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Simultaneous HPLC–MS–MS quantification of 8-iso-PGF_{2 α} and 8,12-iso-iPF_{2 α} in CSF and brain tissue samples with on-line cleanup

Magdalena Korecka^a, Christopher M. Clark^b, Virginia M.-Y. Lee^{b,c}, John Q. Trojanowski^{b,c}, Leslie M. Shaw^{a,*}

^a Department Pathology and Laboratory Medicine, University of Pennsylvania, 3400 Spruce Street, Philadelphia, PA 19104, USA

^b Center for Neurodegenerative Disease Research, University of Pennsylvania, 3600 Spruce Street, Philadelphia, PA 19104, USA

^c Institute on Aging, University of Pennsylvania, 3600 Spruce Street, Philadelphia, PA 19104, USA

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ABSTRACT

Quantitation of isoprostanes such as 8-iso-PGF_{2 α} and 8,12-iso-iPF_{2 α}-VI in biological fluids has been proposed as a reliable test of oxidant stress and inflammation in a variety of disorders. This paper presents a liquid chromatography method with tandem mass spectrometry detection for the simultaneous analysis of these two isoprostanes in human CSF and brain tissue samples. An API 5000 triple quadrupole instrument (AB Sciex, Foster City, CA, USA) with an APCI ion source was used in this study. Aliquots of CSF samples (0.25 mL) were treated with a methanol:zinc sulfate mixture followed by on-line cleanup on an extraction column (Validated-C18) with 0.1% formic acid. The brain tissue samples were homogenized and lipids were extracted using Folch solution. Solid-phase extraction columns (C_{18}) were used for the purification of the brain isoprostane fraction. Chromatographic separation was achieved using an analytical column (Synergi C₁₈ HydroRP) with 0.1% formic acid in water and a mixture of methanol: acetonitrile under isocratic conditions. The mass spectrometer was operated in the MRM scan and negative ion mode. The quadrupoles were set to detect the molecular ions $[M-H]^-$ and high mass fragments of isoprostanes: m/z 353 \rightarrow 193 amu (8-iso-PGF_{2 α}) and m/z 353 \rightarrow 115 amu (8,12-iso-iPF_{2 α}-VI) and their deuterated internal standards: m/z 357 \rightarrow 197 amu (8-iso-PGF₂ α -d₄) and m/z 364 \rightarrow 115 amu (8,12-iso-iPF₂ α -VI-d₁₁). The lower limit of quantification was 2.5 pg/mL for 8-iso-PGF_{2 α} and 5.0 pg/mL for 8,12-iso-PF_{2 α}-VI for the CSF method and 10.0 pg/0.1 g of tissue and 30.0 pg/0.1 g of tissue for 8-iso-PGF_{2 α} and 8,12-iso-iPF_{2 α}-VI, respectively, for the brain tissue method. No ion suppression or enhancement of the detection of 8-isoPGF_{2 α}, 8,12-isoPF_{2 α}-VI or both internal standards was found.

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1. Introduction

 F_2 -isoprostanes are prostaglandin (PG)-like compounds produced primarily from esterified arachidonic acid, a polyunsaturated fatty acid that is highly oxidizable and abundant in the brain, by non-enzymatic reactions catalyzed by free radicals. F_2 isoprostanes are a group of 64 compounds isomeric in structure to $F_{2\alpha}$ prostaglandin. It has been proposed that measurement of F_2 isoprostanes is the most reliable approach to assess oxidative stress

* Corresponding author. Tel.: +1 215 662 6575; fax: +1 215 662 7529. *E-mail addresses:* korecka@mail.med.upenn.edu (M. Korecka), status *in vivo*, providing an important tool to explore the role of oxidative stress in the pathogenesis of neurological, cardiovascular, pulmonary, renal and liver diseases [1–4].

F₂-isoprostanes have been measured in body fluids such as urine, blood, bile, cerebrospinal fluid (CSF) and lung condensate, as well as in brain and liver tissues [1,5,6]. Currently, gas chromatography (GC)–mass spectrometry [7–15] or liquid chromatography (HPLC)–mass spectrometry [6,16–22] are the most accurate methods for identifying and quantifying isoprostanes in biological fluids. However sample preparation is generally a labor-intensive, time-consuming procedure requiring purification by solid-phase extraction with [12] or without derivatization [7,8,10,11,13,14,16–18,21,23], or liquid–liquid extraction [20] sometimes followed by derivatization [9,19]. For the majority of GC methods SPE is combined with thin layer chromatography (TLC) in order to obtain highly purified samples [8,9,12–15].

Abbreviations: ME, matrix effect; RE, recovery.

trojanow@mail.med.upenn.edu (J.Q. Trojanowski), shawlmj@mail.med.upenn.edu, Les.Shaw@uphs.upenn.edu (L.M. Shaw).

