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Analytical and biological variation of biomarkers of oxidative stress during the menstrual cycle

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Abstract

Little information is available on the intra-individual variability of oxidative stress biomarkers in healthy individuals and even less in the context of the menstrual cycle. The objective of this study was to characterize the analytical and biological variability of a panel of 21 markers of oxidative damage, antioxidant defence and micronutrients in nine healthy, regularly menstruating women aged 18–44 years. Analyses included measurement of lipid peroxidation, antioxidant enzymes and antioxidant vitamins. Blood specimens were collected, processed and stored using standardized procedures on days 2, 7, 12, 13, 14, 18, 22 and 28 in one cycle for each subject. Replicate analyses of markers were performed and two-way nested random effects ANOVA was used to describe analytical, intra-individual and inter-individual variability. No statistically significant differences at $\alpha = 0.05$, or temporal effects across the menstrual cycle were observed. Analytical variability was the smallest component of variance for all variables. The ICC among replicates ranged from 0.80 to 0.98. Imprecision based on quality control materials ranged from 1 to 11%. The critical differences between serial results varied greatly between assays ranging from 6 to 216% of the mean level. These results provide important initial information on the variability of biomarkers of oxidative stress, antioxidant defence and micronutrients across the menstrual cycle.

Keywords

Oxidative stress; antioxidants; biological variation; menstrual cycle

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Introduction

Oxidative stress (OS) is an imbalance between free radicals and reactive oxygen species (ROS) and protective radical scavenging antioxidants (AOX) resulting from either an overproduction of ROS or a deficit in AOX protection (Terada 2006). Biomarkers of OS are increasingly being evaluated in experimental, clinical and epidemiological studies and have been implicated in the pathogenesis of numerous diseases including atherosclerosis (Olinski et al. 2002), cancer (Olinski et al. 2002), diabetes (Dandona 2002), respiratory disease (Paredi et al. 2002) and others (Multhaup & Masters 1999, Mathur et al. 2002, Soory 2002, Spector 2000, Agar & Durham 2003, Basu 2003, Dawson & Dawson 2003, Kruidenier et al. 2003), including conditions associated with premature birth (Matsubasa et al. 2002).

Many methods for the measurement of OS have been proposed and several reviews have been written (Morrow et al. 1992, Spiteller & Spiteller 1997, Davies et al. 1999, De Zwart et al. 1999, Hensley et al. 2000, Collins et al. 2004, Pratico et al. 2004). However, there is no current consensus on which methods are the most useful, reliable, accurate or specific for different types of oxidative insults (Pryor 1999). Despite successful application of these markers in some studies, laboratory variability has been a concern during the evaluation of published studies. Little information is available on the variability of OS markers in healthy individuals, especially as they vary across the menstrual cycle. Further *in vivo* studies assessing the variability and reliability of OS measures are needed (Pryor & Godber 1991).

There is increased interest in investigating the role of OS in female reproduction, including the aetiology of infertility and endometriosis (Agarwal et al. 2005). While the mechanisms that relate OS with female fertility are not completely understood, recent animal and human studies have suggested that OS may have an important role in spontaneous abortion (Samborskaia & Ferdman 1966, Jenkins et al. 2000), infertility in men and women (Rayman 2000, Xu et al. 2001), birth weight (Kim et al. 2005), follicular growth and development (Murray et al. 2001), endometriosis (Murphy et al. 1998) and regulation of endometrial angiogenesis (Maas et al. 2001). Investigation of the relationships between OS and female fertility and reproductive outcomes may be masked by normal menstrual cycle variation.

To evaluate the relation between OS and female fertility we selected a panel of biomarkers for longitudinal study across the menstrual cycle. As markers of oxidative damage we measured malondialdehyde (MDA) as thiobarbituric acid reactive substances (TBARS) and a high-performance liquid chromatography (HPLC) lipid peroxidation profile of linoleic acid and its hydroxy and hydroperoxy oxidation products including 13-hydroxy octadecadieneoic acid (13-HODE), 13-hydroperoxy octadecadieneoic acid (13-HpODE), 9-hydroxy octadecadieneoic acid (9-HODE) and 9-hydroperoxy octadecadieneoic acid (9-HpODE). As measures of antioxidant defence we measured the erythrocyte activity of superoxide dismutase (SOD), glutathione peroxidase (GPx) and glutathione reductase (GR), the plasma activity of glutathione peroxidase (pGPx) and the serum activity of paraoxonase 1 (PON1). As measures of micronutrient antioxidants we measured an HPLC fat-soluble vitamin profile (retinol, α , γ , and δ tocopherol and carotenoids) and total vitamin C.

