------------------|------------------------|--------------------------|----------------------|--------------------------|----------------------|---|-------------------------|----------------------|---|
| | Αβ ₁₋₃₈ | Αβ ₁₋₄₀ | Αβ ₁₋₄₂ | Αβ ₁₋₄₃ | $A\beta_{1-38/1-42}$ | $A\beta_{1-38/1-43}$ | $A\beta_{1-40/1-42}$ | $\begin{array}{c} A\beta_{1\text{-}40}/A\beta\\ _{1\text{-}42}+A\beta_{1\text{-}43}\end{array}$ | $A\beta_{1-40}/_{1-43}$ | $A\beta_{1-42/1-43}$ | $\begin{array}{c} A\beta_{(1-40/1-43)} \\ (_{1-38/1-42}) \end{array}$ |
| ADD/ ADMCI | 498 ± 166 | 1552 ± 527 | $71.93 \pm 27.7^{**a}$ | $3.15 \pm 2.66^{****b}$ | $7.38 \pm 2.28^{*e}$ | $250.2 \pm 184.2^{***f}$ | 22.5 ± 6.8 | 21.6 ± 6.6 | $759 \pm 520^{**g}$ | 32.5 ± 19.8 | 101 ± 61 |
| ALS | 647 ± 260 | 1986 ± 615 | 115.8 ± 44.6 | 6.68 ± 2.75 | 6.06 ± 2.55 | 107.4 ± 48.3 | 19.1 ± 8.3 | 18.0 ± 7.8 | 333 ± 138 | 18.0 ± 4.8 | 57 ± 13 |
| CBD | 474 ± 160 | 1634 ± 526 | 90.4 ± 50.1 | 4.82 ± 2.65 | 6.24 ± 2.54 | 125.1 ± 71.2 | 22.0 ± 10.5 | 20.6 ± 9.1 | 452 ± 324 | 21.7 ± 11.4 | 78 ± 50 |
| ENC | 503 ± 156 | 1571 ± 405 | 107.8 ± 54.4 | $4.39 \pm 2.49^{*c}$ | 5.05 ± 1.19 | 133.7 ± 53.2 | 15.9 ± 3.7 | 15.3 ± 3.6 | 429 ± 199 | $26.2\pm2.1^{*h}$ | 85 ± 30 |
| FTD | 671 ± 62 | 2132 ± 438 | 146.4 ± 63.1 | 9.07 ± 4.46 | 5.26 ± 2.36 | 91.4 ± 53.0 | 16.2 ± 5.9 | 15.3 ± 5.6 | 279 ± 134 | 16.8 ± 2.1 | 53 ± 3 |
| CU | 666 ± 249 | 2199 ± 725 | 153.7 ± 79.7 | 9.78 ± 4.58 | 4.66 ± 0.94 | 71.7 ± 15.8 | 15.7 ± 3.7 | 114.7 ± 3.3 | 241 ± 58 | 15.7 ± 3.4 | 53 ± 13 |
| MS | 468 ± 138 | 1604 ± 554 | 97.2 ± 37.6 | 5.58 ± 3.07 | 5.27 ± 2.14 | 120.2 ± 121.7 | 17.8 ± 6.6 | 16.9 ± 6.4 | 393 ± 352 | 20.6 ± 8.5 | 69 ± 26 |
| MSA | 498 ± 191 | 1531 ± 702 | 83.2 ± 33.9 | 5.41 ± 1.61 | 6.42 ± 2.72 | 93.6 ± 33.2 | 19.5 ± 7.9 | 18.2 ± 7.1 | 282 ± 82 | 15.3 ± 4.0 | 46 ± 13 |
| PD | 489 ± 127 | 1702 ± 518 | 97.6 ± 33.9 | $4.98 \pm 2.17^{*d}$ | 5.29 ± 1.23 | 110.7 ± 49.1 | 18.2 ± 4.2 | 17.3 ± 3.9 | 385 ± 181 | 21.2 ± 8.7 | 74 ± 34 |
| PN | 464 ± 165 | 1546 ± 491 | 114.4 ± 48.7 | 7.54 ± 1.94 | 4.30 ± 1.04 | 60.7 ± 10.3 | 14.3 ± 2.9 | 13.3 ± 2.5 | 202 ± 17 | 14.7 ± 4.2 | 50 ± 14 |
| PSP | 537 ± 260 | 1729 ± 830 | 129.6 ± 87.4 | 7.07 ± 5.31 | 4.61 ± 1.42 | 106.5 ± 81.3 | 14.8 ± 4.0 | 14.0 ± 3.8 | 342 ± 262 | 22.6 ± 15.1 | 73 ± 49 |
| SCD | 550 ± 124 | 1759 ± 352 | 118.1 ± 27.2 | 6.67 ± 2.15 | 4.71 ± 0.60 | 86.4 ± 17.8 | 15.2 ± 2.3 | 14.3 ± 2.2 | 284 ± 96 | 18.6 ± 4.7 | 62 ± 29 |
| total | 546 ± 199 | 1771 ± 601 | 111.1 ± 54.8 | 6.27 ± 3.56 | 5.44 ± 1.96 | 111.2 ± 89.1 | 17.7 ± 6.3 | 16.7 ± 5.9 | 370 ± 272 | 20.45 ± 10.2 | 67 ± 35 |

