

Development of a High-Sensitivity Method for the Measurement of Human Nasal A β ₄₂, Tau, and Phosphorylated Tau

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Abstract. Cost-effective and feasible methods for early diagnosis of Alzheimer's disease (AD) are needed. We present two methods to measure AD-related biomarkers simultaneously from one nasal smear for the purpose of diagnosing AD. Japanese men and women aged 63–85 years old were recruited in 2015–2016 for this case–control study. A total of 25 AD cases and 25 controls (22 men and 28 women) participated in this research. Nasal smears were collected from the common nasal meatus, inferior concha, middle nasal meatus, and olfactory cleft, and the proteins in the samples were analyzed by two methods, which we named PGD (Pre-treatment with guanidine- n-Dodecyl-beta-D-maltoside solution) method 1 (PGD-I) and 2 (PGD-II). The PGD-I method measured total tau and amyloid- β (A β)₄₂, but no differences in median levels of total tau and A β ₄₂ between AD cases and controls were found in any of the nasal locations. The PGD-II method measured A β ₄₂, total tau, and phosphorylated tau, but levels of A β ₄₀ in all nasal locations of both groups were near zero. Median levels of phosphorylated tau to total tau (p-tau/t-tau) ratios in the middle nasal meatus and in the olfactory cleft were significantly higher in AD cases than in controls, and could significantly predict AD. To assess diagnostic reliability, areas under the ROC curve were 0.74 (95% CL = 0.52–0.95, $p = 0.030$) for the middle nasal meatus and 0.72 (95% CL = 0.52–0.92, $p = 0.029$) for the olfactory cleft. Thus, PGD-I and PGD-II can detect AD-related biomarkers in nasal smears and PGD-II may be a useful tool for diagnosing AD.

Keywords: Amyloid- β , biomarker, nasal smear, phosphorylated tau, tau

INTRODUCTION

According to a report by Alzheimer's Disease International, in 2013 the number of people with dementia was estimated at 4.4 million, and was expected to rise to 76 million in 2030 and 135 million in 2050. Since about 60%–70% of people

with dementia have Alzheimer's disease (AD), it is important to find effective methods for diagnosing and treating AD. The pathological hallmarks of AD are senile plaques, consisting of amyloid- β (A β) peptides, and neurofibrillary tangles, formed from abnormally phosphorylated tau within the cytoplasm of neurons [1]. Recent advances in AD research show that amyloid pathology begins some 10–20 years before the clinical onset of AD [2–4]. Therefore, it is very important to use not only clinical assessments but also biomarkers for AD diagnosis [5].

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Recently, diagnostic tools such as structural magnetic resonance imaging (sMRI) and molecular neuroimaging with positron emission tomography (PET) have been under development, and are being applied in clinical settings [6]. However, MRI and PET are expensive. An alternate diagnostic approach is biomarker analysis via cerebrospinal fluid (CSF) [7]. Studies have shown that increased levels of total tau (t-tau) and phosphorylated tau (p-tau) proteins, and decreased CSF A β ₄₂ levels can be used to differentiate between AD cases and non-demented, neurologically normal people (“controls”) [7–9]. These CSF markers may also be useful in predicting progress from mild cognitive impairment to AD [10]. CSF biomarker analysis also has its drawbacks—it requires lumbar puncture, which is an invasive procedure with potentially adverse side effects. Therefore, alternative, more cost-effective, less-invasive, convenient, and feasible methods for early diagnosis of AD are needed.

The nasal mucosa is a target tissue for measuring AD biomarkers such as A β ₄₂, tau, and phosphorylated tau (p-tau). Olfactory dysfunction is an early sign of AD, as most patients suffer from anosmia [11]. A recent study found that A β deposition in olfactory epithelium can predict mild cognitive impairment and is associated with AD and dementia [12]. The olfactory bulb is often severely affected at an early stage in AD cases [13]. In addition, A β and tau deposits have been found in the olfactory epithelium of AD cases at autopsy [14, 15]. Animal experiments have shown that ventricle-injected isotope-labelled beta-amyloid peptide (¹²⁵I-A β 40) is transported to the nasal cavity through a non-hematogenous pathway [16]. Deposits of A β peptides in the nasal mucosa in a transgenic mouse model of AD have been found to be well correlated with the amounts of A β depositions in the brain [17]. These studies suggest that the nasal cavity may be a useful site for early diagnosis of AD.