The aim of this pilot study was to characterize the components of biological and measurement variability for the 21 OS markers measured by the laboratory. This included evaluation of laboratory measurement error (i.e. analytical acceptability index), the ability to distinguish significant changes in biomarker levels within (i.e. critical difference) and between (i.e. intraclass correlation coefficient) subjects, the feasibility of employing reference ranges of 'normal' to characterize individuals in terms of populations (i.e. the index of individuality), as well as estimation of the number of specimens required for characterization of the homeostatic set point. These data will be used to set analytical quality specifications, as well as to select

those biomarkers most appropriate, for consideration in a larger observational study investigating the effects of menstrual cycle hormones on OS.

Materials and methods

Participants and specimens

Specimens were collected from nine regularly menstruating, premenopausal, female volunteers. Major exclusion criteria included: current use of oral contraceptives, vitamins, supplements, and prescription or over-the-counter medications; pregnancy in the past 6 months or actively trying to conceive in the next 3 months; history of certain gynaecological disorders (including abnormal Pap smear) and/or surgery; positive screen (IgM) for *Chlamydia* infection at screening; measured body mass index (BMI; kg m^{-2}) <18.5 or >35.0 ; consuming a restricted or supplemented diet; history of physician-diagnosed chronic illness; or recent infection or X-ray exposure.

Fasting blood and urine specimens were collected between 7:00 and 8:30 a.m. on (approximately) days 2, 7, 12, 13, 14, 18, 22 and 27 of a 28-day cycle, to correspond to menstruation, early and late follicular phase, ovulation and early and late luteal phase. Fertility monitors (Clear Blue™) were used beginning on day 6 of the menstrual cycle to assist in timing of specimen collection. Collection and handling protocols were designed to minimize variability in pre-analytical factors (Murphy et al. 2000). Briefly, each participant observed 10 min of seated resting in order to allow fluid shifts to equilibrate. Phlebotomy tourniquets were applied for less than 1 min and removed immediately upon establishing blood flow. Tubes were protected from light, placed on ice and delivered to the processing laboratory within 15 min. All specimens were centrifuged at 4–8°C, portioned into multiple aliquots containing sufficient volume to provide for duplicate analysis for each assay and frozen within 90 min of phlebotomy.

Laboratory methods

Unless otherwise noted all chemicals were from Sigma Chemical Co. (St Louis, MO, USA). All laboratory measurements (144 determinations) were made in a single analytical batch using one set of reagents. For the HPLC linoleic acid peroxidation profile simultaneous measurement of all 144 determinations was impossible due to time requirements; therefore all specimens from one participant were measured together in nine consecutive sets.

HPLC lipid peroxidation profiles—Total lipid-derived linoleic acid, 13-HODE, 9-HODE, 13HpODE and 9-HpODE were determined simultaneously by HPLC with diode array detection (Browne & Armstrong 2000). The HPLC instrumentation consisted of a Shimadzu LC-10Avp pump, FCV 10 flow control valve, DGU14 degassing unit, SPD-M10A photodiode array, SIL-7A autosampler and a PC equipped with EZ Start chromatography data software (Shimadzu Scientific Instruments, Inc., Columbia, MD, USA). All solvents were HPLC grade from Fischer Scientific (Fair Lawn, NJ, USA). Linoleic acid, 13-HpODE, 9-HpODE, 13-HODE and 9-HODE, and 5-hydroxy-eicosahexaenoic acid methyl ester (internal standard) were purchased from Cayman Chemical Co. (Ann Arbor, MI, USA). For analysis, 150 μL of the prepared sample was injected onto a SupelcoSil LC-18 (25 cm $\times$ 4.6 mm I.D., 5 μm particle size, ODS column (Supelco, Inc., Bellefonte, PA, USA) and eluted isocratically with a mobile phase of acetonitrile/0.1% acetic acid/tetrahydrofuran (41:41:18 v/v/v) at a flow rate of 1.3 mL min^{-1} . The column eluent was monitored by the photodiode array from 200–300 nm.

Thiobarbituric acid reacting substances—TBARS were measured using OxiTech reagent kits from ZeptoMetrix Corp. (Buffalo, NY, USA) and expressed in nmol mL^{-1} MDA equivalents (Armstrong & Browne 1994). TBARS pigment was measured at excitation of 535

nm and emission of 552 nm on a RF-5000U spectrofluorometer (Shimadzu Scientific Instruments Inc.).

