



Published in final edited form as:

Steroids. 2015 April ; 96: 140–152. doi:10.1016/j.steroids.2015.01.014.

Ultrasensitive quantification of serum estrogens in postmenopausal women and older men by liquid chromatography-tandem mass spectrometry

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Abstract

An ultrasensitive stable isotope dilution liquid chromatography-tandem mass spectrometry method (LC-MS/MS) was developed and validated for multiplexed quantitative analysis of six unconjugated and conjugated estrogens in human serum. The quantification utilized a new derivatization procedure, which formed analytes as pre-ionized N-methyl pyridinium-3-sulfonyl (NMPS) derivatives. This method required only 0.1 mL of human serum, yet was capable of simultaneously quantifying six estrogens within 20 min. The lower limit of quantitation (LLOQ) for estradiol (E2), 16 α -hydroxy (OH)-E2, 4-methoxy (MeO)-E2 and 2-MeO-E2 was 1 fg on column, and was 10 fg on column for 4-OH-E2 and 2-OH-E2. All analytes demonstrated a linear response from 0.5 to 200 pg/mL (5–2000 pg/mL for 4-OH-E2 and 2-OH-E2). Using this validated method, the estrogen levels in human serum samples from 20 female patients and 20 male patients were analyzed and compared. The levels found for unconjugated serum E2 from postmenopausal women (mean 2.7 pg/mL) were very similar to those obtained by highly sensitive gas chromatography-mass spectrometry (GC-MS) methodology. However, the level obtained in serum from older men (mean 9.5 pg/mL) was lower than has been reported previously by both GC-MS and LC-MS procedures. The total (unconjugated + conjugated) 4-MeO-E2 levels were

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significantly higher in female samples compared with males ($p < 0.05$). The enhanced sensitivity offered by the present method will allow for a more specific analysis of estrogens and their metabolites. Our observations might suggest that the level of total 4-MeO-E2 could be a potential biomarker for breast cancer cases.

Keywords

Estrogen; 17 β -estradiol; breast cancer; endometrial cancer; prostate cancer; stable isotope dilution

1. Introduction

Over the last seven years, gas chromatography (GC)-and liquid chromatography (LC)-mass spectrometry (MS)-based methodology has become increasingly reliable for the quantification of extremely low concentrations of unconjugated estrogens present in the serum and plasma of postmenopausal women (Table 1) [1–3]. Estrone (E1), 17 β -estradiol (E2) and 16 α -hydroxy (OH)-E2 are the major unconjugated estrogens that are present in serum (Fig. 1). They are largely bound to protein in the circulation, and their free concentrations are estimated from the steroid hormone binding globulin levels [4–6]. Reliable assays for unconjugated circulating estrogens are important because even though their concentrations are extremely low, increased levels are thought to be an important risk factor for breast [7–10] and endometrial cancer [11–13]. Increased circulating estrogens are also a potential risk factor for prostate cancer in men [14–16], although no systematic studies have been conducted to date to test this possibility. Unfortunately, serum concentrations for many of the unconjugated estrogens in postmenopausal women are close to the reported lower limits of quantification (LLOQs), raising concerns that the values could represent an over-estimation of the true values (Table 1). This means that the corresponding sulfate and β -glucuronide conjugates (Fig. 1), which are present in higher concentrations (Table 2), might have some predictive value in determining cancer risk [17–19]. Furthermore, the sulfate conjugates (Fig. 1) could be a potential source of their corresponding unconjugated forms through action of tissue sulfatases [19].

The very low concentrations of circulating estrogens and their metabolites mean that analysis by MS-based methodology is very challenging. Importantly, LC-MS can overcome potential problems of cross-reactivity that usually occur with more sensitive but less specific immunoassay-based methodology [20–22]. Furthermore, MS-based methods make it possible to quantify multiple estrogens in a single analytical run, which allows for more comprehensive analyses to be conducted. Conventional positive ion GC-MS and LC-MS generally have inadequate sensitivity for routine analyses of endogenous estrogens in the serum of postmenopausal women and older men. Therefore, numerous derivatization methods have been developed to improve the sensitivity of both [1]. High sensitivity GC-MS procedures usually employ electron capture negative chemical ionization using pentafluorobenzoyl (PFBO) derivatives [12,23–28] (Table 1). The corresponding LC-MS method known as electron capture atmospheric pressure chemical ionization, which uses the pentafluorobenzyl (PFB) derivative [29], is widely used for the quantification of lipids [30] but has found only limited utility for the analysis of estrogens and their metabolites [1].

A second LC-MS approach, which has been used much more widely, involves the use of estrogen derivatives that enhance the electrospray ionization (ESI) signal and therefore improves overall sensitivity during LC-ESI/MS analysis. This approach is exemplified by derivatization of the estrogen phenolic moiety to a dansyl ester [31–34]. The dansyl derivative has been used in a number of studies to quantify unconjugated estrogens [35–39] (Table 1) and conjugated estrogens after hydrolysis with β -glucuronidase/arylsulfatase [35,38–40] (Table 2) in serum samples from postmenopausal women. Alternative approaches to improve ESI signal have included the use of picolinoyl [41] and pyridine-3-sulfonyl [42] derivatives. A third LC-MS approach involves the preparation of pre-ionized (quaternized) derivatives, so that protonation of the estrogen derivative is not required. Therefore, suppression of ionization in the ESI source of the mass spectrometer is minimized. This approach has been reported in studies that utilized the N-methyl-2-pyridyl [43], N-methyl-nicotinyl, and 1-(2,4-dinitro-5-fluorophenyl)-4,4-dimethylpiperaziny [44] derivatives attached to the estrogen 3-phenolic moiety. Our group has also used pre-ionized derivatives to improve sensitivity by adding a Girard P (GP) derivative to the 17-oxo moiety of estrone and its metabolites [45] as well as by adding a Girard T (GT) derivative to the 17-oxo-moiety of androgens [46]. However, this approach cannot be employed for the analysis of E2 and its metabolites that lack a 17-oxo-moiety. We now report development of the pre-ionized N-methyl pyridinium-3-sulfonate (NMPS) derivative (Fig. 2), which provides extremely high sensitivity for LC-ESI/MS/MS analysis of E2 and its metabolites. We demonstrate the utility of this new derivatization for the quantification of estrogens in the serum of postmenopausal women and older men.

2. Experimental

2.1. Reagents and Materials

The six estrogens analyzed in this study, E2, 16 α -OH-E2, 4-methoxy (MeO)-E2, 2-MeO-E2, 4-hydroxy (OH)-E2 and 2-OH-E2 were purchased from Steraloids Inc. (Newport, RI). [13,14,15,16,17,18-¹³C₆]-E2 ([¹³C₆]-E2), [2,3,4-¹³C₃]-16 α -OH-E2 ([¹³C₃]-16 α -OH-E2), [13,14,15,16,17,18-¹³C₆]-4-MeO-E2 ([¹³C₆]-4-MeO-E2), [13,14,15,16,17,18-¹³C₆]-2-MeO-E2 ([¹³C₆]-2-MeO-E2), and [13,14,15,16,17,18-¹³C₆]-2-OH-E2 ([¹³C₆]-2-OH-E2) with an isotopic purity of 99% were purchased from Cambridge Isotope Laboratories (Cambridge, MA). [1,4,16,16,17-²H₅]-4-OH-E2 ([²H₅]-4-OH-E2) with an isotopic purity of 98% was obtained from C/D/N Isotopes, Inc. (Pointe-Claire, Quebec, Canada). β -glucuronidase/arylsulfatase (*Helix pomatia*) was obtained from Roche (Indianapolis, IN). E2-3-(β -D-glucuronide)-17-sulfate (E2-3G-17S) was obtained from Sigma-Aldrich (St. Louis, MO). Pyridine-3-sulfonyl chloride (97%) was obtained from Matrix Scientific (Columbia, SC). Dry acetonitrile was purchased from Acros Organic (New Jersey, USA). Methyl-*tert*-butyl-ether (MTBE), iodomethane, methanol, acetone, L-ascorbic acid, formic acid, hydrochloric acid (HCl), sodium chloride, sodium acetate and sodium bicarbonate were obtained from Sigma-Aldrich (Milwaukee, WI). Double charcoal-stripped human serum was obtained from Golden West Biologicals, Inc (Temecula, CA). All solvents used in this study were HPLC Optima grade unless otherwise noted and were purchased from Fisher Scientific (Pittsburgh, PA).

2.2. Clinical study

Twenty postmenopausal women ages 56 to 65 (mean \pm SD, 60.0 \pm 2.66) and twenty men ages 51 to 69 (mean \pm SD, 61.6 \pm 5.41) were recruited for the study. Menstrual and menopausal status of the women was based on self-report. All participants were healthy and the women were not taking exogenous hormones. The blood collection protocol was approved by the University of Pennsylvania Review Board (Protocol # 800924). After the blood was collected, it was allowed to clot for 1 h at room temperature, serum was separated and aliquots were stored at -80°C . Serum samples were allowed to thaw at room temperature and aliquots of 0.1 mL were used for the estrogen analyses.