*** $^{a}p = 0.0066$; **** $^{b}p < 0.0001$; * $^{c}p = 0.0297$; * $^{d}p = 0.0281$; * $^{e}p = 0.021$; *** $^{f}p = 0.0004$; ** $^{g}p = 0.0033$; * $^{h}p = 0.0123$. ADD, Alzheimer's disease with dementia; ADMCI, AD due to MCI; ALS, anyotrophic lateral sclerosis; CBD, corticobasal degeneration; ENC, encephalopathy; FTD, frontotemporal dementia;), CU, cognitively unimpaired; MS, multiple sclerosis; MSA, multiple system atrophy; PD, Parkinson's disease; PN, peripheral neuropathy; PSP, progressive supranuclear palsy; SCD, spinocerebellar degeneration.



Fig. 1. Correlation among A β species measured by LC-MS/MS. First order regression analysis between A β_{1-38} and A β_{1-40} (a), between A β_{1-38} and A β_{1-42} (b), between A β_{1-42} (c), between A β_{1-42} (d), between A β_{1-40} and A β_{1-43} (e), and between A β_{1-42} and A β_{1-43} (f).

was Y = 0.003858*X+4.017 (r² = 0.2857, p < 0.0001; Fig. 2d), suggesting that this ratio is almost constant. The ratios of 8 out of 10 AD patients were outside the linear regression curve (Fig. 2d: Red circles indicate ADD and triangles indicate ADMCI). Although correlation analysis between measured levels by LC-MS/MS and ELISA in A $\beta_{1-40}/A\beta_{1-42}$ and A $\beta_{1-40}/A\beta_{1-42} + A\beta_{1-43}$ showed close relationships, most values were present in the leftmost regression curve (Fig. 1e, f).

We then examined the efficacy of LC-MS/MS assay values as biomarkers for AD by comparisons with assay values among ADD/ADMCI, CU, and other neurogenerative diseases. No significant differences were observed in $A\beta_{1-38}$ and $A\beta_{1-40}$ levels among ADD/ADMCI and other neurological



Fig. 2. Correlation between LC-MS/MS and ELISA. $A\beta_{1-40}$, $A\beta_{1-42}$, and $A\beta_{1-42} + A\beta_{1-43}$ values measured by LC-MS/MS were strongly correlated with those obtained by ELISA (a: Y=0.9726*X+165.9, $r^2=0.8127$, p<0.0001 for $A\beta_{1-40}$ values; b: Y=0.8584*X+2.545, $r^2=0.6207$, p<0.0001 for $A\beta_{1-42}$; and c: comparisons between $A\beta_{1-42} + A\beta_{1-43}$ levels by LC-MS/MS and $A\beta_{1-42}$ values by ELISA: Y=0.9100*X+2.389, $r^2=0.6245$, p<0.0001). The regression curve between the $A\beta_{1-38}/_{1-42}$ ratio and $A\beta_{1-40}/A\beta_{1-43}$ ratio measured by LC-MS/MS was almost constant. The ratios of 8 out of 10 AD patients were outside the linear regression curve (d: Red circles • represent sporadic ADD, red triangles • represent sporadic ADMCI, brown circles • represent familial AD). Correlation analysis between measured levels by LC-MS/MS and ELISA in $A\beta_{1-40}/A\beta_{1-42}$ and $A\beta_{1-40}/A\beta_{1-42} + A\beta_{1-43}$ showed close relationships, but most values were present in the leftmost regression curve (e and f).

diseases and CU (Fig. 3a, b). $A\beta_{1-42}$ levels were significantly lower in ADD/ADMCI than in CU (p = 0.0066; Fig. 3c). CSF A β_{1-43} values were significantly lower in ADD/ADMCI (Fig. 3d; p < 0.0001), encephalopathy (p = 0.0297), and Parkinson's disease (p = 0.0281) than in CU. The A $\beta_{1-38}/A\beta_{1-42}$ and A $\beta_{1-38}/A\beta_{1-43}$ ratios were significantly higher in ADD/ADMCI than in CU (p = 0.021 in A $\beta_{1-38/1-42}$, Fig. 3e; p = 0.0004 in A $\beta_{1-38/1-43}$, Fig. 3f). Increased ratios of A $\beta_{1-40}/A\beta_{1-42}$ and A $\beta_{1-40}/A\beta_{1-42} +$ A β_{1-43} were not significant (p = 0.1226 in A $\beta_{1-40}/A\beta_{1-42} +$ A β_{1-42} ratio, Fig. 3g; p = 0.0895 in A $\beta_{1-40}/A\beta_{1-42} +$ A β_{1-43} ratio). The A $\beta_{1-40}/A\beta_{1-43}$ ratio was significantly increased (p = 0.0033, Fig. 3h). The