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In this paper we describe a reliable, accurate and sensitive method together with a very simple and rapid sample preparation procedure for the simultaneous quantification of 8-iso-PGF_{2 α} and 8,12-iso-iPF_{2 α}-VI in human CSF and brain tissues. We also present here a new procedure for preparing biological fluids free of endogenous isoprostanes for the preparation of standards and quality control samples and a modified method for measurement of matrix effect that is developed specifically for endogenous compound analysis. The potential applicability of this new method for detection of oxidative stress in brain tissue was demonstrated in a pilot study where both isoprostanes were measured in brain tissue samples collected from rats to test the hypothesis that oxidative stress is activated after seizures. The finding that 8-iso-PGF_{2 α} concentration in rat hippocampus of animals with induced status epilepticus was significantly higher when compared with control animals [24] supports this hypothesis and provided initial experience in the applicability of this new analysis procedure.

2. Experimental procedures

2.1. Materials

Two isoprostanes 8-iso-PGF_{2 α} and 8,12-iso-iPF_{2 α}-VI and their internal standards (IS), tetradeuterated 8-iso-PGF_{2 α} and 8,12-iso $iPF_{2\alpha}$ -VI-d₁₁ were obtained from Cayman Chemical Co (Ann Arbor, MI, USA). Acetonitrile, methanol, ethanol, heptane, chloroform and ethyl acetate, all HPLC purity grade, zinc sulfate, sodium chloride, potassium hydroxide and hydrochloric acid, all A.C.S. purity grade, were from Fisher Scientific (Pittsburg, PA, USA). Formic acid and 3,5-di-tert-butyl-4-hydroxytoluene (BHT) were purchased from Sigma-Aldrich (Milwaukee, WI, USA). Solid-phase extraction columns (SPE C18, 200 mg, 4 mL) were obtained from Alltech Associates Inc. (Deerfield, IL, USA). All solutions were prepared with type one distilled water (Mar Cor Medical System; Harleysville, PA, USA). Artificial CSF (Na, 150 mM; K, 3.0 mM; Ca, 1.4 mM; Mg, 0.8 mM; P, 1.0 mM; Cl, 155 mM) was purchased from Harvard Apparatus (Holliston, MA, USA). All working solutions of 8-iso-PGF_{2 α} and 8,12-iso-iPF_{2 α}-VI for standards and quality control (QC) samples preparation, and working solutions of two ISs were prepared by diluting stock solutions of isoprostanes (1 mg/mL, each) and ISs (0.1 mg/mL, each) with ethanol for CSF analysis and with methanol for brain tissue analysis since methanol was used in the lipid hydrolysis procedure for the latter tissue. The spiking solutions for the calibration standard and quality control samples contained both 8-iso-PGF_{2 α} and 8,12-iso-iPF_{2 α}-VI. Internal standard spiking solution also contained the two deuterated internal standards in equal concentrations (1 ng/mL of ethanol for CSF analysis and 100 ng/mL of methanol for brain tissue analysis).

All calibrators and quality control samples for quantification of isoprostanes in CSF were prepared in artificial CSF, which we validated as a suitable matrix for these samples. The most important limitation for using human CSF as a matrix for standards and QC samples preparation is lack of sufficient volume of this fluid. Additionally, human CSF must be pre-treated to remove endogenous isoprostanes to be used as a biological matrix for standard and QC sample preparation. Thus using artificial CSF for this purpose makes the procedure much simpler and shorter.

All calibrators and quality control samples for quantification of isoprostanes in brain tissues were prepared by spiking brain homogenates, previously treated with SPE columns to remove endogenous isoprostanes, with appropriate concentrations of isoprostanes and the respective internal standards.

All CSF samples were obtained by lumbar puncture after informed consent from subjects with a variety of late-life neurodegenerative dementing disorders and cognitively normal elderly undergoing clinical evaluations in the Penn Memory Center. The samples were aliquoted without centrifugation into polypropylene cryovials and stored at -80 °C.

Post-mortem frontal lobe brain specimens were obtained from patients who came to autopsy at the Center for Neurodegenerative Disease Research at the University of Pennsylvania following informed consent from the decedent's family. The samples were stored at -80 °C, and only gray matter was used for isoprostane analysis.

2.2. Sample preparation

2.2.1. CSF analysis

CSF sample preparation was limited to a protein precipitation step, achieved by mixing the sample aliquot with a combination of methanol and 0.1 M zinc sulfate (7:3, v/v) followed by a centrifugation step [6]. Six standards that contained both isoprostanes with concentrations ranging from 2.5 to 80 pg/mL (8-iso-PGF_{2 α}) and from 5.0 to 250 pg/mL (8,12-iso-iPF_{2 α}-VI) together with blank sample were prepared on the day of analysis by spiking artificial CSF (0.25 mL) with an aliquot of standard and internal standard spiking solutions (0.05 mL, each). Quality control sample aliquots (8-iso-PGF_{2α}: 10, 25 and 50 pg/mL; 8,12-iso-iPF_{2α}-VI: 15, 30 and 100 pg/mL) were prepared in bulk by spiking artificial CSF, and then stored in 1.5 mL Eppendorf tubes at -80 °C until the day of analysis. On the day of analysis the quality control sample aliquots and patient CSF sample aliquots stored at -80°C, were thawed at +4°C and then 0.25 mL of CSF and QC samples were spiked with 0.05 mL of ethanol and 0.05 mL of internal standards spiking solution. Protein-free supernatant was prepared by adding 0.35 mL of the precipitation mixture to the standards, QC, and subjects' CSF samples followed by mixing for 10s and then two consecutive centrifugation steps at $9500 \times g$ (Abbot centrifuge, model 3531) for 15 min. Aliquots of supernatant (370 µL) were injected into the HPLC-MS/MS system.