2.3. Preparation of stock solution and working standard solutions

Estrogen standard and internal standard stock solutions were individually prepared in methanol containing 0.1% (w/v) L-ascorbic acid at a concentration of 1 mg/mL then stored at -20°C . A mixed stock solution of six estrogens or the corresponding internal standards at 1 $\mu\text{g}/\text{mL}$ was prepared by adding 10 μL of each estrogen standard stock solution to a 10 mL volumetric flask with methanol containing 0.1% L-ascorbic acid. Working standard solutions of mixed estrogens and working standard solutions of internal standards (1 ng/mL) were prepared by dilutions of the stock solutions with methanol containing 0.1% (w/v) L-ascorbic acid. Stock solution and working solution of catechols were prepared at concentrations 10 times higher than the other estrogens.

2.4. Preparation of calibration standards and quality controls

Charcoal-stripped human serum was used for preparation of calibration standards and quality controls (QCs). Calibration standards were prepared by spiking appropriate amounts of the working standard solution to charcoal-stripped human serum to make the concentrations of 0.5, 1, 2, 5, 10, 20, 50, 100, 200 pg/mL (10 times higher concentration for 4-OH-E2 and 2-OH-E2). The preparation procedures for QC samples at concentrations of 1.5, 75, and 175 pg/mL were the same as that of the calibration standards. 10 μL of working standard solution of internal standards) was added to each calibration standard and QC sample.

2.5. Sample preparation procedure

The sample preparation procedure was designed to determine unconjugated and total estrogens in serum, including hydrolysis, extraction, derivatization and re-suspension. All serum samples were thawed at room temperature and centrifuged at 5300g at 4°C for 10 min (Z216MK High Capacity Refrigerated Microcentrifuge, Hermel Labortechnik). For the determination of total estrogens, 10 μL of internal standards working solution was spiked into a 0.1 mL aliquot of serum using a calibrated syringe, followed by the addition of 0.1 mL water, 0.1 mL 0.5% L-ascorbic acid, 0.2 mL sodium acetate buffer (200 mM, pH 5.0), and 10 μL of β -glucuronidase/arylsulfatase. Samples were incubated at 37°C for 19 h. After hydrolysis, samples were acidified with 5 μL of 1N HCl followed by addition of 50 μL saturated sodium chloride. Samples underwent liquid-liquid extraction (LLE) with 1.3 mL of MTBE by vortex-mixing for 10 min, followed by centrifugation at 3400g at 4°C for 15 min

(D37520 Osterode Benchtop Centrifuge, Thermo Electron). The top organic layer was transferred to a clean glass tube and evaporated to dryness under nitrogen.

To each dried sample, 100 μ L acetone, 100 μ L sodium bicarbonate (100 mM, pH 9), and 10 μ L of pyridine-3-sulfonyl chloride (10 mg/mL in acetone) were added and vortex mixed for 5 sec. Samples were then incubated in a water bath (Isotemp 210, Fisher Scientific) at 60 °C for 30 min. The derivatives were extracted with 1 mL of MTBE and the organic layer was evaporated to dryness under nitrogen. The N-methyl derivatization reaction was carried out by adding 100 μ L iodomethane (20% in acetonitrile) and followed by incubation at 80 °C for 30 min. The sample was evaporated to dryness under nitrogen. The dry residue was then dissolved in 50 μ L 20% aqueous methanol and 1 μ L was injected for LC-MS/MS analysis. For the analysis of unconjugated estrogens, identical sample preparation procedures were followed with the exclusion of the hydrolysis step. For analysis of PS derivatives the methylation step was not conducted.

2.6. LC-MS/MS

A Vantage TSQ mass spectrometer (Thermo Scientific, San Jose, CA) equipped with a CaptiveSpray™ ion source (Michrom Bioresources, Inc., Auburn, CA) was used for all analysis. The mass spectrometer was interfaced with a Waters nanoAcquity UPLC system (Waters Corporation, Milford, MA) equipped with an autosampler and sample thermo-controller (set at 4°C). Both the UPLC and mass spectrometer were controlled by Xcalibur software (Thermo Scientific). The syringe was washed with water/acetonitrile (95:5, v/v) containing 0.1% formic acid after every injection. Separations were performed using a Waters BEH130 C18 column (150 μ m x 100 mm, 1.7 μ m, 130 Å) (Waters Corporation, Milford, MA) at 50°C using a partial loop injection. Samples were eluted with a linear gradient at a flow rate of 2 μ L/min. For NMPS derivatives, solvent A was water/acetonitrile (99.5:0.5, v/v) containing 0.1% formic acid, and solvent B was acetonitrile/water (98:2, v/v) containing 0.1% formic acid. The gradient started with 30% B, held for 3 min, then linearly increased to 80% B over 17 min. After washing with 90% B for 5 min, the column was re-equilibrated with 30% B for 10 min prior to the next injection. For separation of the PS derivatives, solvent A was water containing 0.1% formic acid, and solvent B was methanol/acetonitrile (40:60, v/v). The gradient started with 50% B, held for 3 min, then linearly increased to 60% B over 30 min. After washing with 80% B for 10 min, the column was re-equilibrated with 50% B for 10 min.

The MS operating conditions were as follows: spray voltage, 1800 V; ion transfer capillary temperature, 300 °C; collision gas, argon at 1.5 mTorr; ion polarity, positive; scan type, selected reaction monitoring (SRM); chrom filter peak width, 15 s; S-lens, 135 v; cycle time, 1 sec; Q1 peak width (FWHM), 0.7 u; Q3 peak width, 0.7 u; DCV, 5 V. The SRM transitions for all the analytes and internal standards are showed in Table 3.

2.7. Method validation

2.7.1. Linearity and lower limit of quantification (LLOQ)—To evaluate linearity of standard curves, calibration standards were prepared at concentrations of 0.5, 1, 2, 5, 10, 20, 50, 100 and 200 pg/mL of each estrogen in charcoal stripped human serum, except for 4-

OH-E2 and 2-OH-E2, which were prepared at 10 times higher concentration. 10 μ L of internal standard working solution was added to each sample. Calibration curves were generated by plotting the area ratios of the analyte to internal standard peak using linear regression with 1/x weighting. The lower limit of quantification (LLOQ) was defined as the lowest calibration level, which could be fitted to the calibration curve with a residual of less than 10% and peak area ratio deviating less than 25%.

2.7.2. Accuracy and precision—Quality control (QC) samples were prepared at the low quality control (LQC; 1.5 pg/mL for E2, 16 α -OH-E2, 4-MeO-E2 and 2-MeO-E2, 15 pg/mL for 4-OH-E2 and 2-OH-E2), medium quality control (MQC; 75 pg/mL for E2, 16 α -OH-E2, 4-MeO-E2 and 2-MeO-E2, 750 pg/mL for 4-OH-E2 and 2-OH-E2) and high quality control (HQC; 175 pg/mL for E2, 16 α -OH-E2, 4-MeO-E2 and 2-MeO-E2, and 1750 pg/mL for 4-OH-E2 and 2-OH-E2) samples. The accuracy and precision were determined on 5 replicate serum samples run on the same day (intra-day) and on 3 different days (inter-day).

2.7.3. Stability—Stability of the NMPS derivatives was assessed by allowing validation samples to sit in the autosampler and re-analyzing after 24 h. Stability of estrogens in the serum was also assessed following three freeze-thaw cycles for the LQC, MQC and HQC.

2.7.4. Recoveries of unconjugated estrogens—The recovery of each unconjugated estrogen was determined at the LQC, MQC and HQC (n=5). One set of QC samples was spiked with internal standard working solution followed by extraction and derivatization procedures described previously. A second batch of QC samples underwent the same preparation protocol, except the internal standard working solution was added after the extraction procedure. Both sets of samples were derivatized and analyzed by consecutive LC-MS analyses. The recoveries of estrogens were calculated by comparing the ratios of analyte/internal standard of both batches.

2.7.5. Recoveries of conjugated E2—The efficiency of enzymatic hydrolysis was determined using β E2-3G-17S standard as the substrate followed by quantification of the liberated E2 by LC-MS/MS. E2-3G-17S (0.037 pmol/mL, equal to 10 pg/mL of unconjugated E2; and 0.37 pg/mL, equal to 100 pg/mL of unconjugated E2) as well as the internal standards were spiked to 0.1 mL of water, then incubated with 0.1 mL of 0.5% (w/v) L-ascorbic acid, 0.2 mL of sodium acetate buffer (200 mM, pH 5.0), and 10 μ L of β -glucuronidase/arylsulfatase at 37 °C for 19 h. After hydrolysis, samples were extracted, derivatized and analyzed by LC-MS/MS. The unconjugated estrogen was quantified from the relevant standard.

2.8. Data Analysis—Serum concentrations of E2 and its metabolites were calculated using Xcalibur software (version 2.6) from Thermo Fisher Scientific. Statistical analyses were performed using GraphPad Prism (v 5.01, GraphPad Software Inc., La Jolla, CA).