Antioxidant enzymes—Kinetic enzyme assays were adapted to the Cobas Fara II autoanalyzer (Roche Diagnostic Systems, Inc., Basel, Switzerland) (Pippenger et al. 1998). Plasma glutathione peroxidase (GPx), erythrocyte glutathione peroxidase (EGPx) and erythrocyte glutathione reductase (EGSHR) were performed using OxiTek reagent kits from ZeptoMetrix (Buffalo, NY, USA). Erythrocyte superoxide dismutase (ESOD) activity was determined by the inhibition of the oxidation of cytochrome C by xanthine/xanthine oxidase under standard conditions. One unit of SOD activity was defined as the amount of enzyme which induced a 50% inhibition of the reaction. Calibration curves of % inhibition vs SOD activity were generated using purified SOD. Erythrocyte enzyme activities were normalized per gram or milligram of haemoglobin (Hb). The activity of human serum paraoxonase was measured as both arylesterase activity (PONa) and paraoxonase activity (PONp) (Browne et al. 2007). Diethyl *p*-nitrophenyl phosphate (paraoxon), 98.0%, was obtained from Chem Service Inc. (Westchester, PA, USA). PONa activity was measured as the rate of formation of phenol using 4 mmol l⁻¹ phenylacetate in 20 mM TRIS-HCl, pH 8.0, with 1.0 mM CaCl₂. PONp activity was determined by the rate of formation of *p*-nitrophenol using 1 mmol l⁻¹ paraoxon as the substrate in 50 mmol l⁻¹ glycine buffer, pH 10.5, with 1.0 mmol l⁻¹ CaCl₂. Water blanks were used to correct for non-enzymatic hydrolysis.

Fat-soluble antioxidant vitamin profiles—Retinol, vitamin E (α , γ and δ tocopherol) and carotenoids (β -carotene, β -cryptoxanthin, lycopene and lutein/zeaxanthin) were measured in serum using HPLC (Browne & Armstrong 1998). Vitamin-A-alcohol (>95% *cis*-retinol) was purchased from Fluka Chemical Co. (Milwaukee, WI, USA). Lutein, zeaxanthin and β -cryptoxanthin were purchased from Indofine Pure Biochemicals (Hillsborough, NJ, USA). Results are expressed in $\mu\text{g ml}^{-1}$. Lutein and zeaxanthin were quantified together as a single peak and reported as a total lutein/zeaxanthin.

Total vitamin C—Total ascorbic acid was determined by the dinitrophenylhydrazine (DNPH) method (Chalmers & McWhinney 1986). Samples for analysis were stabilized immediately following phlebotomy and centrifugation by adding 0.5 ml of heparin plasma to 2.0 ml of 6% MPA and centrifuging at 3000g for 10 min. Clear supernatant was decanted and frozen at -80°C for analysis. The absorbance of each DNPH derivatized sample was determined at 520 nm on a Shimadzu 160U spectrophotometer (Shimadzu Scientific Instruments, Inc.).

Quality control—As quality control (QC) material is not commercially available for many of the assays, 'in-house' controls were generated. These specimens were used to estimate method performance characteristics and assess imprecision. Intra-assay reproducibility was calculated from 20 determinations in the same run. Long-term interassay reproducibility and control ranges were generated by running five samples per day over a period of 20 days. The mean $\pm$ two standard deviations was then calculated as the acceptable range for control materials which were analyzed in duplicate with all patient determinations. Performance was monitored using Levy-Jennings quality control plots. Daily quality and analytical run acceptability were evaluated according to Westgard multirule charts (Westgard et al. 1974). Day to day reproducibility was evaluated by direct limit Cusum Plots (Westgard et al. 1977). Fat-soluble vitamin control specimens consisted of 500 μl aliquots of a serum pool stored at -80°C in amber, 1.0 ml cryovials. Standard reference material (SRM) 968C1 and 968C2 from the National Institute of Standards and Technology (NIST) were included in all analytical runs, providing three-level quality control. Further laboratory quality assurance was maintained

through participation in the Micronutrient Measurement Quality Assurance Program round-robin proficiency testing from NIST.

For HPLC lipid peroxidation profiles, TBARS and PGPx control specimens consisted of aliquots of an EDTA plasma pool stored at -80°C in amber, 1.0 ml cryovials in the presence of 10 μM indomethacin and 0.005% butylated hydroxyl toluene as antioxidant preservatives. Total ascorbic acid control specimens were prepared by adding dithiothreitol to heparinized plasma. A final concentration of 1 g l^{-1} has been shown to prevent the oxidative loss of vitamin C and samples are stable for at least 1 year at -80°C (Margolis & Duewer 1996). Antioxidant enzyme control material was provided by ZeptoMetrix Corp. as lyophilized, human blood plasma and human erythrocyte lysates.