2.2.2. Brain tissue analysis

Preparation of brain tissue gray matter for analysis of isoprostanes includes three primary steps: lipid extraction, followed by hydrolysis that releases total isoprostanes, and their purification. The extraction and hydrolysis procedure followed the method described by Morrow and Roberts [25]. Briefly, brain tissue samples were homogenized with a blade homogenizer and lipids were extracted using Folch solution (chloroform:methanol, 2:1, v/v) containing 0.005% BHT to prevent auto-oxidation for both procedures. Following evaporation to dryness under nitrogen, lipids were hydrolyzed with KOH (15%) to release $F_{2\alpha}$ -isoprostanes. Solidphase extraction columns (C_{18}) were next used for the purification of the isoprostane fraction which was subsequently analyzed by HPLC with tandem mass spectrometry detection.

In the purification process 2 mL of homogenate was spiked with 0.08 mL of methanol and 0.06 mL of the spiking solution of internal standards and loaded onto pre-conditioned (with 6 mL of methanol, followed by 6 mL of water) C_{18} solid-phase extraction columns. Next the columns were washed with 9 mL of water (pH 3.0) and 9 mL of heptane. Isoprostanes were then eluted with 9 mL of a mixture of ethyl acetate and methanol (1:1, v/v). The eluent from the C_{18} SPE columns was then evaporated to dryness under a stream of nitrogen at 37 °C and the isoprostanes were re-dissolved in 0.5 mL of ethanol and analyzed using the HPLC-tandem mass spectrometry system.

The calibration curve for brain tissue sample analysis included blank sample and five calibrators with the concentration range from 10 to 100 pg/0.1 g of tissue for 8-iso-PGF_{2 α} and from 30 to 200 pg/0.1 g of tissue for 8,12-iso-iPF_{2 α}-VI. Three QC samples (8iso-PGF_{2 α}: 25, 40 and 80 pg/0.1 g of tissue; 8,12-iso-iPF_{2 α}: 30, 80 and 160 pg/0.1 g of tissue) were included into each run. Brain tissue homogenate free of endogenous isoprostanes, was prepared by running 2 mL of the mixture of homogenate through C_{18} SPE columns, previously pre-conditioned with methanol and water, 6 mL of each. The homogenate collected from the columns, free of the endogenous isoprostanes which were adsorbed onto the columns, was used as the standard and OC samples' matrix. For the preparation of standards 2 mL of the matrix was spiked with 0.08 mL of spiking solutions of standards and 0.06 mL of spiking solution of internal standards and loaded on pre-equilibrated SPE columns, in the same way like CSF from patient, for purification process. Quality control samples were prepared in bulk by spiking the matrix, free of endogenous isoprostanes, with QC sample spiking solutions. Then the aliquots of QC samples were stored in 1.5 mL polypropylene Eppendorf tubes at -80 °C. On the day of analysis the quality control samples were thawed at +4 °C and 2 mL of QC was spiked with 0.08 mL of methanol and 0.06 mL of internal standards spiking solution and loaded onto pre-conditioned C₁₈ solid-phase extraction columns. The next steps of standard and QC sample preparation are the same as for the individual sample preparation.

2.3. HPLC-tandem mass spectrometer and conditions

Analyses of 8-iso-PGF_{2α} and 8,12-iso-iPF_{2α}-VI were carried out on an API 5000TM triple quadrupole mass spectrometer equipped with a QJetTM ion guide and accelerated by a LINAC[®] collision cell (AB Sciex, Foster City, CA, USA) with an atmospheric pressure chemical ionization probe in a Turbo VTM ion source, interfaced with an Agilent 1100 liquid chromatography system including an autosampler G1313A, isocratic pump G1310A, binary pump G1312A and degasser G1379A. Analyst 1.4.2 software (AB Sciex, Foster City, CA, USA) was used for system control and data processing. The mass spectrometer was operated in the negative ion mode and MRM scan type. The first and third quadrupoles were set to detect the negatively charged molecular ions $[M-H]^-$ and a high mass fragment of 8-iso-PGF_{2α} (m/z 353 \rightarrow 193 amu) and 8,12-isoiPF_{2α}-VI (m/z 353 \rightarrow 115 amu) and their ISs (m/z 357 \rightarrow 197 amu (8-iso-PGF_{2 α}-d₄) and m/z 364 \rightarrow 115 amu (8,12-iso-iPF_{2 α}-VI-d₁₁). The fragmentation process was described previously [16,21].