We recently reported a highly sensitive method for measuring the contents of A β ₄₂ in human nasal epithelium [18]. However, when we applied this method to measurement of A β ₄₂ in nasal smears from AD cases and controls, the data showed very high variability (data not shown here). Since nasal epithelium bears variable amounts of nasal discharge, it is difficult to standardize the values for total protein concentrations. Therefore, we tried to develop a new method to simultaneously measure AD biomarkers such as A β ₄₂, A β ₄₀, tau, and phosphorylated tau from a single nasal smear. We now present two novel methods for simultaneously detecting

biomarkers in human nasal smears. Each of these two methods was separately applied to the nasal smear samples, collected at four different sites from AD cases and from controls. Then, biomarker levels were compared between AD patients and controls.

MATERIALS AND METHODS

Study population

Elderly men and women were recruited from Shiga University of Medical Science Hospital in Kusatsu, Japan from 2015 to 2016. All participants provided written informed consent. This study complies with the Declaration of Helsinki and was approved by the Ethics Committee of the Shiga University of Medical Science (Approval number = 25–218). Finally, swabs from 19 people (8 control cases and 11 AD cases) were measured by PGD (Pre-treatment with guanidine-n-Dodecyl-beta-D-maltoside solution) method 1 (PGD-I), and swabs from 31 people (17 control cases and 14 AD cases) were measured by PGD method 2 (PGD-II).

Participants’ medical records, including history of disease, medication, and complications, were collected. Blood samples were taken from participants, and magnetic resonance imaging (MRI) was performed. To confirm the presence or absence of dementia, we examined all subjects following established hospital diagnostic protocols including the Mini-Mental State Examination (MMSE), Clinical Dementia Rating, Wechsler Memory Scale (WMS)-RI, WMS-RIL, Alzheimer’s Disease Assessment Scale-cog (ADAS-cog), and Global Deterioration Scale (GDS) (Table 1).

Human nasal smear collection and pre-treatments

A designated otolaryngologist collected nasal smears from the left inferior concha, middle nasal meatus, olfactory cleft, and common nasal meatus using cotton swabs (HUBY-COTIX EN3-50S; Sanyo Co., Ltd., Osaka, Japan) (Fig. 1). All nasal smears were placed in microtubes, and nasal samples were extracted using the following methods.

Crude extraction

Each cotton swab was added to a microtube containing 100 μ l of 50 mM Tris HCL buffer (pH 8.0)

Table 1
Characteristics of AD cases and controls

Characteristics	AD case (n = 25)		Control (n = 25)	
	Median	IQR	Median	IQR
Age (y)	76	73–79	71	67–77
Sex (% male)		48.0		40.0
Diabetes (%)		20.0		16.0
Hypertension (%)		24.0		20.0
Hyperlipidemia (%)		24.0		24.0
Glucose Intolerance (%)		4.0		0.0
Clinical Dementia Rating Composite Rating (%)				
0 (None)		0		100
0.5 (Very Mild)		40		0
1 (Mild)		40		0
2 (Moderate)		20		0
3 (Severe)		0		0
MMSE	23.0	18.5–26.0	29.0	27.5–30.0
WMS-R I	5.0	3.0–8.0	16.0	13.5–22.0
WMS-R II	0.0	0.0–0.5	12.0	7.5–17.5
ADAS-cog	14.0	10.5–19.0	5.0	3.5–6.0
GDS	1.0	0.0–3.0	2.0	0.0–7.0

Data are presented as medians and interquartile range (IQR) or as percentages of individuals having the characteristic, as indicated.