Statistical analysis

Statistical analyses were conducted using SAS version 9.01 (SAS Institute Inc., Cary, NC, USA). An unabridged statistical description is included as a supplemental file. Distributions were considered for each of the 21 analytes, stratified by day and log-transformed where this improved the approximation to a normal. Two-way nested random effects ANOVA was used to describe the single menstrual cycle variability for each analyte. An exchangeable correlation matrix, which assumes constant correlation coefficient between any two error terms within the same subject, was specified for error terms to accommodate the clustered nature of the data. Models were specified as:

$$Y_{ijk} = \mu + (sub)_i + (day)_{j(i)} + e_{k(ji)}$$

where Y_{ijk} described the analyte value for the k th replicate ($k = 1, 2$), on the j th day ($j = 1, 2, 3, \dots, 8$), for the i th subject ($i = 1, 2, 3, \dots, 9$); μ the grand mean for the analyte; $(sub)_i$ the random effect of the i th subject; $(day)_{j(i)}$ the random effect of the j th day nested in the i th subject; and $e_{k(ji)}$ the k th replicate nested in the j th day and the i th subject (Littell et al. 1998). Random effects and errors were assumed to be normally distributed with mean zero and a constant variance σ^2 .

The proportion of the total variance attributable to each effect was estimated by dividing the components of variance by the total variance where σ_B^2 corresponds to $(sub)_i$, σ_W^2 to $(day)_{j(i)}$ and σ_A^2 to $e_{k(ji)}$. The total variance for an analyte was defined as $\sigma_T^2 = \sigma_B^2 + \sigma_W^2 + \sigma_A^2$. The index of individuality (I_I), was calculated as $\sqrt{(\sigma_W^2 + \sigma_A^2) / \sigma_B^2}$ (Fraser & Harris 1989). Coefficients of variation (CVs) for each analyte were calculated from the components of variance specified as $CV_I = \sqrt{\sigma_I^2} / x_I$; where σ_I^2 represents a variance component and x_I an analyte mean. The analytical acceptability index (A_I), was calculated as CV_A / CV_W where <0.25 is optimal performance, <0.5 is desirable performance and <0.75 is minimal performance (Cotlove et al. 1970).

Homogeneity of within-subject variance, across subjects, was evaluated using the index of heterogeneity (I_H) defined as $CV_W + CV_A / (\sqrt{2/(n-1)}) \times 100$, with rejection of the homogeneity assumption where $I_H > 1.35$. The critical difference between serial results (CD), was estimated as $2.77 \times \sqrt{(CV_W)^2 + (CV_A)^2}$ under the homogeneity of variances assumption (Fraser & Harris 1989). The minimum number of specimens, K , required to reduce the index of individuality to that observed for PONp (i.e. smallest observed in the current study) was defined as $K >$

$(I_{\text{Mean}} / I_{\text{PONp}})^2$. The intraclass correlation coefficient (ICC), defined as $\sigma_B^2 / (\sigma_B^2 + \sigma_A^2)$, describes the proportion of variability in observed values due to biological variation (Lachin 2004) across the range of measured values.

Results

The demographic characteristics of the study participants are listed in Table I. The distribution and number of measurements available for each marker are shown in Table II. The maximum number of specimens ($9 \text{ participants} \times 8 \text{ days} \times 2 \text{ replicates} = 144$) were available for vitamin C (Vit C), pGPx and TBARS. For HPLC vitamins 142 replicates were available due to two HPLC autosampler instrumental errors (one on day 7 for one subject and day 27 for another). Two replicates were similarly missing for linoleic acid and lipid peroxide analyses (two participants on day 12). Among the HPLC analytes, 14 samples from two participants had visibly detectable analyte peaks which were below the limit of quantification (set by chromatographic peak width and threshold). In these cases, values were generated via manual forced peak integration (11 values for 9-HODE, two for 9-HpODE and one for 13-HpODE only). An identical procedure was employed for the δ -tocopherol measures with values below the limit of quantification. This strategy preserved sample size while avoiding the effect of censoring (Schisterman et al. 2006). One participant had both replicates unavailable for the day 7 serum AOX enzymes analyses and another had both replicates unavailable for the day 2 erythrocyte AOX enzymes analyses.

Analytical performance

The QC performance for all assays is shown in Table III and list the mean, within-run and between-run CV. Markers are divided into groups based on the QC material that was used. For fat-soluble vitamins, all within- and between-run CVs were below 10.0% with the exception of δ -tocopherol where levels of the 'in-house' and SRM 968C1 QC materials were below the limit of quantification of our system for this analyte ($0.005 \mu\text{g mL}^{-1}$). Lipid and lipid peroxide imprecision ranged from 4.7 to 11.2%. Although measured simultaneously, lower imprecision was noted for linoleic acid owing to inherently higher concentration. Plasma TBARS demonstrated imprecision between 8.0 and 9.0%. Serum PONA and PONP determinations demonstrated CVs $<2.0\%$ across a broad range of enzyme activity as did PGPx with CVs $<5.0\%$. Erythrocyte antioxidant enzymes eSOD, eGPx and eGSHR demonstrated considerably higher between-run relative to within-run imprecision. In-house QC materials demonstrated no statistically significant changes over the study period and are in our experience are stable for at least 18 months when stored at -80°C (data not shown).