3. Results

3.1. Reaction standardization of estrogens and their metabolites with the NMPS derivatives

A scheme showing the NMPS derivatization reaction is shown in Fig. 2. Each of the derivatization steps was essentially complete within 30 min. E2, 16 α -OH-E2, 4-MeO-E2 and 2-MeO-E2 contain only one phenolic group and reacted with one molar equivalent of the pyridine-3-sulfonyl chloride, generating pyridine-3-sulfonyl (PS) derivatives that were then converted to NMPS derivatives with iodomethane. The catechol estrogens, 4-OH-E2 and 2-OH-E2, which each contains two phenolic hydroxyl groups, reacted with two molar equivalents of the pyridine-3-sulfonyl chloride generating *bis*-PS derivatives. Interestingly, each catechol PS derivative only formed mono-NMPS derivatives as a mixture of two regioisomers (Fig. 2).

3.2. LC-MS/MS analysis of estrogen NMPS derivatives

Each of the estrogen derivatives exhibited intense ions corresponding to $[M^+]$ ion of the relevant NMPS derivative. Collision induced dissociation (CID) and MS/MS analysis revealed the formation of three major product ions (Fig. 3) from each of the estrogen derivatives. Unique product ions were observed for E2, 16 α -OH-E2, isomeric 4-OH-E2 and 2-OH-E2 derivatives at m/z 364.2, 380.2 and 379.2, respectively (Fig. 4). This corresponded to the unusual attachment of the pyridine moiety to the parent molecule after losing SO₂ from the derivative (fragment c, Fig. 4). However, this predominant ion was not observed in 4-MeO-E2 and 2-MeO-E2, where the major product ions arose from the loss of the E2 moiety from both of the derivatives (m/z 158.2; fragment b, Fig. 4). The product ion corresponding the pyridinium moiety (m/z 93.1; fragment a, Fig. 4) was observed for all of the estrogen NMPS-derivatives. As expected, the corresponding internal standards gave similar product ions to NMPS derivatives of the endogenous metabolites (Fig. 3).

3.3. Method validation

Typical LC/SRM-MS chromatographic profiles of the six estrogen-NMPS derivatives together with their corresponding six internal standards are shown for the LLOQ standard of 0.5 pg/mL for E2, 16 α -OH-E2, 4-MeO-E2 and 2-MeO-E2, and 5 pg/mL for 4 and 2-OH-E2 (Fig. 5A). A chromatogram is shown for the HQC standard of 175 pg/mL for E2, 16 α -OH-E2, 4-MeO-E2 and 2-MeO-E2, 1750 pg/mL for 4-OH-E2 and 2-OH-E2 (Fig. 5B). The six estrogen derivatives were baseline separated within a 20 min chromatographic run time. The two co-eluting regioisomers from 4-OH-E2 and 2-OH-E2 are shown with an asterisk (Figs. 5 and 6). Comparison of the NMPS and PS derivatives revealed an increase in sensitivity and signal-to-noise for analysis of unconjugated estrogens in the same serum sample (Fig. 7).

3.3.1. Calibration curve and limit of quantification—Calibration curves for each estrogen were constructed from the ratios of the peak area of the NMPS derivatives to corresponding internal standard with 1/x weighting. Satisfactory linearity was observed over 400-fold concentration with linear regression correlation coefficients all better than 0.99. (Table 4) The lower LLOQ was defined as the lowest concentration on the calibration curve

that could be reliably and reproducibly measured with accuracy and precision of less than 20% and a signal-to-noise ratio great than 10.

3.3.2. Assay accuracy, precision, recovery and stability of free estrogens—

Overall, excellent accuracy and precision were obtained for the analysis of all three QC serum samples (Table 5). For the LQC, MQC and HQC, the intra-day accuracies were 102–118%, 95–108%, and 98–105%; the intra-day precisions were 4.5–13.9%, 0.9–3.4%, 0.6–4.5%; the inter-day accuracies were 91–124%, 102–111%, and 95–106%, and the inter-day precisions were 4.4–19.9%, 1.1–4.4%, 4.2–5.8%. The recovery of six estrogens after extraction, derivatization, and purification from LQC, MQC, and HQC samples (n=5) was 84–98% for E2, 92–106% for 16 α -OH-E2, 99–109% for 4-MeO-E2, 98–112% for 2-MeO-E2, 95–109% for 4-OH-E2, and 90–100% for 2-OH-E2 (Table 6). The precision and accuracy data for analysis of the LQC, MQC, and HQC samples (n=5) after three freeze thaw cycles were similar to that shown in Table 5 for the inter-day validation (data not shown). LQC, MQC, and HQC samples that were re-analyzed after 24 h standing in the autosampler (4 °C) gave essentially identical data to that obtained from the original analyses.

3.3.3. Recovery of estrogens after hydrolysis and extraction procedure—

Quantification of total (unconjugated + conjugated) estrogens was performed by way of enzymatic hydrolysis of E2-3G-17S in water, following by extraction, derivatization and analysis by LC-SRM/MS. The hydrolysis efficiency was > 90% and was complete after 19 h. Complete hydrolysis under this condition was confirmed by analyzing triplicate control samples with 0.037 pmol/mL of E2-3G-17S (equivalent of 10 pg/mL of free E2) and 0.37 pmol/mL of E2-3G-17S (equivalent of 100 pg/mL of free E2) (Table 7).

3.4. Analysis of estrogens in patient serum samples

A total of 40 serum samples (20 female and 20 male) were processed using the LC-SRM/MS/MS assay. All estrogen concentrations were within the range of standard curves or below the LLOQs. Representative chromatograms of serum estrogens for a typical female subject are shown in Fig. 6. Unconjugated E2 was quantified in female (Fig. 6A) and male samples (chromatograms not shown). However, 16 α -OH-E2, 4-MeO-E2, 2-MeO-E2, 4-OH-E2 and 2-OH-E2 were below the LLOQ in female (Fig. 6A) and male samples (chromatograms not shown). Total E2 and 16 α -OH-E2 could be quantified in the female samples (Fig. 6B), whereas only E2 and 16 α -OH-E2 could be quantified in the male samples (chromatogram not shown). The mean concentrations (\pm SEM) of unconjugated E2 in female and male serum were 2.7 pg/mL (\pm 0.6 pg/mL) (Fig. 8A) and 9.5 pg/mL (\pm 0.8 pg/mL) (Fig. 8D), respectively. All other unconjugated estrogens were below the LLOQ (0.5 pg/mL for E2, 16 α -OH-E2, 4-MeO-E2 and 2-MeO-E2, 5 pg/mL for 4-OH-E2 and 2-OH-E2). The mean concentration (\pm SEM) of total E2, 16 α -OH-E2, and 4-MeO-E2 in the female samples were 20.7 pg/mL (\pm 4.1 pg/mL), 32.5 pg/mL (\pm 8.1 pg/mL), and 8.2 pg/mL (\pm 1.9 pg/mL), respectively (Fig. 8C). The mean concentration (\pm SEM) of total E2 and 16 α -OH-E2, in the male samples were 17.4 pg/mL (\pm 1.1 pg/mL) and 36.5 pg/mL (\pm 12.9 pg/mL), respectively (Fig. 8F). There were significant differences in unconjugated E2 and total 4-MeO-E2 concentrations between the female and male groups ($p < 0.05$).

4. Discussion

Estrogens play an important role in the etiology of breast cancer and elevated concentrations of circulating E2 are thought to be associated with an increased risk for breast cancer in postmenopausal women [10,38,47–55]. E2 could increase breast cancer risk through mechanisms in which E2 acts either as a hormone to stimulate aberrant cell proliferation or as the precursor to the formation of genotoxic catechol metabolites [52,55,56]. E2 is metabolized through three hydroxylation pathways - 4-hydroxylation, 2-hydroxylation, and 16 α -hydroxylation (Fig. 1) [13,57]. Catechols derived from both the 4- and 2-hydroxylation pathways can be readily oxidized to form electrophilic quinone derivatives. These reactive electrophiles can form stable or depurinating adducts with DNA, which if not repaired could lead to mutations in the DNA that induce tumorigenesis [58]. Methylation of the 4-OH-E2 and 2-OH-E2 (Fig. 1) would prevent their conversion to reactive quinones. In addition, the methylated metabolites 4-MeO-E2 and 2-MeO-E2 could potentially serve as biomarkers of catechol formation.

Over the last decade, MS-based methods for the quantification of unconjugated serum E2 in postmenopausal women have become more specific and accurate so that more recently GC-MS and LC-MS methods have generally given values that are < 10 pg/mL (Table 1). Multiplexed assays for multiple estrogens are easier to conduct by LC-MS and so most of the reported values have employed this technique. Unfortunately, the reported values of all of the unconjugated serum estrogens apart from E2 are close to the limits of sensitivity of the assays and so it is not clear that these concentrations have any diagnostic value. This impedes statistical efficiency for epidemiologic studies, preventing accurate stratification and correlation analysis. Moreover, there is still considerable variability between laboratories in the analysis of individual samples [59]. These issues, coupled with general concerns over the reliability of E2 measurements [60], and the need for improved sensitivity to monitor aromatase inhibition [61], suggested the need for a more specific, sensitive, and reliable method for the analysis of E2 and its major metabolites to complement the high sensitivity method we developed for estrone and its metabolites [62].