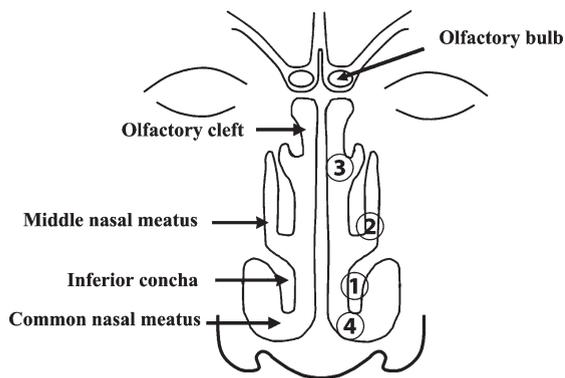


Fig. 1. Anatomy of the nasal cavity. Nasal smears were taken from the inferior concha (1), middle nasal meatus (2), olfactory cleft (3), and common nasal meatus (4) in that order.

containing 8M guanidine HCL, 200 mM NaCl, 2 mM EDTA, and 0.05% n-Dodecyl-beta-D-maltoside and was vortexed vigorously (we refer to this solution as the guanidine solution). The cotton swab was then removed from the microtube. The solution in the microtube was heated for 10 min at 70°C. The solution was put into a 0.2 µm PTFE spin filter (Ultrafree-MC 0.2, Millipore Japan) set in a new microtube. The solution was then filtered by centrifugation (14,000 × g, 30 min, 25°C). A volume of 90 µl of the filtered solution was divided into two tubes containing 45 µl each for separate Aβ and tau measurements.

Aβ measurements

Each 45 µl sample of filtered solution was placed in a 300 kDa Nanosep® Centrifugal Filter (Paul Japan Corp., Tokyo, Japan) and filtered by centrifugation at 14,000 × g for 15 min at 25°C. After centrifugation, 90 µl of 50 mM Tris HCL buffer (pH 8.0) containing 200 mM NaCl, 1 mM EDTA and 0.05% n-Dodecyl-beta-D-maltoside (we call this the wash buffer) was added onto the same filter membrane. The solution was filtered into the same tube by centrifugation at 14,000 × g for 15 min at 25°C. The filtered solution was collected and mixed well. A volume of 120 µl of the filtered solution and then 40 µl of the wash buffer were placed in 1 ml of the G-10 column set in a Bio-Spin Chromatography Column (Bio-Rad, Hercules, CA, USA) in order to remove guanidine HCl. The Bio-Spin Column was spun for 1 min at 800 × g and the filtered solution (~160 µl) was collected. Triplicates of 5 µl (total 15 µl) were used for the measurements of total protein concentration using the Protein Assay Rapid Kit WAKO (Wako Pure Chemical Industries, Ltd., Osaka, Japan), and the remaining solutions for the measurements of Aβ₄₂ and Aβ₄₀ concentrations were applied to Human/Rat Beta Amyloid (42) ELISA Kit Wako, High Sensitive (Wako Pure Chemical Industries, Ltd.) and Human Beta Amyloid (40) ELISA Kit Wako II (Wako Pure Chemical Industries, Ltd.).

Tau measurements

We used two different methods: PGD-I and PGD-II.

PGD-I

A volume of 45 μ l of the filtered solution was added to 45 μ l of the guanidine solution and mixed well. The mixed solution (80 μ l) and the wash buffer (80 μ l) were placed into 1 ml of the G-10 column set in a Bio-Spin Chromatography Columns (Bio-Rad), in order to remove guanidine hydrochloride (HCl). The Bio-Spin Column was spun for 1 min at 800 \times g and the filtered solution (\sim 160 μ l) was collected. Triplicates of 5 μ l (total 15 μ l) were used for the measurements of total protein concentration using the Protein Assay Rapid Kit WAKO (Wako Pure Chemical Industries). The remaining solution was used to measure total tau concentrations using TAU (Total Human ELISA Kit (Invitrogen)).

PGD-II

A volume of 40 μ l of the filtered solution and then 10 μ l of the wash buffer were placed in 0.7 ml of G-10 column set in a Bio-Spin Chromatography Column (Bio-Rad), in order to remove guanidine HCl. The Bio-Spin Column was spun for 1 min at 800 \times g and the filtered solution (\sim 50 μ l) was collected. Triplicates of 2 μ l (6 μ l total) were used to measure total protein concentration using the Protein Assay Rapid Kit WAKO (Wako Pure Chemical Industries, Ltd.). Volumes of 10 and 25 μ l of the filtered solution were used to measure total tau and phosphorylated tau (p-tau) concentrations using the TAU (Total Human ELISA Kit (Invitrogen) and TAU [pT181] Human ELISA Kit (Invitrogen), respectively.