Biological variability

The mean and standard deviation of variable levels for the study relative to the mean and standard deviation of levels for each participant are depicted in Appendix Figures 1-3. Prior to analysis, a log transformation was applied to α -tocopherol, β -cryptoxanthin, β -carotene, lycopene, Vit-C, TBARS, 13-HpODE, linolenic and linoleic acids, SOD, GSHR to improve approximation to a normal distribution. Only six of nine study participants had levels of δ -tocopherol in the quantifiable range limiting variability estimates for this micronutrient. The components of variance, the ICC among replicate analyses and the calculated values for analytical, intra-individual and inter-individual%CVs are shown in Table IV. In general, analytical variability was small relative to both intra- and inter-individual variability, comprising less than 10% of total variability for all analytes except 13-HpODE which was 12.35%. The ICCs are greater than 0.9 for all analytes except δ -tocopherol which was 0.76.

Table V shows the calculated parameters of biological variation A_I , I_I , k , I_H , and CD. A generally held criterion sets an A_I threshold <0.5 ; however, performance of each analyte was evaluated according to ranks such that $A_I < 0.25$ was 'optimal', $A_I < 0.5$ was 'desirable', $A_I < 0.75$ was minimal, and $A_I \geq 0.75$ was insufficient (Fraser et al. 1997). All study variables achieved at least minimal analytical acceptability except for linoleic acid which failed to achieve at least minimal performance. Substantial variation in the individuality of analytes (i.e. I_I) was

observed, ranging from 0.24 for PONp to 3.52 for SOD. The smaller the I_I , the greater the individuality and repeated measurements give little additional information. This is evidenced in the number of measurements needed to estimate the homeostatic set point (k) which correlates strongly with I_I and ranges from 1 (retinol, β -cryptoxanthin and PONp) to nearly 15 (eSOD) measurements required to achieve a 95% chance that the mean is within $\pm 5\%$ of the true value. Five variables (δ -tocopherol, 13-HODE, 13-HpODE and 9-HpODE) failed to meet homogeneity assumptions demonstrating μH values greater than 1.35 while 9-HODE was borderline at 1.28. The range of critical difference values varies greatly from less than 1% to greater than 200% between two successive serial values. The largest CD values are associated with the 13-HODE, 9-HODE, 13-HpODE and 9-HpODE where large, heterogeneous intra-individual variability results make it difficult to assess the significance of changes in serial results.

No statistically significant differences, at $\alpha = 0.05$, in temporal effects across the menstrual cycle (i.e. 'day'), were observed. The lack of findings may be due to the small sample size of the current study which may have insufficient power to detect significant differences over the time period studied. A larger study should be done to evaluate these possible effects.

Discussion

Data on biological variation of biomarkers associated with OS are generally lacking, in relation to the menstrual cycle in particular. Limited studies have been conducted to address these methodological issues (Kadiiska et al. 2000, 2005a,b, Kato et al. 2006) and there is still controversy over which biomarkers to use. It has been suggested that multiple biomarkers of OS should be utilized for research purposes until more sensitive and specific assays are developed (Chait 1994). The current consensus in laboratory medicine is that quality specifications should be based on the parameters of biological variation (Fraser & Hyltoft Petersen 1999, Hyltoft Petersen et al. 2002). In this study we generated biological variability data for a panel of biomarkers that are commonly measured in the laboratory when assessing OS. Using these data on components of biological variation, we have derived desirable quality specifications, indexed the usefulness of population-based reference ranges and estimated the level of changes in serial results obtained in an individual needed to find statistical significance. Although the number of participants investigated was small, a comparison of studies on biological variation demonstrates that estimates of intra-individual and inter-individual variation are similar regardless of the number of individuals studied (Fraser 1992, Sebastian-Gimbaro 1997, Ricos et al. 2004).

Lipid peroxidation (LPO) is a prominent manifestation of oxidative damage and a number of markers of this process have been developed. We chose to estimate MDA by TBARS reaction. MDA by highly specific GC-NICI-MS has been advocated as a useful marker of lipid peroxidation (Hyltoft Petersen et al. 2002); however, the instrumentation is costly and not widely available. Indirect estimation of MDA can be made by reaction with thiobarbituric acid which forms a fluorescent adduct with MDA. Although criticized for its specificity it is still the most commonly employed biomarker of LPO because of its simplicity (Kadiiska et al. 2005b) and has been shown to correlate well with GC-MS (Liu et al. 1997). Despite its wide application this is, to our knowledge, the first attempt to characterize the components of variability of this assay during the menstrual cycle. One previous study of a reference sample group (Nielsen et al. 1997) estimated reference limits for the group to be between 0.36 and 1.24 $\mu\text{M l}^{-1}$ with intra-individual CVs of 6–30%. The analytical performance of this assay is minimally acceptable ($\text{CV} \approx 8\%$ and $A_I = 0.54$) and has a high degree of intra-individual variability necessitating replicate analysis to estimate the true plasma level which must nearly double to detect significantly different serial results.