Fig. 3. Values of CSF Aβ species and ratios among neurological diseases and controls. No significant differences were observed in A β_{1-38} and A β_{1-40} levels among ADD/ADMCI and other neurological diseases and CU (a, b). A β_{1-42} levels were significantly lower in ADD/ADMCI than in CU (p = 0.0066; c). CSF A β_{1-43} values were significantly lower in ADD/ADMCI (d; p < 0.0001), ENC (p = 0.0297), and PD (p = 0.0281) than in CU. The A $\beta_{1-38}/A\beta_{1-42}$ and A $\beta_{1-38}/A\beta_{1-43}$ ratios were significantly higher in ADD/ADMCI than in CU (e: p = 0.021; f: p = 0.0004). Ratios of A $\beta_{1-40}/A\beta_{1-42}$ and A $\beta_{1-40}/A\beta_{1-42} + A\beta_{1-43}$ were not significant (g: p = 0.1226 in A $\beta_{1-40}/A\beta_{1-42}$ ratio; p = 0.0895 in A $\beta_{1-40}/A\beta_{1-42} + A\beta_{1-43}$ ratio was significantly increased (h: p = 0.0033). The A $\beta_{(1-40)}/A\beta_{(1-38)}/A\beta_{(1-38)}/1-42$) ratio was not significantly changed in ADD/ADMCI and other diseases (i). (a: Blue circles \bullet represent sporadic ADD and ADMCI, cyan circles \bullet represent familial AD).

 $A\beta(_{1-40/1-43})/A\beta(_{1-38/1-42})$ ratio was not significantly changed in ADD/ADMCI and other diseases. These results imply that successive cleavage from $A\beta_{1-43}$ to $A\beta_{1-40}$ and from $A\beta_{1-42}$ to $A\beta_{1-38}$ was enhanced in parallel in AD.

DISCUSSION

Mass spectrometry is a device that is capable of detecting the masses of various substances, as well as accurate peptide sequence lengths. Due to advances in the operability and analytical devices for extremely small amounts, this method has been clinically applied to the measurement of various substances in recent years. MS/MS is the combination of two mass analyzers in one mass spectrometric instrument. The combination with a highly selective separation method by LC allows for more accurate and reliable quantification for clinical application. Therefore, the LC-MS/MS technique has been used to measure A β peptides [7–9, 17–19, 26–29]. This method eliminates time-consuming immune assays with a large number of steps and multiple antibodies or immunoprecipitation steps for pre-clinical studies. This approach also allows for performance of a single assay to measure several different A β species simultaneously in one sample with selectivity, specificity, and in a high-throughput format, while still achieving the high sensitivity required for low level endogenous A β peptides (such as CSF A β_{1-43}), plasma A β species, and modified tau products.

Comparative assays using the same samples between the LC-MS/MS assay and commonly used ELISA systems for A β_{1-40} , A β_{1-42} , and A β_{1-42+} A β_{1-43} confirmed a strong correlation (Y=0.9726* X + 165.9 for $A\beta_{1-40}$; Y = 0.8584 * X + 2.545 for $A\beta_{1-42}$; Y = 0.9100 * X + 2.389 for $A\beta_{1-42} + A\beta_{43}$; Fig. 2a-c). These results suggest that this LC-MS/MS assay is as reliable and reproducible a system as conventional ELISA systems. We did not validate $A\beta_{1-38}$ or A β_{1-43} assay values by comparisons with ELISA systems. This LC-MS/MS assay requires further improvements in sensitivity for lower A β_{1-43} values. However, studies on relationships with the respective AB species showed strong linear correlations among A β_{1-38} , A β_{1-40} , A β_{1-42} , and A β_{1-43} (Fig. 1). Reliable ELISA for $A\beta_{1-38}$ and $A\beta_{1-43}$ are not commonly available. The present results support the application of the LC-MS/MS assay system for clinical usage, similar to conventional ELISA systems.