We measured nasal fluid samples from a total of 11 AD cases and 8 controls using the PGD-I method, and samples from 14 AD cases and 17 controls using the PGD-II method (Table 2).

Table 2
Number of control and AD case participants according to nasal location and method

Location	PGD-I		PGD-II	
	AD case	Control	AD case	Control
Olfactory cleft	11	8	14	17
Middle nasal meatus	11	8	14	17
Inferior concha	11	8	14	17
Common nasal meatus	11	8	10	15

Statistical analysis

Participants were classified by cognitive status, as AD case or control. Characteristic data were presented as medians and interquartile ranges (IQRs) or percentages for prevalence. Measured protein concentrations (standardized to total protein) were compared between AD cases and controls in both PGD-I and PGD-II methods. Protein levels were presented as medians and IQRs. Levels of p-tau/t-tau from PGD-II were compared between AD cases and controls. Data were presented as a column scatter plot using GraphPad Prism Version 7. Receiver operating characteristics (ROC) curves and areas under the curve (AUC) were used to assess the reliability of nasal protein levels for predicting AD. Youden's Index (sensitivity + specificity - 1) was used to determine the optimal cut-offs. Wilcoxon's rank-sum test was used to compare medians between AD cases and controls.

All statistical analyses were performed using SAS version 9.4 (SAS Institute, Cary, North Carolina). Two-tailed *p*-values of <0.05 were considered as significant.

RESULTS

Characteristics of study participants are presented in Table 1. A total of 25 AD cases and 25 controls were recruited. In total, there were 22 men and 28 women. The median ages for AD cases and controls were 76 and 71 years old, respectively. The prevalence of diabetes, hypertension, and hyperlipidemia were not significantly different between AD cases and controls. AD cases exhibited higher composite ratings for clinical dementia and lower scores on cognitive tests including the MMSE, WMS-R I, WMS-R II, ADAS-cog, and GDS (Table 1).

The results of PGD-I and PGD-II tests are shown in Tables 3 and 4, respectively. We collected nasal smears from four regions (Fig. 1). Because of feeling of discomfort, some people refused sample collections in process, and in some cases, we failed to collect a sufficient volume of specimens, so sample numbers in Tables 3 and 4 were different from those in Table 2.

Protein measurements by PGD-I from nasal smears of the inferior concha, middle nasal meatus, olfactory cleft, and common nasal meatus are presented in Table 3. Total tau (t-tau) and A β ₄₂ were identified in AD cases and controls. T-tau levels were highest in the common nasal meatus and lowest in the

Table 3
Proteins measured in different nasal locations by PGD-I method in AD cases and controls

Location	AD case			Control			p-value
	Sample (n)	Median	IQR	Sample (n)	Median	IQR	
t-tau							
Olfactory cleft	10	567.7	517.5–634.8	7	636.4	426.7–675.9	0.887
Middle nasal meatus	9	576.0	529.1–713.9	7	468.6	260.1–751.4	0.370
Inferior concha	10	618.6	418.7–781.8	8	640.8	355.3–726.4	0.762
Common nasal meatus	10	704.4	437.7–858.2	8	721.6	513.9–855.2	1.000
Aβ ₄₂							
Olfactory cleft	8	2.00	1.40–3.20	8	1.60	0.90–2.00	0.525
Middle nasal meatus	8	1.90	0.90–4.80	8	1.80	1.40–2.40	0.623
Inferior concha	8	2.50	0.00–4.00	7	0.40	0.00–1.60	0.326
Common nasal meatus	10	2.40	1.70–3.60	8	1.70	0.60–2.80	0.459

p-values were derived from Wilcoxon test for comparison of medians.