In addition to TBARS we have employed an HPLC technique for the direct determination of linoleic acid peroxidation. Linoleic acid (LA; octadecadienoic acid, ODE) is the most abundant PUFAs in human blood plasma and its LPO products predominate (Spiteller & Spiteller 1997, Browne & Armstrong 2000). HpODEs are rapidly reduced in biological surroundings to corresponding hydroxy acids (HODEs) which are secondary products of LPO but are significantly more stable than HpODEs (Browne & Armstrong 2000). Linoleic acid peroxidation products are of high physiological relevance and in some diseases characterized by inflammation or cell injury are present at 10–100 times higher concentration compared with healthy individuals and been advocated to be excellent biomarkers of oxidative stress (Spiteller & Spiteller 1997).

The analytical performance of the HPLC measurement of each HODE and HpODE was at least analytically desirable ($A_1 < 0.48$) and demonstrated a CV of less than 5% based on external QC material. Our estimates of intra-individual variation for lipid peroxidation products range from 26 to 77% which is not largely dissimilar to those reported for F2 isoprostanes of about 42% (Helmerson & Basu 2001) indicating that there is considerable day-to-day biological variation in LPO. Factors that might cause this variability include differences in the formation and metabolism of these compounds depending on various endogenous regulating factors, i.e. enzymes and β -oxidation systems. Other reasons could be variability of endogenous defence mechanisms against OS and inflammation.

There is limited information with regard to fluctuations in antioxidant enzyme in the context of menstrual cycle with the most consistent finding has been increased pGPx and eGPx activity with increasing oestrogen levels (Massafra et al. 1998, 2000, Serviddio et al. 2002, Ha & Smith 2003). Antioxidant enzymes play an important role in inactivation and removal of toxic ROS. Besides individual differences in gene expression, antioxidant enzyme activities depend on other endogenous and/or environmental factors since within-subject variation is greater than analytical imprecision. The antioxidant activities in erythrocytes and plasma do not necessarily reflect levels of antioxidant defence in the whole organism since antioxidant enzyme activities vary among tissues.

Our results indicate that for PON1, pGPx, eGPx and GSHR, variation between individuals are generally larger than variation within-subject, suggesting enzyme activities in blood may reflect differences in antioxidant defence. The analytical performance for all assays is at least desirable, the analytical imprecision is less than 5%, and our estimate of GPx, GSHR and SOD within-subject variation is comparable to previous reports (Guemouri et al. 1991, Andersen et al. 1997). A previous estimate of the biological variability in PON 1 paraoxonase and arylesterase activities, during a 4-week time period in a larger and more diverse group of men and women, demonstrated comparable analytical variance but smaller intra-individual and larger inter-individual variance than those reported here (Browne et al. 2007).

The components of variation for fat-soluble vitamins in plasma reported in the literature (Talwar et al. 2005) are similar to our values, suggesting that our measurements represent reliable estimates of biological variation. Furthermore, there is no indication of increased variability of these vitamins during the menstrual cycle in our study. Other reports describing the variation of carotenoid and tocopherols levels during the menstrual have produced conflicting results. With regard to carotenoids, Rock et al. (1995) reported that α -carotene and lutein were increased in the mid-leuteal phase and the early follicular phase respectively, in young women, only if uncorrected for total cholesterol. Other carotenoids did not vary across the menstrual cycle, whether corrected or uncorrected for total cholesterol. Tangney et al. (1991) and Palan et al. (2006) also found no variation in carotenoids across the menstrual cycle. In contrast, in a controlled dietary study, individual plasma carotenoid concentrations increased from 8 to 29% at different phases of the menstrual cycle compared with menses (Forman et al.

1996). With regard to tocopherols, several reports have demonstrated a non-significant fluctuation in plasma α -tocopherol levels during the menstrual cycle. Lanza et al. (1998) reported 12% lower levels at menses than during luteal phase, Reimer et al. (2005) reported 12% lower levels during follicular phase compared with the luteal phase and Palan et al. (2006) observed 28% lower levels in the follicular versus luteal phase. It should be noted however that these observations were made under controlled dietary conditions which were not a component of several studies failing to demonstrate menstrual cycle variation and may explain the inability of the present study to detect these changes. Finally, it has been suggested that concentrations of these fat-soluble vitamins vary depending on the amount of lipid present and should be standardized per amount of cholesterol or total lipid. In at least one report this normalization did not affect the parameters of biological variation (Maes et al. 1996). Since a major goal of the present study was to characterize analytical variability, we did not standardize fat-soluble vitamin levels to cholesterol in an effort to avoid the introduction of bias (Schisterman et al. 2005) and additional analytical and biological variability inherent to this procedure.