Since the levels of estrogens and their metabolites are in the low pg/mL range, it was very important to optimize each sample preparation step to obtain as high recovery as possible. The extraction solvents, the effects of reaction heating time and temperature, the concentration of pyridine-3-sulfonyl chloride and iodomethane, and reaction pH were optimized. Solvents were tested for their ability to efficiently extract the estrogens and their metabolites from human serum, including ethyl acetate, ethyl acetate/hexane (60:40), dichloromethane, MTBE, and MTBE/hexane (9/1, 5/5). MTBE was the most efficient solvent to extract all estrogens tested; furthermore, it removed many of the co-eluting interfering peaks. Acidification of serum and addition of saturated brine also improved recovery. There were almost quantitative recoveries of estrogens and PS-estrogens derivatives from both LLE steps. The methylation reaction is a critical step for NMPS derivatization. High quality acetonitrile is recommended as the solvent for the iodomethane reaction. Additionally, the PS derivatives should be evaporated to complete dryness before methylation to ensure reaction efficiency. We compared aliquots of patient samples using both NMPS and PS derivatization. The NMPS derivatives gave a two-fold increase in

response when compared with the PS derivatives. Furthermore, the additional derivatization step lowered the noise in the analyte channel and thus increased the signal/noise ratio (Fig. 7). The highest yield of NMPS derivatives was obtained with sodium bicarbonate at pH 9. When other conditions were held the same, heating sample at 60 °C for 30 min for pyridinium-3-sulfonate derivatization followed by 30 min at 80 °C for N-methyl derivatization, gave the best yield of NMPS derivatives for all estrogens and their metabolites. The major catechol derivatives formed under the conditions that were employed were mono-N-methyl derivatives (Figure 3). However, any minor amounts of *bis*-N-methyl derivatives that might be formed would not have affected the ratio of analyte to stable isotope-labeled internal standard.

Micro-LC and capillary LC have been widely used as a routine technique in various bioanalytical laboratories. Theoretically, nanospray LC should increase the sensitivity of analysis and so should allow for further improvements in detection limits [63]. Consequently, we compared the sensitivity of a Waters 2695 system (0.2 mL/min) with a Waters nanoAcquity UPLC system (2 μ L/min) in the early stages of method development. The signal-to-noise increased by 2.5-fold at the lower flow rate and so method validation and subsequent analyses were conducted with a flow rate of 2 μ L/min. In order to obtain both reproducible separation and high sensitivity for estrogens and the internal standards within a reasonable separation time, the nanospray UPLC conditions were carefully optimized. Tests showed that a mobile phase consisting of water and acetonitrile, both containing 0.1% formic acid, was the best mobile phase system.

The NMPS derivatives allowed E2 and five of its metabolites to be analyzed with extremely high sensitivity but less sample consumption. The limit of detection for E2 was 1 fg on column and the LLOQ for serum E2 was 500 fg/mL with only 100 μ L of serum. Different to dansyl derivatives [31–34], NMPS derivatives also provided high specificity due to their analyte-dependent product ions, which arise from main structure of estrogen and its metabolites. This new derivative in combination with nano-LC-MS provided a mean value of 2.7 pg/mL (n=20) for unconjugated postmenopausal serum E2 (Table 1, Fig. 8A), which was close to the mean value of 4.4 pg/mL reported for GC-MS assays and lower the mean value of 7.3 pg/mL reported for LC-MS assays that used the dansyl derivative (Table 1). Furthermore, it agrees well with the mean value of 3.3 pg/mL that was reported recently from a novel method, which analyzed underivatized E2 using fluoride-enhanced negative ESI coupled with LC-MS [64]. This confirms that the NMPS derivative confers high specificity as well as high sensitivity for E2 analysis. The mean value of unconjugated serum E2 found for 20 older men of 9.6 pg/mL (Fig. 8C) was significantly higher than that found in postmenopausal women (Fig. 8A). It is noteworthy that higher levels of unconjugated E2 found in men compared with postmenopausal women is similar to many other studies, although the absolute level was somewhat lower. For example, a mean value of 20.6 pg/mL was reported for unconjugated serum E2 in six studies of older men, which used the PFBO derivative coupled with GC-MS-based methodology [27,65–69]. The mean unconjugated serum E2 level we found in men was also lower than studies that employed LC-negative ESI/MS (mean 17.2 pg/mL) [70], LC-negative atmospheric pressure photoionization/MS (mean 20 pg/mL) [71], and regular LC-ESI/MS of the E2 dansyl

derivative (mean 22.9 pg/mL) [72–74]. This provides further evidence that the NMPS derivative in combination with LC-MS confers high sensitivity and specificity for E2 analysis. The five other unconjugated serum estrogens were all below the LLOQ for both postmenopausal women (Fig. 8A; Table 1) and older men (Fig. 8D). This together with previous reports of values that were close to the reported assay LLOQs suggests that such measurements have little diagnostic value.

The mean level obtained for total serum E2 of 20.7 pg/mL in postmenopausal women (Table 2, Fig. 8C) was similar to the mean value of 19.7 pg/mL that was obtained using LC-MS of the dansyl derivative (Table 2). A relatively low value for serum E2-3G of 5.5 pg/mL was found using an excellent stable isotope dilution assay of the intact β -glucuronide [26]. In our study the β -glucuronide and sulfate conjugates (total E2 - unconjugated E2) corresponded to 18 pg/mL (Fig. 8B) suggesting that the sulfate was the major conjugate of E2 (Fig. 1) that was present in the serum. This would potentially be a source for E2 through the action of sulfatases that are present in breast tissue [75,75] and so analysis of total serum E2 seems to be a worthwhile endeavor. Perhaps more specific methodology for intact E2-3S similar to that developed for E2-3G [26] and E1-S [76–78] would be even more meaningful.

The concentrations of total serum E2 in older men of 17.4 pg/mL (Fig. 8F) were, surprisingly, quite similar to the levels found in postmenopausal women (Fig. 8C). This means that the β -glucuronide and sulfate conjugates (7.9 pg/mL, Fig. 8E) were lower than the corresponding unconjugated serum E2 (Fig. 8D) concentration in the male subjects. A previous LC-MS study using the dansyl derivative reported a mean serum total E2 of 62.3 pg/mL in older men [79], which was significantly higher than that found in the present study. However, it is noteworthy that this study also reported total serum estrogen concentrations in postmenopausal women that were much higher than has been reported previously (Table 2). This suggests that there was some interference in the method that was used, and that the correct total serum E2 concentrations are closer to value of 17.4 pg/mL determined in the present study.

There is no consensus in the literature on the correct levels of total serum 16 α -OH-E2 – reported values span a wide range from 27.9 pg/mL to 126.0 pg/mL (Table 2). The value obtained in the present study of 32.5 pg/mL (Table 2, Fig. 8C) seems to be more realistic in view of the undetectable levels of the unconjugated form in serum. In fact our data would suggest that 16 α -OH-E2 is only present as β -glucuronide and/or sulfate conjugates (Fig. 8B). It is conceivable that the differences between the various studies could be due to differences in efficiency of hydrolysis by glucuronidase/sulfatase enzyme and so it will be important in the future to validate our finding through analysis of the intact β -glucuronide and sulfate conjugates. The mean concentration of total serum 16 α -OH-E2 of 36.5 pg/mL obtained for older men (Fig. 8F) was very similar to the value observed in postmenopausal women (Table 1, Fig. 8C).

Previous studies have suggested that levels of total 4-MeO-E2 in serum from postmenopausal women are below 1 pg/mL and so analysis of this molecule cannot really be justified. The present study has revealed that the mean level is in fact slightly higher at 8.2 pg/mL (Table 2, Fig. 8C), suggesting that total serum 4-MeO-E2 might be a useful

LC	liquid chromatography
HQC	high quality control
LLE	liquid-liquid extraction
LLOQ	lower limit of quantification
LQC	low quality control
MQC	medium quality control
MS	mass spectrometry
<i>m/z</i>	mass to charge ratio
NMPS	N-methyl pyridinium-3-sulfonyl
OH	hydroxy
OMe	methoxy
PS	pyridine-3-sulfonyl
S	sulfate

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Highlights

- Estrogen NMPS derivatives facilitate high sensitivity LC-ESI/MS analysis.
- Stable isotope dilution LC-MS method developed for six serum estrogens.
- Mean unconjugated serum E2 was 2.9 pg/mL in postmenopausal women
- Mean unconjugated serum E2 was 9.1 pg/mL in older men
- Conjugated serum 4-MeO-E2 is potential cancer biomarker.