CSF sample cleanup, and the additional sample cleanup for homogenate samples, was achieved on-line with 0.1% formic acid in water (extraction solution) at a flow rate of 1.5 mL/min using an extraction column (Validated-C₁₈, 5 μ m; PerkinElmer, MA, USA). After 1.8 min of washing the switching valve was activated and the sample extract was back flushed from the extraction column onto the analytical column (Synergi C₁₈ 4 μ m Hydro-RP 250 mm × 3 mm, Phenomenex, USA) maintained at 40 °C (Fig. 1). The mobile phase consisted of 0.1% formic acid in water (solvent A) and a mixture of acetonitrile: methanol (20:80, v/v, solvent B) (solvent A:solvent B-40:60, v/v) adjusted to pH 6.15 with ammonium hydroxide. The flow rate of the mobile phase was 0.6 mL/min and total run time was 17 min.

2.4. Assay validation

Validation of our method followed FDA Guidance for Industry–Bioanalytical Method Validation, May 2001 [26] and was done using standards, QC samples, human CSF and homogenate from post-mortem human brain tissues.

2.4.1. LLOQ and linearity

The lower limit of quantification (LLOQ) was defined as the lowest concentration of 8-iso-PGF_{2α} and 8,12-iso-iPF_{2α}-VI that could be measured with a reproducibility of \leq 20% and an accuracy of 80–120% [26]. The linearity of the isoprostanes assay was assessed over the concentration range 2.5–300 pg/mL (8-iso-PGF_{2α}) and 5–1000 pg/mL (8,12-iso-PGF_{2α}-VI) for CSF analysis and over the concentration range 10–300 pg/0.1 g of tissue (8-iso-PGF_{2α}) and 30–1000 pg/0.1 g of tissue (8,12-iso-iPF_{2α}-VI) using a linear through zero non-weighted regression model.

2.4.2. Between- and within-day precision and accuracy

Between-day precision and accuracy was evaluated for all standards and QC samples for CSF and brain tissue analyses. The within-



Fig. 1. Switching valve set-up for simultaneous analysis of 8-iso-PGF_{2α} and 8,12-iso-iPF_{2α}-VI: (A) position for on-line cleanup and (B) position for eluting analyte from the extraction column to the analytical one and then to the mass spectrometer.

day precision and accuracy was evaluated for three CSF QC samples. All data were collected from a replication experiment in which each sample was analyzed multiple times (10–20) on the same day and in duplicate on different days (5 days). Mean, SD and CV (%) were calculated for the results from the same and different days.

2.4.3. Absolute recovery and matrix effect

The possibility of a matrix effect (ME) on detection of the two isoprostanes and their internal standards, and absolute recovery (RE) of these four compounds in human CSF and brain tissue homogenate were assessed based on the procedure established by Matuszewski et al. [27] using a mixture of 5 human CSF samples from different patients (to obtain necessary volume) or a mixture of 5 different brain homogenates (to obtain necessary volume) as a biological matrix. The original procedure was modified since we were working with endogenous compounds that are always present in biological material, so in the ME/RE assessment experiment we had to deal with both endogenous and exogenous isoprostanes (see the procedure below and Fig. 2). For correct calculation of ME on detection of isoprostanes it was necessary to distinguish exogenous from endogenous isoprostanes. Briefly, three sets of three samples obtained by spiking the CSF or homogenate mixture with standard spiking solutions ranged from 5 to 80 pg/mL (CSF) or 30-70 pg/0.1 g of tissue (homogenate)(8iso-PGF_{2\alpha}) and from 12.5 to 250 pg/mL (CSF) or 60–150 pg/0.1 g of tissue (homogenate) (8,12-iso-iPF_{2 α}-VI) and the mixture of both internal standards were prepared. Set 1, without biological matrix, contained standards (exogenous isoprostanes) with internal standards in the extraction solution. We used the extraction solution (0.1% formic acid in water) as a solvent for set 1, instead of mobile phase suggested by Matuszewski, since our procedure includes a switching valve technique and 2-dimensional chromatography that

incorporates as a first step the binding of the compounds on the extraction column. Thus, addition of mobile phase to set 1 would change this binding. Set 2 contained CSF or homogenate extract spiked with standards and internal standards spiking solutions after protein precipitation (CSF) or cleanup on C₁₈ SPE columns (brain homogenate). This set contained both exo- and endogenous isoprostanes. Additionally, an endogenous control (exogenous blank) sample was included into set 2 since this set was subsequently compared with set 1 that contained only exogenous isoprostanes. This sample contained besides endogenous isoprostanes, and the two internal standards, ethanol/methanol instead of the standard spiking solution. Set 3 contained regular samples i.e. CSF or homogenate spiked with standards and internal standards spiking solutions before protein precipitation (CSF) or cleaning on C₁₈ SPE columns (homogenate). Set 3 contained exo- and endogenous isoprostanes, similar to set 2, permitting direct comparison of set 3 with set 2. This experiment was run on 2 separate days; all samples were run in triplicate. The peak area for standards and internal standards in these 3 sets was used for the calculations of ME and RE for standards and internal standards according to the following equations:for standards.