Table 4
Proteins measured in different nasal locations by PGD-II method in AD cases and controls

Location	AD case			Control			p-value
	Sample(n)	Median	IQR	Sample(n)	Median	IQR	
t-tau							
Olfactory cleft	13	265.5	143.5–391.3	17	367.7	256.6–735.8	0.250
Middle nasal meatus	12	194.8	54.8–539.9	17	335.1	266.0–562.0	0.328
Inferior concha	14	301.0	154.0–909.4	16	522.0	185.9–985.9	0.727
Common nasal meatus	9	327.9	194.0–1290.0	15	304.5	253.5–699.4	1.000
p-tau							
Olfactory cleft	13	31.0	24.5–47.5	16	11.7	0.00–63.8	0.459
Middle nasal meatus	12	29.9	0.00–52.3	17	0.00	0.00–41.5	0.176
Inferior concha	14	0.00	0.00–0.00	16	0.00	0.00–0.00	0.581
Common nasal meatus	9	0.00	0.00–0.00	15	0.00	0.00–0.00	0.884
Aβ ₄₂							
Olfactory cleft	13	0.00	0.00–1.29	17	0.00	0.00–0.72	0.469
Middle nasal meatus	13	1.02	0.00–1.44	17	0.00	0.00–0.00	0.250
Inferior concha	14	0.00	0.00–1.38	17	0.00	0.00–1.42	1.000
Common nasal meatus	9	0.00	0.00–0.00	15	0.00	0.00–1.08	1.000
Aβ ₄₀							
Olfactory cleft	13	0.00	0.00–0.01	17	0.01	0.00–0.01	0.117
Middle nasal meatus	13	0.00	0.00–0.01	17	0.00	0.00–0.01	0.987
Inferior concha	14	0.00	0.00–0.01	17	0.01	0.00–0.01	0.481
Common nasal meatus	8	0.00	0.00–0.01	15	0.00	0.00–0.01	0.765
p-tau/t-tau*							
Olfactory cleft	12	0.20	0.00–0.20	15	0.00	0.00–0.10	0.047
Middle nasal meatus	9	0.20	0.00–0.20	14	0.00	0.00–0.10	0.048
Inferior concha	13	0.00	0.00–0.00	13	0.00	0.00–0.00	0.864
Common nasal meatus	9	0.00	0.00–0.00	12	0.00	0.00–0.00	1.000

p-values were derived from Wilcoxon test for comparison of medians. *p-tau/t-tau is a ratio (has no units).

olfactory cleft for AD cases. In controls, t-tau levels were highest in the common nasal meatus and lowest in the middle nasal meatus. Median t-tau levels were not found to be significantly different between AD cases and controls in any of the nasal locations.

Aβ₄₂ levels were highest in the inferior concha and lowest in the middle nasal meatus in AD cases, whereas the opposite was found in controls—Aβ₄₂ levels were highest in the middle nasal meatus and lowest in the inferior concha. However, no

significant differences were observed in median levels of Aβ₄₂ between AD cases and controls in any of the locations.

Table 4 shows protein measurements by PGD-II of t-tau, p-tau, Aβ₄₂, and Aβ₄₀. T-tau was highest in the common nasal meatus and lowest in the middle nasal meatus in AD cases. Unlike PGD-I, PGD-II found that t-tau was highest in the inferior concha and lowest in the common nasal meatus of controls. Levels of PGD-II-measured Aβ₄₂ were low in most

locations in AD cases and controls. No significant differences in median levels of t-tau and A β_{42} in any of the locations were found from PGD-II. Median A β_{40} levels were near zero in all nasal locations and were not significantly different between AD cases and controls in any of the nasal locations. Median p-tau was identified only in the middle nasal meatus and olfactory cleft in AD cases and controls. Although median p-tau levels appeared higher in AD cases than in controls, these differences were not statistically significant in any of the nasal locations.