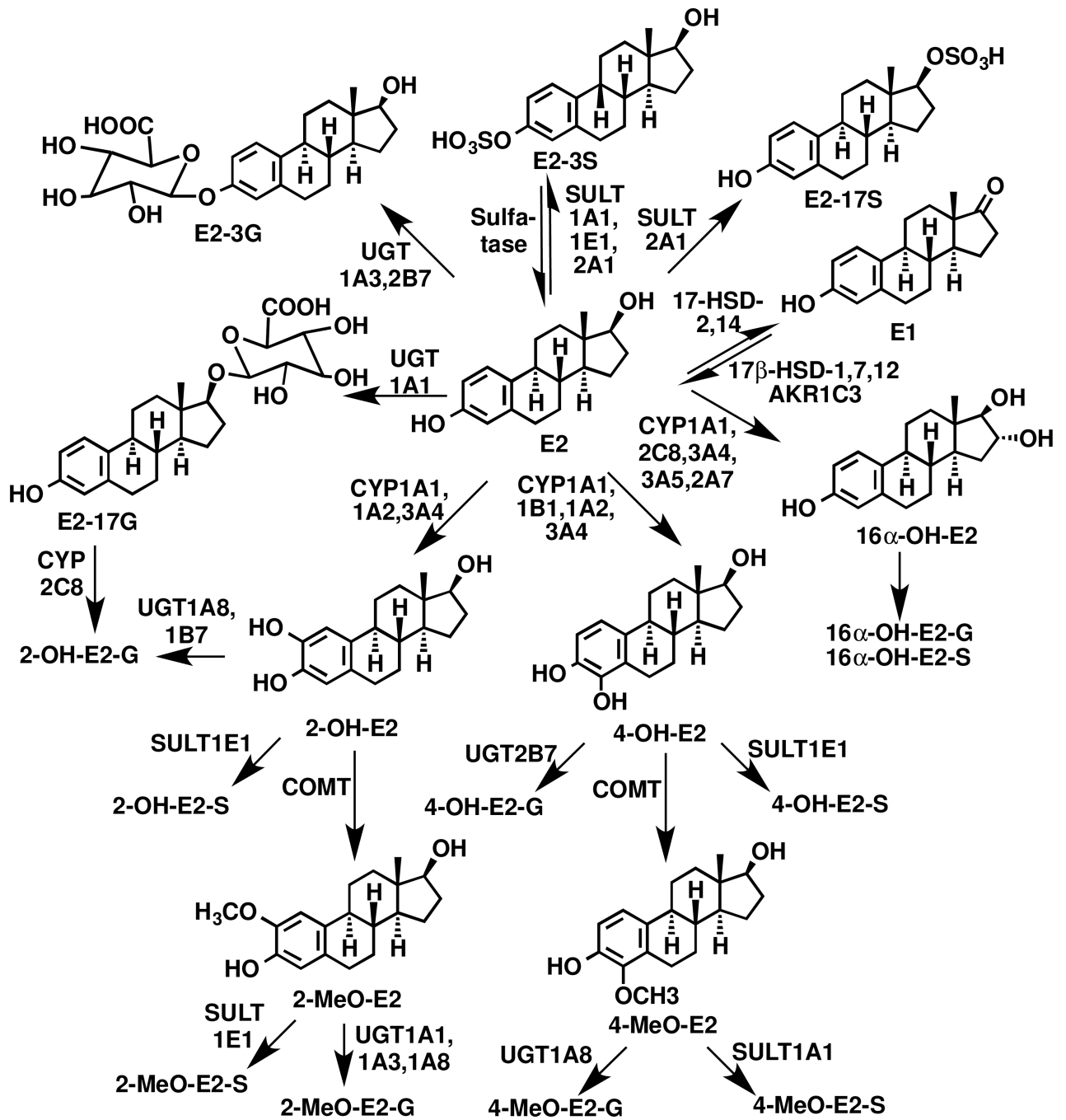


Figure 1.
Enzymatic pathways involved in estradiol metabolism.

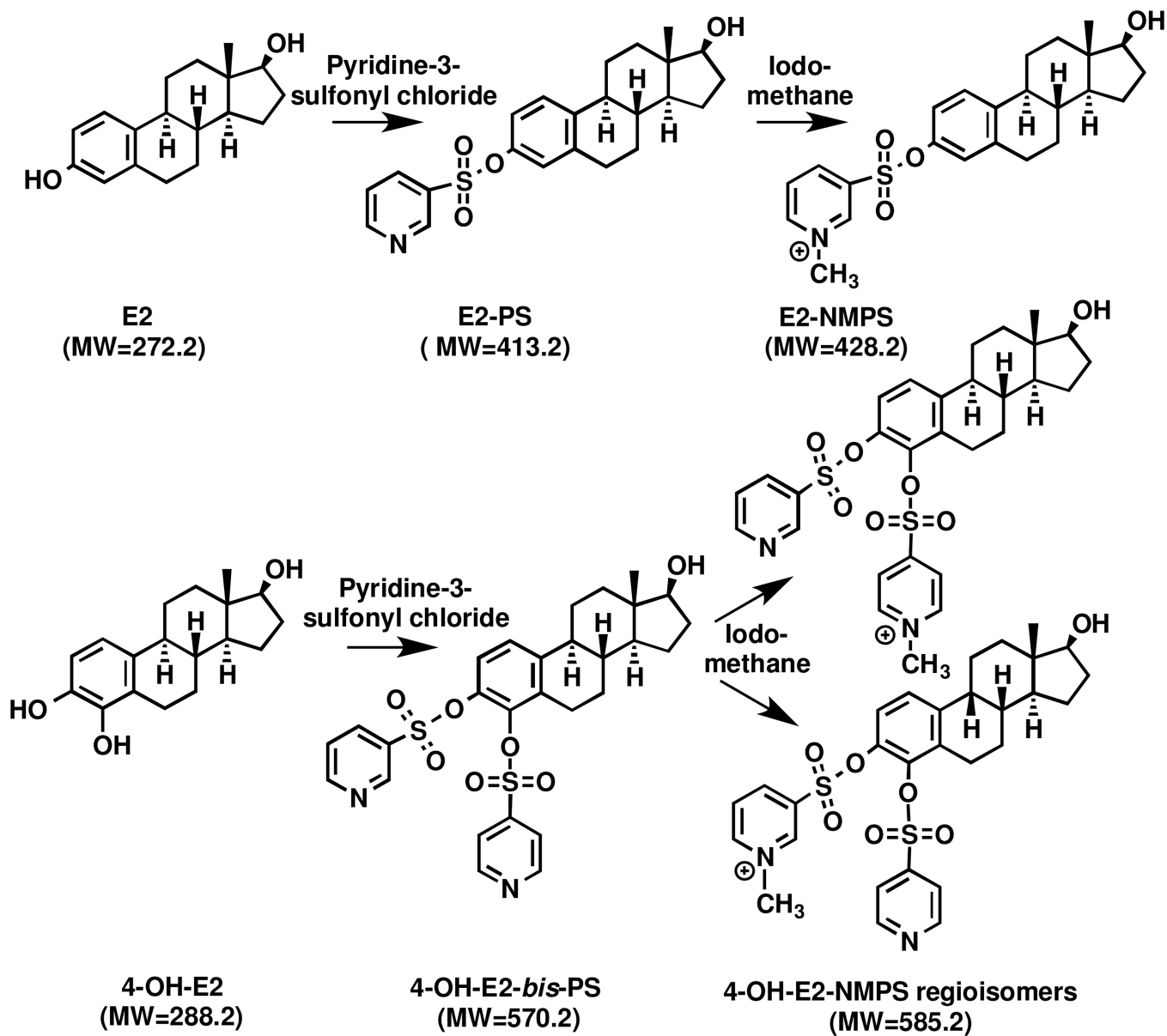


Figure 2.

Two step reaction of pre-ionized N-methyl pyridine-3-sulfonyl (NMPS) derivatives and their structures.