ME (%) =
$$\frac{\text{set2} - \text{endogenous control of set2}}{\text{set1}} \times 100$$

RE (%) = $\frac{\text{set3}}{\text{set2}} \times 100$

ME (%) =
$$\frac{\text{set2}}{\text{set1}} \times 100$$

$$RE (\%) = \frac{set3}{set2} \times 100$$



Fig. 2. Sample preparation for matrix effect and absolute recovery assessment.

where set 1—peak area of standard (exogenous isoprostanes) or internal standard from set 1, set 2—peak area of standard (exo-and endogenous isoprostanes) or internal standard from set 2, endogenous control of set 2—peak area of endogenous isoprostanes in endogenous control sample from set 2, set 3—peak area of standard (exo- and endogenous isoprostanes) or internal standard from set 3.

To determine the ME for artificial CSF and homogenate extract, without endogenous isoprostanes, on their detection, we used a post column infusion technique [28,29]. An infusion pump was used to deliver a constant flow (10 μ L/min) of isoprostanes standard contained 50 pg/mL of 8-iso-PGF_{2 $\alpha}$ and 8,12-iso-iPF_{2 α}-VI each, to HPLC eluent. An artificial CSF sample, or homogenate extract treated to remove endogenous isoprostanes, was injected under the conditions of our method and the response from the infused analyte was recorded.}

2.4.4. Analytical recovery of analytes from human CSF and brain tissue homogenate

In addition the analytical recovery of isoprostanes from human CSF and brain tissue homogenates was tested in a spiking experiment. Ten human CSFs and brain tissue homogenate were spiked separately with three different spiking solutions that contained both 8-iso-PGF_{2α} and 8,12-iso-iPF_{2α}-VI (each at concentrations of: 12.5, 25 and 50 pg/mL) for CSF analysis or one spiking solution that contained both 8-iso-PGF_{2α} (40 pg/0.1 g of tissue) and 8,12-iso-iPF_{2α}-VI (80 pg/0.1 g of tissue) for brain tissue analysis. After preparation (protein precipitation for CSF and C₁₈ SPE column cleanup for brain tissue analysis) the samples before and after spiking were run using our method. Analytical recovery of the isoprostanes was expressed as the percentage of the nominal spiked-in concentrations. The final results were calculated as the mean, SD and coefficient of variation.

2.4.5. Carry over

Sample carryover was checked by running blank samples of CSF and brain tissue homogenate after the highest standard and

3. Results

3.1. Linearity and LLOQ

A representative mass chromatogram of 8-iso-PGF_{2α} and 8,12iso-iPF_{2α}-VI together with their internal standard in a sample of CSF is shown in Fig. 3. The retention times of 8-iso-PGF_{2α} and of 8,12-iso-iPF_{2α}-VI both in CSF and brain tissue sample, were 8.62 ± 0.2 min and 15.56 ± 0.4 min, respectively (n = 13 days). The assay was linear for 8-iso-PGF_{2α} over the range 2.5–300 pg/mL (CSF; y = 0.0267(±0.0034)x, r = 0.9997) or 10–300 pg/0.1 g of tissue (brain tissue; y = 0.0032(±0.0004)x, r = 0.9989, n = 10), and for 8,12-iso-iPF_{2α}-VI over the range 5–1000 pg/mL (CSF; y = 0.0126(±0.0008)x, r = 0.9995) or 30–1000 pg/0.1 g of tissue (brain tissue; y = 0.0022(±0.0002)x, r = 0.9985, n = 10). The LLOQ was 2.5 pg/mL and 5.0 pg/mL for 8-iso-PGF_{2α} and 8,12-iso-iPF_{2α}-VI vI respectively, for the CSF method, and 10.0 pg/0.1 g of tissue and 30.0 pg/0.1 g of tissue for 8-iso-PGF_{2α} and 8,12-iso-iPF_{2α}-VI respectively, for the brain tissue method.

3.2. Between- and within-day precision and accuracy for standards and quality control samples

3.2.1. Standards

Between-day precision (CV) for 8-iso-PGF_{2 α} standards ranged from 1.8% to 8.5% (CSF) and from 2.4% to 8.9% (brain tissue). For 8,12-iso-iPF_{2 α}-VI standards between-day precision ranged from 1.1% to 10.6% (CSF) and from 1.5% to 10.2% (brain tissue) (*n*=5 days). Between-day accuracy for 8-iso-PGF_{2 α} standards ranged



Fig. 3. Mass chromatogram of endogenous 8-iso-PGF_{2 α} (A) (5.61 pg/mL) and 8,12-iso-iPF_{2 α}-VI (C) (16.0 pg/mL) detected in human CSF. Additional isomers and prostaglandin peak identified using commercial available standards are labeled. Each internal standard (B and D) concentration is 100 pg/mL CSF.