The ratios of p-tau/t-tau are shown in Fig. 2. Median values of p-tau/t-tau were significantly higher in AD cases than in controls in the middle nasal

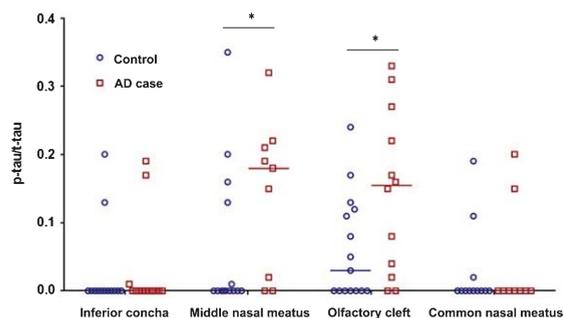


Fig. 2. P-tau/t-tau ratios in different nasal locations in AD cases and controls, measured by the PGD-II method. Asterisks represent significant differences ($p < 0.05$) between AD case and control median values. Wilcoxon's p -values are 0.048 and 0.047 for the middle nasal meatus and olfactory cleft, respectively.

meatus and in the olfactory cleft. No significant differences in median p-tau/t-tau levels were observed in the inferior concha and the common nasal meatus between AD cases and controls.

The receiver operating characteristics (ROC) curves for the ratio of p-tau/t-tau in predicting dementia are displayed in Fig. 3. The middle nasal meatus p-tau/t-tau ROC curve has an AUC of 0.738, with 95% confidence limits (95% CL)=0.523–0.953, p -value (p)=0.030 (Fig. 3A). The ROC curve AUC for the ratio in the olfactory cleft was 0.722, 95% CL=0.523–0.9214, p =0.029. Cut-off scores for each test were identified using Youden's Index (sensitivity + specificity—1) as this is often used when there is no preference for sensitivity over specificity [19]. Cut-off scores with the highest Youden's Index value were 0.02 (sensitivity=0.78 and specificity=0.71) for the middle nasal meatus and 0.15 (sensitivity=0.58 and specificity=0.87) for the olfactory cleft. A table of cut-off values with their associated sensitivity, specificity, and Youden's Index values can be found in Supplementary Tables 1 and 2.

DISCUSSION

Using PGD-II, we could successfully measure levels of t-tau, p-tau, and A β_{42} simultaneously in nasal smears. Levels of A β_{40} in many nasal locations were not detectable. We had previously developed a

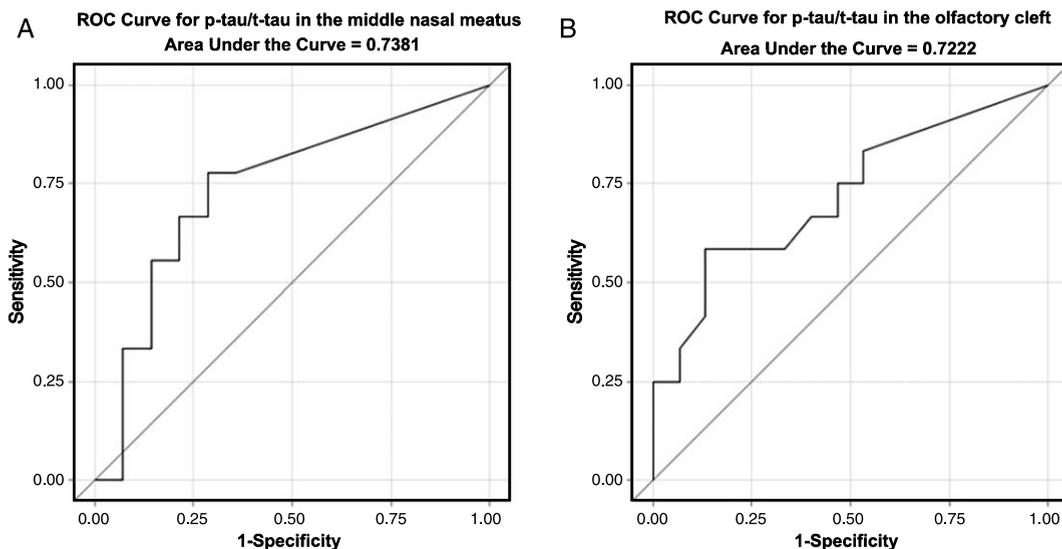


Fig. 3. Receiver operating characteristic (ROC) curves for p-tau/t-tau in (A) the middle nasal meatus and (B) the olfactory cleft, for predicting AD. The area under the curve (AUC) is 0.738, 95% Confidence Limits (95% CL)=0.523–0.953, p =0.030 for A and 0.722, 95% CL=0.523–0.921, p =0.029 for B.