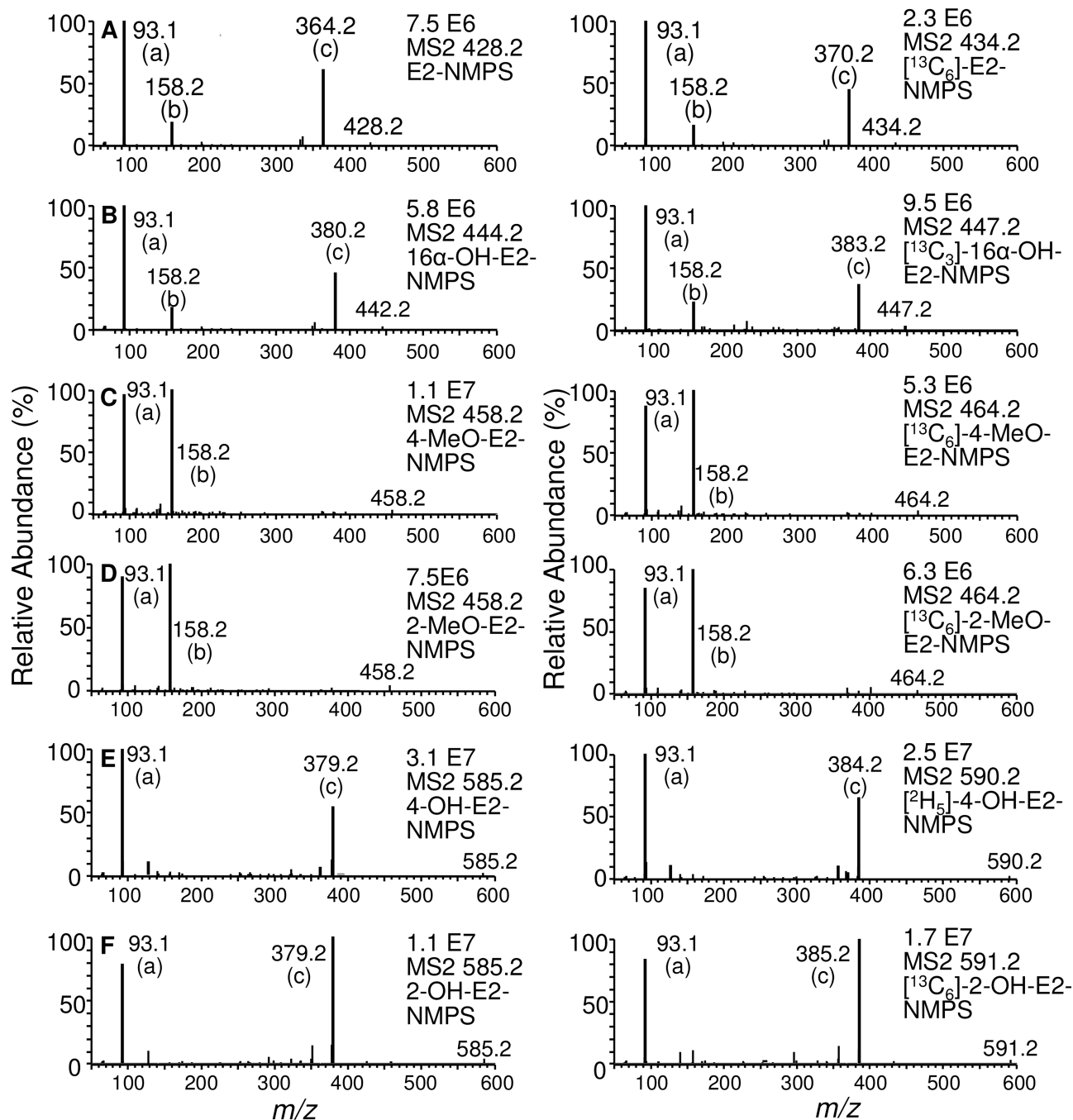
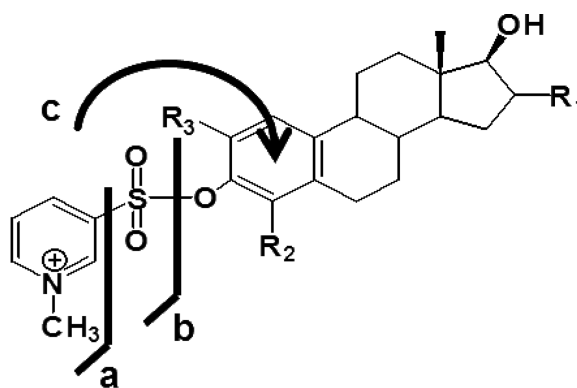
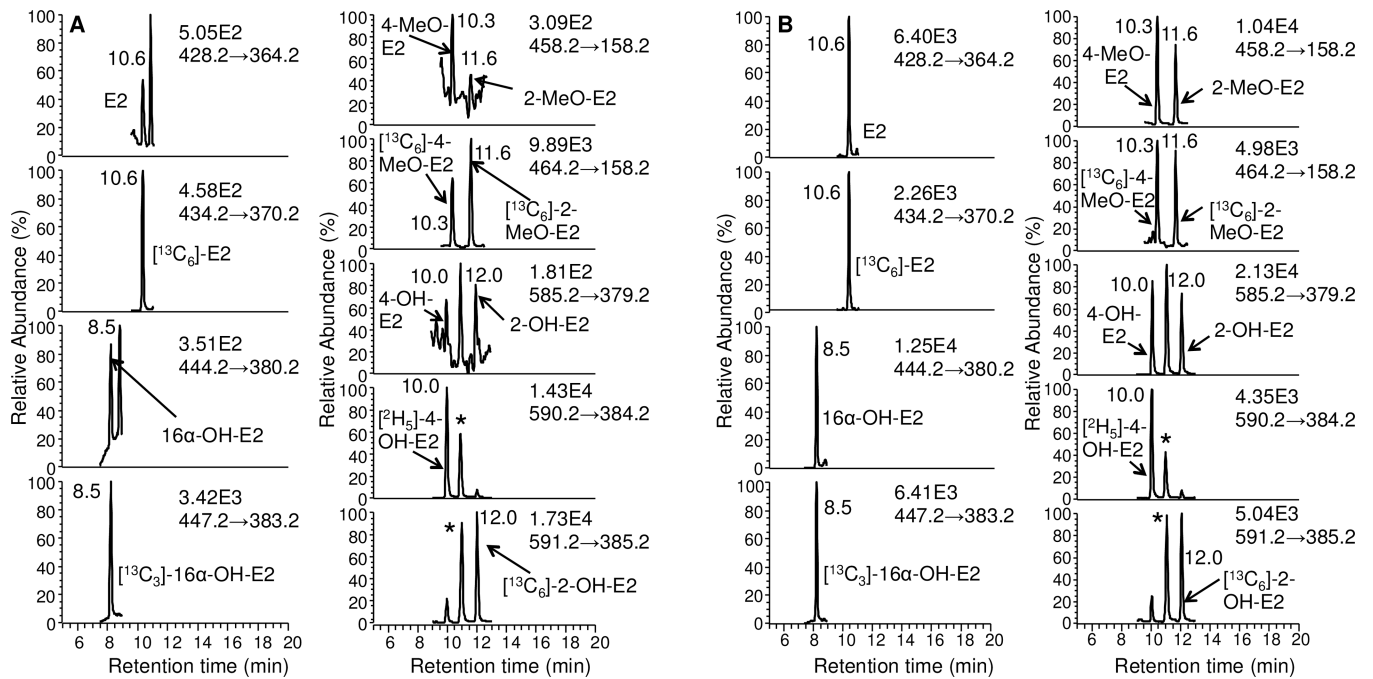


Figure 3. Full scan MS/MS analysis of product ions of E2-NMPS, 16 α -OH-E2-NMPS, 4-MeO-E2-NMPS, 2-MeO-E2-NMPS, 4-MeO-E2-NMPS and 2-OH-E2-NMPS derivatives. The most intense product ions were selected for the SRM analysis.



Estrogen	Substituent			Product ions		
	R1	R2	R3	a (m/z)	b (m/z)	c (m/z)
E2	H	H	H	93.1	158.2	364.2
[¹³ C ₆]-E2	H	H	H	93.1	158.2	370.2
16α-OH-E2	OH	H	H	93.1	158.2	380.2
[¹³ C ₃]-16α-OH-E2	OH	H	H	93.1	158.2	383.2
4-MeO-E2	H	OCH3	H	93.1	158.2	ND
[¹³ C ₆]-4-MeO-E2	H	OCH3	H	93.1	158.2	ND
2-MeO-E2	H	H	OCH3	93.1	158.2	ND
[¹³ C ₆]-2-MeO-E2	H	H	OCH3	93.1	158.2	ND
4-OH-E2	H	OH	H	93.1	ND	379.2
[² H ₅]-4-OH-E2	H	OH	H	93.1	ND	384.2
2-OH-E2	H	H	OH	93.1	ND	379.2
[¹³ C ₆]-2-OH-E2	H	H	OH	93.1	ND	385.2

Figure 4.
Assignment of product ions from LC-MS/MS analysis of estrogen NMPS derivatives.

**Figure 5.**

LC-SRM chromatograms for analysis of estrogens and their metabolites extracted from double charcoal-stripped human serum as NMPS derivatives. (A) LLOQ samples (0.5 pg/mL for E2, 16α-OH-E2, 4-MeO-E2 and 2-MeO-E2; 5 pg/mL for 4-OH-E2 and 2-OH-E2). (B) HQC samples (175 pg/mL for E2, 16α-OH-E2, 4-MeO-E2 and 2-MeO-E2; 5 pg/mL for 4-OH-E2 and 2-OH-E2). Asterisks show co-eluting second regioisomers from 4-OH-E2 and 2-OH-E2.