Table 1

Between days (n = 5) precision and accuracy for 8-iso-PGF_{2 α} and 8,12-iso-iPF_{2 α}-VI standards and quality control samples for CSF and brain tissues analysis.

		Concentration	Found amount ± SD (pg/mL)	CV (%)	Accuracy (%)
CSF	Standards 8-iso-PGF _{2α}	2.5	2.73 ± 0.2	7.42	109.3 ± 7.9
	(pg/mL)	5.0	5.45 ± 0.5	8.51	109.0 ± 8.9
		10.0	9.69 ± 0.4	3.76	96.9 ± 3.6
		25.0	25.4 ± 0.9	3.51	101.5 ± 3.5
		50.0	51.3 ± 2.0	3.94	102.6 ± 4.2
		80.0	79.1 ± 1.4	1.76	99.0 ± 1.8
	QC samples 8-iso-PGF _{2α}	10.0	10.5 ± 0.8	7.78	105.1 ± 8.2
	(pg/mL)	25.0	26.6 ± 1.1	4.23	106.4 ± 4.4
		50.0	54.1 ± 2.3	4.25	108.1 ± 4.6
	Standards 8,12-iso-iPF $_{2\alpha}$ -VI	5.0	4.97 ± 0.5	10.60	99.2 ± 10.4
	(pg/mL)	12.5	13.0 ± 0.6	4.36	104.5 ± 4.4
		25.0	24.4 ± 0.8	3.23	97.6 ± 2.9
		50.0	47.3 ± 3.9	8.34	94.5 ± 7.8
		125.0	120 ± 4.6	3.82	95.8 ± 3.7
		250.0	253 ± 2.9	1.14	101.5 ± 1.3
	QC samples 8,12-iso-iPF _{2α} -VI	15.0	15.9 ± 1.3	8.10	105.7 ± 8.4
	(pg/mL)	30.0	30.4 ± 1.5	4.80	101.3 ± 4.9
		100.0	93.8 ± 3.3	3.50	93.8 ± 3.3
Brain tissue	Standards 8-iso-PGF $_{2\alpha}$	10.0	10.4 ± 0.9	8.90	104.4 ± 9.3
	(pg/0.1 g of tissue)	30.0	29.7 ± 1.0	3.60	99.1 ± 3.6
		50.0	50.1 ± 1.4	2.89	100.3 ± 2.9
		70.0	64.7 ± 3.8	5.80	92.3 ± 5.4
		100.0	101.4 ± 2.5	2.44	101.4 ± 2.5
	QC samples 8-iso-PGF _{2α}	25.0	26.6 ± 2.2	8.20	106.5 ± 8.8
	(pg/0.1 g of tissue)	40.0	41.8 ± 1.2	2.80	104.4 ± 3.0
		80.0	89.0 ± 4.5	5.10	111.3 ± 5.7
	Standards 8,12-iso-iPF $_{2\alpha}$ -VI	30.0	32.5 ± 3.3	10.20	108.3 ± 11.1
	(pg/0.1 g of tissue)	60.0	61.2 ± 2.6	4.24	101.9 ± 4.3
		100.0	98.0 ± 5.8	5.95	98.0 ± 5.8
		150.0	151.3 ± 4.9	3.33	100.7 ± 3.4
		200.0	199.7 ± 2.3	1.53	99.7 ± 1.5
	QC samples 8,12-iso-iPF _{2α} -VI	30.0	31.4 ± 2.8	8.90	104.6 ± 9.3
	(pg/0.1 g of tissue)	80.0	85.7 ± 9.2	9.60	107.1 ± 11.5
		160.0	171.0 ± 17.6	10.30	106.9 ± 11.0

from 96.9% to 109.3% (CSF) and from 92.3% to 104.4% (brain tissue) and for 8,12-iso-iPF_{2α}-VI standards from 94.5% to 104.5% (CSF) and 98.0% to 108.3% (brain tissue). See Table 1 for more details.

3.2.2. Quality control samples

Between-day (n = 5 days) precision for three CSF QC samples of 8-iso-PGF_{2 α} were 7.8%, 4.2% and 4.3%, with accuracy range 105.1–108.1% and for 8,12-iso-iPF_{2 α}-VI were 8.1%, 4.8% and 3.5%, with accuracy range 93.8–105.7%. Between-day (n = 5 days) precision for three brain tissue QC samples of 8-iso-PGF_{2 α} were 8.2%, 2.8% and 5.1%, with accuracy range 104.4–111.3% and of 8,12-iso-iPF_{2 α}-VI were 8.9%, 9.6% and 10.3%, with accuracy range 104.6–107.1%. See Table 1 for more details. Within-day precision (n = 10–20) for CSF 8-iso-PGF_{2 α} QC samples was lower than 7% with a mean accuracy of 103.6 ± 5.3%, and for 8,12-iso-iPF_{2 α}-VI QC samples was lower than 5% with a mean accuracy of 98.3 ± 3.2%.