Appendix

Graphs of the grand mean and standard deviation of study variable levels for all study subjects (All) and the mean and standard deviation of levels for each of nine individual subjects (1–9).

1. .

PONa, PON1 arylesterase activity; PONp, PON1 paraoxonase activity; EGSHR, erythrocyte glutathione reductase activity; PGPx, plasma glutathione peroxidase activity; EGPx, erythrocyte glutathione peroxidase activity and ESOD, erythrocyte superoxide dismutase activity; TBARS, thiobarbituric acid reactive substances; 9-HODE, 9 hydroxy octadecadieneoic acid; 13-HODE, 13 hydroxy octadecadieneoic acid; 13-HpODE, 13 hydroperoxy octadecadieneoic acid and 9-HpODE, 9 hydroperoxy octadecadieneoic acid.

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Table I

Characteristics of the study participants (n = 9).

Characteristic	
Age (years)	
Mean	38.5±4.7
Min–max	30.9–44.8
Cycle length (days)	
Mean	28.4±1.3
Min–max	26–30
Height (cm)	
Mean	165.1±8.9
Min–max	149.2–179.5
Weight (kg)	
Mean	71.1±18.5
Min–max	53.7–111.9
Body mass index (kg cm ⁻²)	
Mean	25.9±5.1
Min–max	20.9–34.7
Race, <i>n</i> (%)	
White	8 (88.9)
African-American	1 (11.1)
Education	
High school graduate	0
College, no degree	2 (22.2)
Associate's degree	3 (33.3)
Bachelor's degree	4 (44.4)

Table II

Number of repeated measurements available for each marker in nine study participants.

	Total replicates	Mean value	Range	
			Min	Max
Vitamins and micronutrients				
Retinol (μg ml ⁻¹)	142	0.39	0.230	0.530
δ-Tocopherol (μg ml ⁻¹)	142	0.03	NQ	0.172
γ-Tocopherol (μg ml ⁻¹)	142	2.17	0.76	3.2
α-Tocopherol (μg ml ⁻¹)	142	10.69	7.06	17.01
Lutein (μg ml ⁻¹)	142	0.14	0.07	0.20
β-Cryptoxanthin (μg ml ⁻¹)	142	0.11	0.044	0.261
β-Carotene (μg ml ⁻¹)	142	0.28	0.079	0.897
Lycopene (μg ml ⁻¹)	142	0.59	0.312	1.131
Vitamin C (mg dl ⁻¹)	144	1.97	1.047	3.344
Lipids and lipid peroxides				
TBARS (μmol l ⁻¹)	144	1.46	1.061	2.096
13-HODE (μmol l ⁻¹)	142	0.15	0.034	0.561
9-HODE (μmol l ⁻¹)	142	0.16	NQ	0.413
13-HpODE (μmol l ⁻¹)	142	0.26	NQ	1.507
9-HpODE (μmol l ⁻¹)	142	0.35	NQ	2.199
Linoleic (μg ml ⁻¹)	142	233.7	107.7	350.1
Plasma AOX enzymes				
pGPx (U l ⁻¹)	144	850.36	616.4	1130.5
Serum AOX enzymes				
PONa (kU l ⁻¹)	140	69.28	51.4	249.3
PONp (U l ⁻¹)	140	294.21	117	711
Erythrocyte AOX enzymes				
eSOD (U g ⁻¹ Hgb)	140	7.93	5409	14816
eGSHR (U g ⁻¹ Hgb)	140	7.53	4.92	12.40
eGPx (U g ⁻¹ Hgb)	140	57.8	35.8	99.0

PONa, PON1 arylesterase activity; PONp, PON1 paraoxonase activity; eGSHR, erythrocyte glutathione reductase activity; pGPx, plasma glutathione peroxidase activity; eGPx, erythrocyte glutathione peroxidase activity; eSOD, erythrocyte superoxide dismutase activity; TBARS, thiobarbituric acid reactive substances; 9-HODE, 9 hydroxy octadecadieneoic acid; 13-HODE, 13 hydroxy octadecadieneoic acid; 13-HpODE, 13 hydroperoxy octadecadieneoic acid; 9-HpODE, 9 hydroperoxy octadecadieneoic acid.

NQ, not quantified (below limit of quantification).

Table III
Study variable means, within-run and between-run imprecision based on quality control materials expressed as percentage coefficient of variation (%CV).