high-sensitivity method for measuring levels of $A\beta_{42}$ from human nasal smears [18]. Since $A\beta_{42}$ in nasal epithelium is aggregated and insoluble, we used formic acid treatments in order to dissolve the aggregated $A\beta_{42}$ [18]. However, we needed to neutralize the solution and then concentrate the neutralized solution before ELISA measurements [18]. In addition, t-tau and p-tau are degraded by formic acid preparation. This requires another nasal smear sample from the individual just to measure levels of tau and p-tau. A major disadvantage of this is that nasal fluid amount is limited and the extraction of many samples may contaminate subsequent samples. Instead of formic acid, we used guanidine HCl to dissolve aggregated proteins in this study. We use a G-10 column to remove guanidine HCl, and then tested samples by ELISA methods.

The PGD-I method measured $A\beta_{42}$, and t-tau, and PGD-II measured $A\beta_{42}$, t-tau, and p-tau, but median levels of $A\beta_{40}$ in all nasal locations of both groups were very low (0 or 0.01). Neither method found that levels of t-tau, p-tau, $A\beta_{42}$, and $A\beta_{40}$ were significantly different between AD cases and controls in any of the nasal locations. However, the p-tau/t-tau ratios in samples from the olfactory cleft and middle nasal meatus were significantly higher in AD cases than in controls. We found that p-tau/t-tau in both locations can significantly predict AD. One study has shown high sensitivity and specificity when using the p-tau/t-tau ratio in CSF to diagnose AD [7]. Hyperphosphorylation of tau is a major pathological change in AD and is critical in the formation of neurofibrillary tangles, and tau proteins have been found to be abnormally deposited in the dystrophic olfactory epithelium of AD patients [14, 15]. The increased p-tau/t-tau ratio in the nasal cavity could be the result of pathologic changes in the olfactory epithelium in AD.

We collected nasal smears from four regions. Because of feelings of discomfort, some people refused sample collections in processes, and in some cases, we failed to collect a sufficient volume of specimens. However, since we found that we could measure levels of t-tau, p-tau, and $A\beta_{42}$ from a single sample, the discomfort of sampling can be minimized.

Some studies have shown that the plasma $A\beta_{42}/A\beta_{40}$ ratio can be a good marker to identify AD [20–22]. In our study, the $A\beta_{42}/A\beta_{40}$ ratio in nasal smears did not significantly differ between AD cases and controls. However, the $A\beta_{42}/A\beta_{40}$ ratio could only be calculated in a few cases and control

samples as the level of $A\beta_{40}$ in many nasal locations seemed to be zero. Thus, due to the small sample size, our $A\beta_{42}/A\beta_{40}$ ratio results may not be conclusive. Further studies with increased sample sizes are needed.

Although $A\beta$ peptides and tau proteins are found in nasal epithelium [14], the sources of the detected biomarkers in the nasal cavity are yet to be determined. Ayala-Grosso et al. reported that patient-derived olfactory stromal cells from biopsies of olfactory mucosa expressed $A\beta$ peptides [23]. Our previous study using a rat model showed that $A\beta$ peptides injected in the brain were transported from the brain into the nasal cavity [16]. The present study showed that the p-tau/t-tau ratio in the nasal cavity was significantly higher in AD cases than in controls, in the olfactory cleft and middle nasal meatus. Interestingly, the olfactory cleft and middle nasal meatus are deeper in the nasal cavity where the biomarker levels are closer to those of the olfactory epithelium and cranial cavity. Although the sources of the detected biomarkers in the nasal cavity are yet to be determined, biomarkers in nasal smears may be a useful tool for diagnosing AD.

Conclusion

Our novel PGD-II method detected $A\beta_{42}$, $A\beta_{40}$, p-tau, and t-tau from nasal smears, but levels of $A\beta_{40}$ in all nasal locations of both groups were near zero. This method may be useful in distinguishing between AD patients and cognitively healthy individuals by assessing the p-tau/t-tau ratio in the olfactory cleft and middle nasal meatus.

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SUPPLEMENTARY MATERIAL

The supplementary material is available in the electronic version of this article: <http://dx.doi.org/10.3233/JAD-170962>.

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