3.3. Absolute recovery and matrix effect

The mean absolute RE of 8-iso-PGF_{2α} assessed for the concentration range 5–80 pg/mL (CSF) or 30–70 pg/0.1 g of tissue (brain tissue) was $103.2 \pm 9.0\%$ (CV=0.1%) or $98.8 \pm 13.3\%$ (CV=6.1%), respectively, and of 8,12-iso-iPF_{2α}-VI for the concentration range 12.5-250 pg/mL (CSF) or 60-150 pg/0.1 g of tissue (brain tissue) was 98.6 ± 6.7 pg/mL (CV=0.1%) or $111.0 \pm 11.6\%$ (CV=10.5%), correspondingly (Table 2). Minimal, clinically not significant ME was found for the detection of 8-iso-PGF_{2α} (-6%, CV=0.1% for CSF and -2%, CV=6.6% for brain tissue), and very small ion enhancement for the detection of 8,12-iso-iPF_{2α}-VI in CSF (+1.1%, CV=0.1%). For brain tissue, we found very small, but insignificant ion suppression

for detection of 8,12-iso-iPF_{2α}-VI (-2%, CV = 7.4%). Data for the RE and ME experiment for both internal standards are summarized in Table 2. Lack of ME using artificial CSF for detection of 8-iso-PGF_{2α} and 8,12-iso-iPF_{2α}-VI is shown in Fig. 4. There is also no ME for brain tissue homogenate, previously run through the SPE columns, on detection of 8-iso-PGF_{2α} and 8,12-iso-iPF_{2α}-VI (data not shown).

3.4. Analytical recovery of analytes from human CSF and brain tissue homogenate

Results of the analytical recovery of 8-iso-PGF_{2 α} and 8,12-iso-iPF_{2 α}-VI after spiking human CSF with three different spiking solutions of the two isoprostanes in each solution are presented in Table 3. Mean analytical recoveries of 8-iso-PGF_{2 α} and 8,12-iso-iPF_{2 α}-VI from CSF were respectively, from 102 ± 5.3% to 107 ± 6.9% and 95.6 ± 3.1% to 103.1 ± 5.3%. The respective analytical recoveries from brain tissue homogenates were 100.9 ± 12.9% and 103.5 ± 7.2% for the two isomers.

3.5. Carry over

No carryover effect was observed. There were no detectable peaks of 8-iso-PGF_{2 α} or 8,12-iso-iPF_{2 α}-VI in the chromatogram of blank samples of artificial CSF or CSF pre-treated to remove endogenous isoprostanes and brain tissue homogenate (pre-treated to remove endogenous isoprostanes) run immediately after the highest standard. Accuracy of the lowest standard run after the highest one was 98.2 ± 3.9% (CV = 4.0%, *n* = 10) for 8-iso-PGF_{2 α} and 101.9 ± 2.9% (CV = 2.9%, *n* = 10) for 8,12-iso-iPF_{2 α}-VI (data for artificial CSF analysis).

Table 2 ME (%) and RE (%) ($n = 18$) of 8-iso-PGF _{2α} , 8,12-iso-iPF _{2α} -VI and their internal standards for CSF and brain tissue samples.								
	Compound	8-iso-PGF $_{2\alpha}$	8-iso-PGF _{2α} -d ₄	8,12-iso-iPF _{2α} -VI	8,12-iso-iPF _{2α} -VI-d ₁₁			
CSF	Concentration range (pg/mL) ME RE	$5-80 \\ 94.0 \pm 12.5 \\ 103.2 \pm 9.0$	$\begin{array}{c} 100 \\ 105.7 \pm 5.2 \\ 101.3 \pm 3.7 \end{array}$	$\begin{array}{c} 12.5{-}250\\ 101.1{\pm}9.7\\ 98.6{\pm}6.7\end{array}$	$100 \\ 104.3 \pm 4.7 \\ 102.3 \pm 5.7$			
Brain tissue	Concentration range (pg/0.1 g of tissue) ME RE	30-70 98.1 ± 6.5 98.8 ± 13.0	$\begin{array}{c} 1500 \\ 96.4 \pm 15.6 \\ 107.9 \pm 13.2 \end{array}$	60-150 98.1 \pm 7.3 111.0 \pm 11.6	$\begin{array}{c} 1500 \\ 96.8 \pm 4.4 \\ 110.5 \pm 12.7 \end{array}$			

Table 3

Mean analytical recovery (%) of 8-iso-PGF_{2 α} and 8,12-iso-iPF_{2 α}-VI used for spiking human CSF and brain tissue homogenate (*n* = 10).

	CSF recovery of (pg/mL)			Brain tissue recovery of		
8-iso-PGF $_{2\alpha}$		12.5	25.0	50.0	40.0	
	Median	103.5	110.1	100.8	99.0	
	Mean	102.0	107.0	102.5	100.9	
	SD	5.3	6.9	5.8	12.9	
	CV (%)	5.2	6.4	5.7	12.8	
8,12-iso-iPF _{2α} -VI		12.5	25.0	50.0		80.0
	Median	96.7	103.6	95.6		105.0
	Mean	99.7	103.1	95.6		103.5
	SD	6.6	5.3	3.1		7.2
	CV (%)	6.6	5.1	3.3		6.9



Fig. 4. Lack of ME of artificial CSF on detection of 8-iso-PGF_{2 α} (A) and of 8,12-iso $iPF_{2\alpha}$ -VI (B) by the post-column infusion experiment. The experimental conditions are described in Section 2.4.3. The arrows represent the retention time for 8-iso-PGF_{2 α} and 8,12-iso-iPF_{2 α}-VI (A and B, respectively).