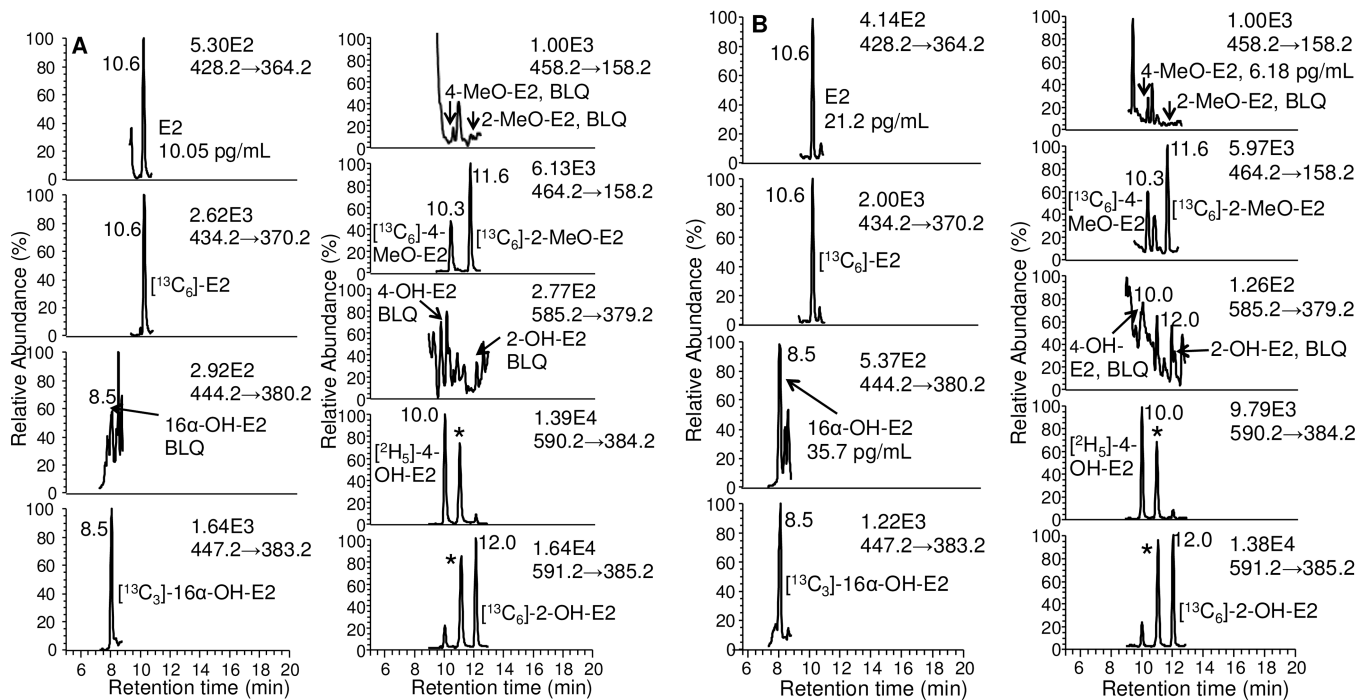


Figure 6. LC-SRM/MS chromatograms obtained from analysis of estrogens and their metabolites as NMPS derivatives in serum from postmenopausal women. (A) unconjugated estrogens (B) total estrogens. Asterisks show co-eluting second regioisomers from 4-OH-E2 and 2-OH-E2.

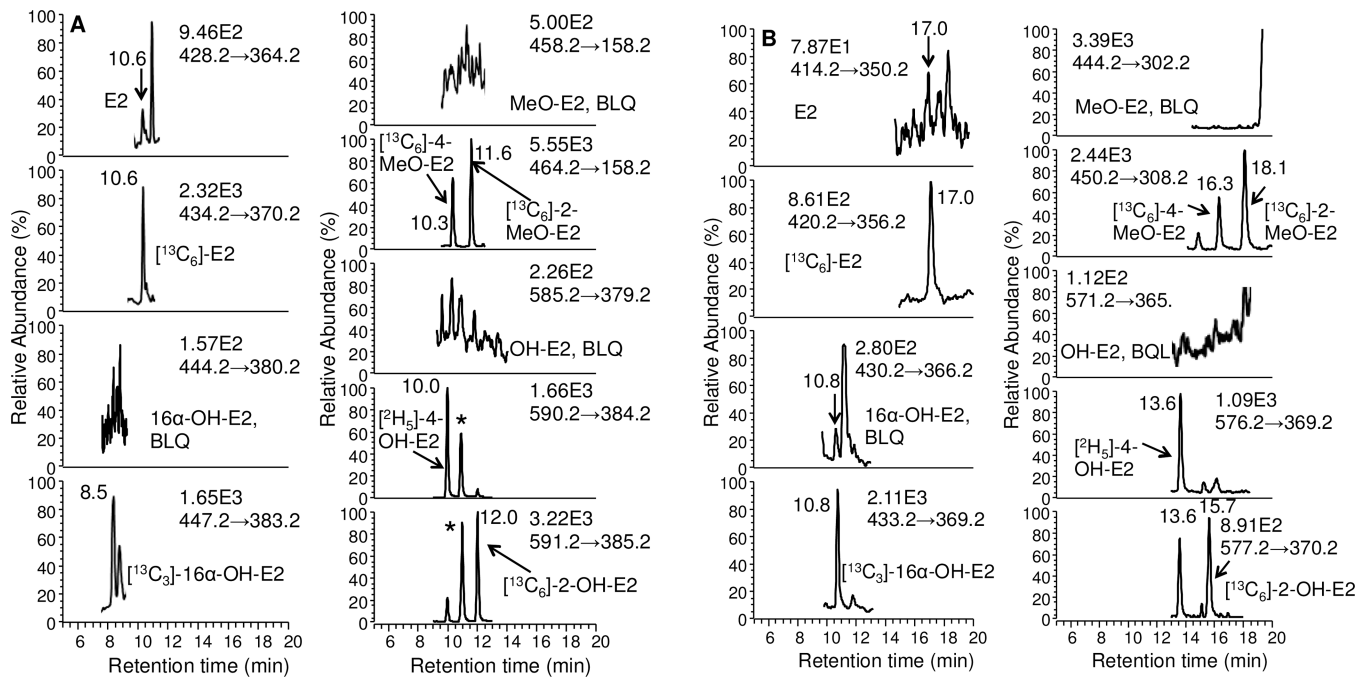


Figure 7. Comparison of derivatives for analysis of unconjugated serum estrogens from the same patient sample. (A) NMPS derivatives. (B) PS derivatives. Asterisks show co-eluting second regioisomers from 4-OH-E2 and 2-OH-E2.

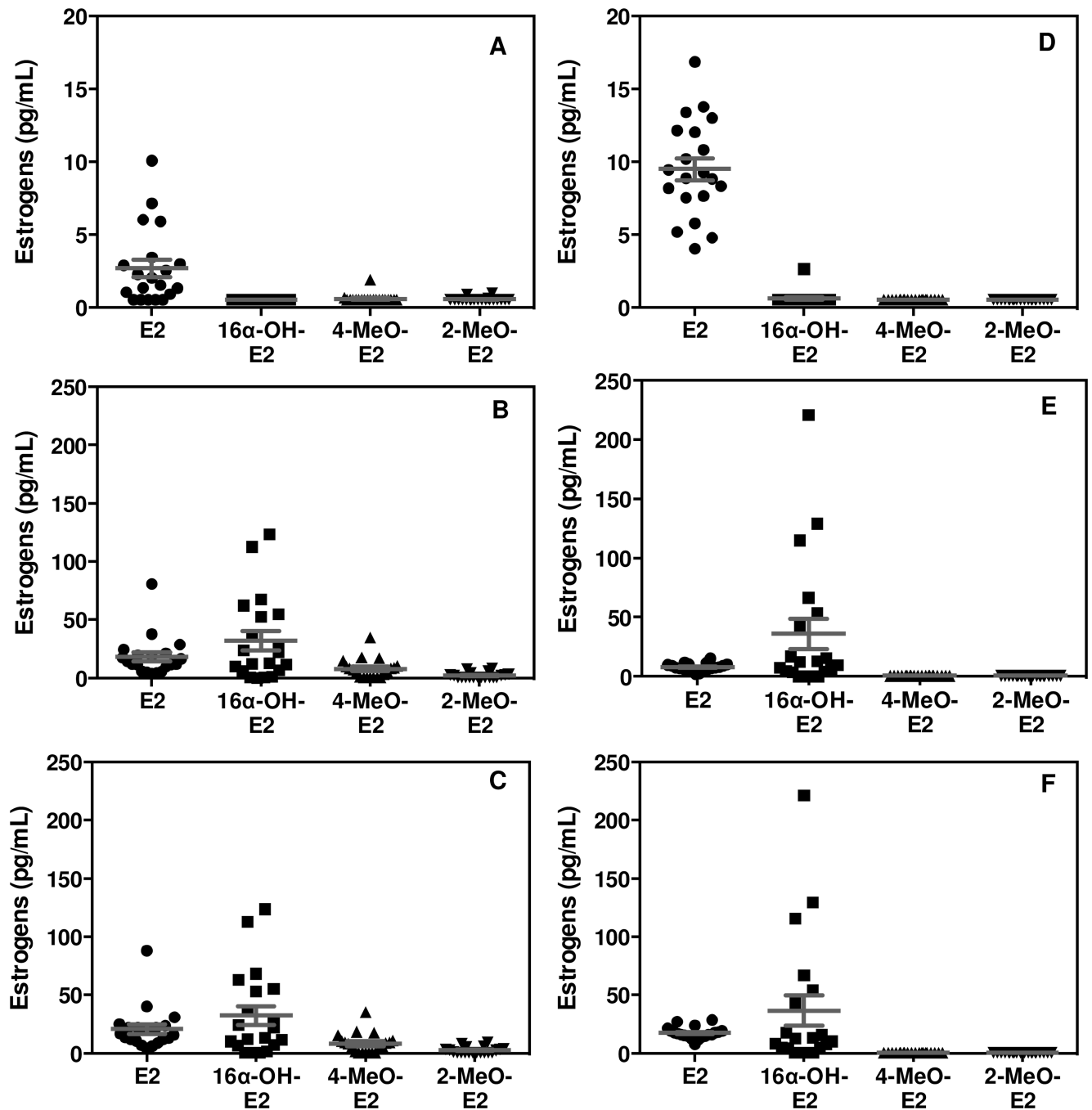


Figure 8.

Mean values for estrogen concentrations (pg/mL) in serum samples from 20 postmenopausal women (mean age 60.0) and 20 older men (mean age 61.6). (A) Unconjugated serum estrogens in postmenopausal women. (B) Conjugated serum estrogens in postmenopausal women. (C) Total serum estrogens in postmenopausal women. (D) Unconjugated serum estrogens in older men. (E) Conjugated serum estrogens in older men. (F) Total serum estrogens in older men. Individual values \pm SEM are shown.