We previously proposed that $A\beta_{1-38/1-42}$ and $A\beta_{1-40/1-43}$ were distinct product/precursor pairs and reflected γ -secretase activity [9]. Significantly decreased CSF levels of A β_{1-42} and A β_{1-43} accompanied by increased A $\beta_{1-38/1-42}$ and A $\beta_{1-40/1-43}$ ratios and an enhanced proportional ratio of A $\beta_{1-40/1-43}$ versus A_{β1-38/1-42} suggested altered successive cleavage by γ -secretase in AD. In the present study, strong correlations between A β_{1-38} and A β_{1-40} (r² = 0.8395) and between A β_{1-42} and A β_{1-43} (r² = 0.7576) also support this cleavage being processed in parallel. The processing ratio from $A\beta_{1-42}$ to $A\beta_{1-38}$ was 2.686 (Fig. 1b), and that from $A\beta_{1-43}$ to $A\beta_{1-40}$ was 112.8 (Fig. 1f). The accurate ratios of A $\beta_{1-38/1-42}$ were 4.66 in CU and 7.38 in AD whereas those of $A\beta_{1-40/1-43}$ were 241 in CU and 795 in AD. Both cleavage processes appear to have been enhanced in parallel twice in AD (Fig. 2d). Recent studies revealed that CNS-derived A β is cleared into the CSF [7] and peripheral blood [40], and that the clearance rate decreases in late onset AD [8] and is differently regulated by age, APOE isoforms, and the presence of AB amyloidosis [41-44]. Therefore, the measurement of precise amounts of A β_{1-38} , A β_{1-40} , A β_{1-42} , and AB1-43 is relevant for clinical application of biomarkers and understanding of the processing of AB species in AD.

Comparisons between 66 cases of early-onset AD and 25 of late-onset AD by ELISA showed that CSF $A\beta_{1-43}$ significantly varied with age in AD. However, A β_{1-43} levels were not superior to A β_{1-42} for diagnostic accuracy [15]. CSF A β_{1-43} , but not A β_{1-42} , differentiated between patients with amnestic MCI who progressed to AD during a 2-year followup [16]. Goda and Kobayashi evaluated dog CSF $A\beta_{1-38}$, $A\beta_{1-40}$, $A\beta_{1-42}$, and $A\beta_{1-43}$ using peptide adsorption-controlled LC-MS/MS and showed that the A β assay was reproducible [17]. A β quantitation by LC-MS/MS showed satisfactory intra- and interday reproducibility as well as freeze-thaw and room temperature stability [17, 18]. The findings of a recent large study by Janelidze et al. using three different immunoassays revealed that the CSF A $\beta_{42}/A\beta_{40}$ and A $\beta_{42/38}$ ratios were significantly better than CSF AB42 alone for the detection of brain amyloid deposition in prodromal AD and differentiating AD dementia from non-AD dementias [20]. Widely used ELISA for the quantitation of $A\beta_{1-42}$ /total tau, $A\beta_{1-42}/A\beta_{1-40}$, $A\beta_{1-42}/A\beta_{1-38}$, and $A\beta_{1-42}$ showed the greatest accuracy for identifying amyloid-positive subjects. A β_{1-40} and A β_{1-38} showed significantly lower discriminative powers [21]. The first report of ELISA assays for $A\beta_{1-38}$, $A\beta_{1-40}$, and $A\beta_{1-42}$ showed only decreased levels of $A\beta_{1-42}$ compared with $A\beta_{1-38}$ and $A\beta_{1-40}$ in CSF from AD [22]. CSF $A\beta_{1-42}/A\beta_{1-40}$ and $A\beta_{1-42}/A\beta_{1-43}$ were positively correlated with in gait variability and ADL in mild ADD [23]. A UPLC-MS-MS assay of CSF A β_{1-42} showed equivalent diagnostic utility to those of commonly used AlzBio3 ELISA systems [26]. Oe et al. reported that the validated methods had limits of quantitation of 44 fmol/mL for A β_{1-42} and 92 fmol/mL for AB1-40 in CSF. A strong correlation was observed between LC/MS/MS and ELISA for $A\beta_{1-42}$ (r² = 0.915), and those correlations were weak for A β_{1-40} (r² = 0.644) [27]. The accuracy and precision of LC-MS/MS will contribute to the clinical diagnosis and monitoring of therapeutic interventions [27].

The present results suggest that the quantitative LC-MS/MS assay for CSF A β species is as reliable as specific ELISA for the clinical evaluation of CSF biomarkers for AD. Since the present study was not a large scale or prospective control study, further large-scale studies are necessary. However, the reliable and convenient LC-MS/MS multiple assay of A β species is equivalent to specific multiple ELISA assays. Therefore, the LC-MS/MS multiple assay of A β species may be applicable and contribute to kinetic [7, 8] and drug monitoring [44] studies on A β species. The establishment of a multiple A β and tau species assay by LC-MS/MS is warranted.

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