4. Discussion

Our goal here was to develop an analytical method that is reliable, accurate and precise with a straightforward simplified sample preparation procedure for quantification of 8-iso-PGF_{2 α} and 8,12iso-iPF_{2\alpha}-VI in CSF and human brain tissue.

Solid-phase extraction columns have been used in sample preparation procedures in all published HPLC-MS methods for the analysis of isoprostanes. The first study to use a protein precipitation step followed by on-line cleanup for analysis of isoprostanes is our report of this procedure for analysis of isoprostanes in human urine and plasma [6]. The method for CSF and brain tissue analysis presented in this paper is based on that.

Our method uses an automated column-switching technique together with on-line sample cleanup that reduces manual sample preparation to a simple protein precipitation step, and eliminates the need for complex purification steps. We used a 10 port switching valve, instead of a simpler six port valve, since this kind of device was available at our laboratory at the time of method development. The connection that we set, one of at least three available for 10 port switching valve, gave satisfactory results based on the data from method validation. For CSF samples, protein precipitation and on-line cleanup were the only steps involved in sample preparation. For brain tissues it was necessary to perform standard procedures for tissue preparation including tissue homogenization with lipid extraction followed by hydrolysis and purification of released isoprostanes on SPE columns. The samples obtained from SPE columns were suitable for injection into the HPLC/MS/MS system without additional on-line cleanup with a column-switching technique. However, since we run CSF and brain tissues on the same HPLC-mass spectrometer system to avoid the additional technical work with disconnecting the switching valve and changing the tubing set-up we adjusted the CSF method for analysis of brain tissues samples as well.

The lowest 8,12-iso-iPF_{2 α}-VI standard for CSF analysis (the LLOQ) is twice as high as the lowest one for the quantification of 8-iso-PGF_{2 α} (5.0 pg/mL vs 2.5 pg/mL) since based on our results the concentration of 8,12-iso-iPF_{2 α}-VI in CSF is about 3.1 ± 0.97 (*n* = 65) times higher compared to the concentration of 8-iso-PGF_{2 α}. Also the lowest standards for the quantification of both isoprostanes in the brain tissues assay are higher compared to the assay for CSF, since we found that the concentration of these isoprostanes in brain tissues is 10-12 times higher compared to their concentration in CSF. Based on the recommendations of Shah et al. [30] and the FDA guideline for analytical performance validation [26] the described assay has excellent between- and within-day analytical recovery and precision for all standards and QC samples for both isoprostanes in the two studied biological matrices: human CSF and post-mortem human brain tissues. The mean analytical recovery was measured not only for standards and QC samples but also for human CSF and brain tissue homogenates because of the differences in the biological matrices (artificial CSF, used for calibrator and QC samples preparation vs human CSF or brain tissue homogenate "cleaned" of endogenous isoprostanes on SPE columns used for calibrator and QC sample preparation vs brain tissue homogenate containing endogenous isoprostanes). This approach provides an assurance that the different matrices used for standard and QC sample preparation do not result in artifactual isoprostane concentrations in human CSF and brain tissue.

The ion suppression/enhancement and absolute recovery were checked for the CSF (artificial and human CSF) and brain tissue matrices. None of these exerted a significant effect on ion detection or recovery. It has already been shown that for small molecules the atmospheric pressure chemical ionization ion source gives less ion suppression when compared to electrospray ionization (ESI) [31–33]. As we previously reported for the urinary isoprostane analysis method, we observed very high ion suppression using ESI [6], for the current study we selected APCI without taking ESI into consideration. Additionally to further reduce potential inaccuracy caused by changing ionization efficiency from even minimal ion suppression or enhancement from sample to sample we used stable isotope-labeled analytes as internal standards (8-iso-PGF_{2 α}-VI-d₁₁).

5. Conclusions

We have described an HPLC-atmospheric pressure chemical ionization-tandem mass spectrometry method with high sensitivity and selectivity for the simultaneous quantification of 8-iso-PGF_{2α} and 8,12-iso-PF_{2α}-VI in human CSF and brain tissue samples. The developed method showed very good performance in terms of accuracy and repeatability. The pre-treatment procedure of CSF is extremely simple and did not involve costly, time and labor-consuming sample preparation step.

Biological fluids free of endogenous isoprostanes prepared according to the procedure developed here are suitable, inexpensive matrices for standards and quality control preparation. This methodology meets the criteria of the United States Food and Drug Administration for analytical method validation and qualification. We are utilizing this procedure for the investigation of oxidative stress in neurodegenerative diseases.

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