Table 1

Concentrations of unconjugated estrogens in postmenopausal serum determined by GC-MS and LC-MS.

Method	LC-MS		GC-MS		GC-MS		GC-MS		LC-MS		GC-MS		LC-MS		LC-MS		Overall Mean pg/mL	
	Dansyl	Mean	PFBO	Mean	PFBO	Mean	PFBO	Mean	Dansyl	Mean	PFBO	Mean	Dansyl	Mean	Dansyl	Mean		
Reference	35		23	24	25	26	27	36	37	38	28	39						
Derivative	Dansyl		PFBO	PFBO	PFBO	PFBO	PFBO	Dansyl	PFBO	Dansyl	PFBO	PFBO	Dansyl	PFBO	Dansyl	PFBO	Mean	
pg/mL	Mean		Mean	Mean	MeanL	Mean	Mean	Mean	Median	Mean	Mean	Median	Mean	Median	Mean	Mean	Mean	
E2	15.0		3.5	7.3	3.0	6.6	4.2	15.9	3.4	3.1	3.1	3.4	3.1	4.2	2.7	3.1	2.7	5.1
16 α -OH-E2	7.9		x	x	x	x	x	x	x	x	x	x	x	13.5	7.7	x	7.7	10.6
4-MeO-E2	BLQ		x	x	x	x	x	x	x	BLQ	x	x	x	BLQ	BLQ	x	BLQ	BLQ
2-MeO-E2	BLQ		x	x	x	x	x	x	x	1.4	x	x	x	0.7	BLQ	x	0.7	1.0
4-OH-E2	x		x	x	x	x	x	x	x	x	x	x	x	x	BLQ	x	BLQ	BLQ
2-OH-E2	BLQ		x	x	x	x	x	x	x	BLQ	x	x	x	BLQ	BLQ	x	BLQ	BLQ

X = not determined

BLQ = below LLOQ

PFBO = pentafluorobenzoyl

Table 2
Concentrations of conjugated estrogens in postmenopausal serum determined by LC-MS.

Technique	G/S		Intact	Intact		G/S		G/S		G/S		Overall Mean pg/mL
	Dansyl	None		None	12	Dansyl	38	Dansyl	39	Dansyl	40	
Derivative												
Reference	35	26	26	12	38	39	40	40	40	40		
pg/mL	Mean	Mean	Mean	Median	Median	Median	Median	Median	Median	Median	Mean	Mean
E2-T	51.5	x	x	x	9.8	6.2	10.5	6.2	10.5	20.7	20.7	19.7
E2-3G	x	8.4	x	2.5	x	x	x	x	x	x	x	5.5
16 α -OH-E2-T	27.9	x	x	x	126.0	78.4	70.5	78.4	70.5	32.5	32.5	67.1
4-MeO-E2-T	BLQ	x	x	x	0.8	0.6	0.8	0.6	0.8	8.2	8.2	2.6
2-MeO-E2-T	BLQ	x	x	x	2.3	4.7	4.0	4.7	4.0	2.5	2.5	3.4
2-MeO-E2-3G	x	6.9	x	2.5	x	x	x	x	x	x	x	4.7
4-OH-E2-T	x	x	x	x	x	x	x	x	x	BLQ	BLQ	BLQ
2-OH-E2-T	11.1	x	x	x	10.4	6.6	3.5	6.6	3.5	BLQ	BLQ	7.9

X = not determined

BLQ = below LLOQ

T = conjugated + unconjugated

G = glucuronide

G/S = β -glucuronidase/arylsulfatase hydrolysis as described in section 2.5.

Table 3

LC-SRM/MS conditions for estrogen NMPS derivatives.

Analyte	Parent (m/z)	Product (m/z)	Collision energy(V)	Start time (min)	Stop time (min)	S-lens (V)
E2	428.2	364.2	35	9.5	11.5	135
[¹³ C ₆]-E2	434.2	370.2	35	9.5	11.5	135
16 α -OH-E2	444.2	380.2	35	7.5	9.5	135
[¹³ C ₃]-16 α -OH-E2	447.2	383.2	35	7.5	9.5	135
MeO-E2	458.2	158.2	35	9.5	12.5	135
[¹³ C ₆]-MeO-E2	464.2	158.2	35	9.5	12.5	135
OH-E2	585.2	379.2	35	9.0	13.0	135
[² H ₅]-4-OH-E2	590.2	384.2	35	9.0	13.0	135
[¹³ C ₆]-2-OH-E2	591.2	385.2	35	9.0	13.0	135

Table 4

Typical calibration curves for estrogens as NMPS derivatives.

Analyte	Equations	Correlation coefficients (R ²)	LLOQ (pg/mL)
E2	Y=0.015X+0.066	0.9997	0.5
16 α -OH-E2	Y=0.011X+0.063	0.9992	0.5
4-MeO-E2	Y=0.012X+0.016	0.9998	0.5
2-MeO-E2	Y=0.010X+0.0031	0.9995	0.5
4-OH-E2	Y=0.025X+0.039	0.9987	5
2-OH-E2	Y=0.0016X+0.0097	0.9996	5

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Table 5
Accuracy and precision for determination of estrogens as NMPS derivatives in serum (n=5).

Analyte	LQC (1.5 pg/mL)			MQC (75 pg/mL)			HQC (175 pg/mL)					
	Mean	SD	CV	Accuracy	Mean	SD	CV	Accuracy	Mean	SD	CV	Accuracy
Intra-day												
E2	1.5	0.2	10%	102%	80.7	1.7	2%	108%	183.8	3.0	2%	105%
16 α -OH-E2	1.8	0.1	5%	118%	79.2	1.1	1%	106%	180.2	2.7	2%	103%
4-MeO-E2	1.6	0.2	14%	105%	77.9	1.6	2%	104%	176.5	8.0	5%	101%
2-MeO-E2	1.7	0.2	9%	112%	76.0	0.7	1%	101%	179.3	2.5	1%	102%
4-OH-E2	1.5	0.1	6%	102%	80.2	1.3	2%	107%	180.6	1.0	1%	103%
2-OH-E2	1.6	0.1	6%	107%	71.5	2.4	3%	95%	171.4	5.1	3%	98%
Inter-day												
E2	1.4	0.1	4%	91%	78.5	0.9	1%	105%	173.2	9.0	5%	99%
16 α -OH-E2	1.5	0.2	11%	102%	79.0	1.8	2%	105%	174.1	8.1	5%	99%
4-MeO-E2	1.6	0.2	11%	105%	83.2	3.7	4%	111%	185.7	10.8	6%	106%
2-MeO-E2	1.9	0.1	5%	124%	79.0	2.2	3%	105%	173.7	7.6	4%	99%
4-OH-E2	1.6	0.1	8%	104%	77.5	2.4	3%	103%	172.4	7.3	4%	99%
2-OH-E2	1.4	0.3	20%	95%	76.3	1.5	2%	102%	166.0	8.1	5%	95%

SD: standard deviation

CV: coefficient of variation

RE: relative error

RE (%) = (found concentration – actual concentration/actual concentration)×100

Accuracy (%) = RE (%) + 100

Table 6

Recovery of unconjugated estrogens after extraction, derivatization and analysis (n=5).

Analytes	LQC (1.5 pg/mL)				MQC (75 pg/mL)				HQC (175 pg/mL)			
	Post-spike	Pre-spike	Recov. (%)	SD	Post-spike	Pre-spike	Recov. (%)	SD	Post-spike	Pre-spike	Recov. (%)	SD
E2	1.62	1.37	84%	4.0	80.16	78.46	98%	1.1	184.23	173.24	94%	4.9
16 α -OH-E2	1.45	1.53	106%	13.9	86.31	78.98	92%	2.1	173.03	174.07	101%	4.7
4-MeO-E2	1.44	1.57	109%	11.4	78.73	81.77	104%	3.0	186.73	185.67	99%	5.8
2-MeO-E2	1.66	1.85	112%	5.4	78.87	78.96	100%	2.8	177.55	173.65	98%	4.3
4-OH-E2	1.43	1.56	109%	8.6	77.98	77.54	99%	3.1	182.29	172.42	95%	4.0
2-OH-E2	1.42	1.43	100%	18.9	78.71	76.30	97%	1.9	184.42	166.04	90%	4.4

Post-spike: Internal standard spiked into MTBE extracted human serum, followed by derivatization.

Pre-spike: Internal standard spiked into human serum, followed by MTBE extraction and derivatization.

Recov. (%) = [(pre-spike-post-spike)/post-spike] × 100

Table 7

Recovery of E2 from E2-3G-17S after hydrolysis, extraction, derivatization, and analysis (n=5).

E2-3G-17S (pmol/mL)	β -glucuronidase/arylsulfatase	Theoretical E2 from hydrolysis (pg/mL)	Calculated E2 from hydrolysis (pg/mL)	Yield (%)
0.037 (10 pg/mL E2 equivalent)	Without	0.0	0.8	8.0
	With	10.0	10.5	105.0
0.370 (100 pg/mL E2 equivalent)	Without	0.0	0.6	6.0
	With	100.0	96.6	96.6