	Mean	Within% CV (between%CV)	Mean	Within% CV (between%CV)	Mean	Within% CV (between%CV)
Vitamins and micronutrients	SRM 968 C1		SRM 968 C2		In-house QC	
Retinol ($\mu\text{g ml}^{-1}$)	0.74	3.01 (6.09)	0.44	2.96 (5.93)	0.52	6.17 (6.03)
δ -Tocopherol ($\mu\text{g ml}^{-1}$)	0.05	21.23 (40.78)	0.58	8.85 (14.27)	0.03	71.85 (61.90)
γ -Tocopherol ($\mu\text{g ml}^{-1}$)	3.71	2.15 (5.49)	1.65	2.29 (5.26)	1.96	4.55 (7.54)
α -Tocopherol ($\mu\text{g ml}^{-1}$)	6.88	2.34 (4.49)	14.96	2.51 (5.52)	13.52	4.99 (5.83)
Lutein ($\mu\text{g ml}^{-1}$)	0.07	5.55 (5.93)	0.08	5.00 (9.16)	0.11	6.55 (7.75)
β -Cryptoxanthin ($\mu\text{g ml}^{-1}$)	0.06	2.89 (8.00)	0.03	9.68 (9.69)	0.07	5.95 (9.13)
β -Carotene ($\mu\text{g ml}^{-1}$)	0.14	6.52 (7.53)	0.35	6.66 (8.74)	0.17	5.27 (9.86)
Lycopene ($\mu\text{g ml}^{-1}$)	0.35	5.59 (8.25)	0.48	5.99 (8.72)	0.43	7.63 (9.02)
Vitamin C* (mg dl^{-1})					Heparin/DTT plasma	6.88 (9.60)
Lipids and lipid peroxides	EDTA plasma QC1		EDTA plasma QC2			
TBARS ($\mu\text{mol l}^{-1}$)	2.0	8.10 (8.29)	1.10	8.20 (8.90)		
13-HODE ($\mu\text{mol l}^{-1}$)	0.105	9.20 (11.20)				
9-HODE ($\mu\text{mol l}^{-1}$)	0.120	9.00 (9.50)				
13-HpODE ($\mu\text{mol l}^{-1}$)	0.037	8.00 (10.50)				
9-HpODE ($\mu\text{mol l}^{-1}$)	0.053	10.40 (10.30)				
Linoleic ($\mu\text{g ml}^{-1}$)	225	4.70 (4.40)				
Plasma antioxidant enzymes	EDTA plasma QC1		EDTA plasma QC2			
pGPx (U l^{-1})	474	3.30 (4.58)				
Serum antioxidant enzymes	Serum QC1		Serum QC2			
PONa (kU l^{-1})	106.7	1.20 (1.20)	124.3	0.90 (1.10)		
PONp (U l^{-1})	115.1	1.40 (1.60)	457.2	1.20 (1.30)		
Erythrocyte antioxidant enzymes	Erythrocyte lysate QC					
eSOD ($\text{U mg}^{-1} \text{Hgb}$)	4.74	4.64 (8.02)				
eGSHR ($\text{U g}^{-1} \text{Hgb}$)	3.57	3.65 (10.64)				
eGPx ($\text{U g}^{-1} \text{Hgb}$)	36.38	5.01 (7.92)				

For definitions see footnote for Table II.

Table IV

Components of variance, ICCs among replicate analyses, and calculated values for inter-individual (B), intra-individual (W), and analytical (A) coefficients of variation (CV) for study analytes.

Analyte	%S ² _B	%S ² _W	%S ² _A	ICC	95%Io	95%Hi	CV _B	CV _W	CV _A
Vitamins and micronutrients									
Retinol	90.96	6.11	2.93	0.98	0.95	0.98	21.26	5.51	3.82
δ-Tocopherol	16.50	78.33	5.17	0.76	0.65	0.84	52.01	113.31	29.11
γ-Tocopherol	67.62	31.58	3.00	0.99	0.98	0.99	27.59	18.85	3.00
α-Tocopherol	73.36	24.33	2.31	0.97	0.95	0.98	6.65	3.83	1.18
Lutein	80.71	16.60	2.70	0.97	0.95	0.98	22.35	10.14	4.09
β-Cryptoxanthine*	92.72	6.84	0.44	1.00	0.99	1.00	19.73	5.36	1.36
β-Carotene*	93.52	4.18	2.30	0.98	0.96	0.98	48.36	10.23	7.58
Lycopene*	67.93	27.93	4.14	0.94	0.91	0.96	47.90	30.72	11.82
Vitamin C	51.31	44.09	4.59	0.92	0.87	0.95	25.75	23.87	7.71
Lipids and lipid peroxides									
TBARS	49.83	38.80	11.37	0.81	0.72	0.88	30.19	26.64	14.42
13-HODE	14.23	84.43	1.34	0.91	0.87	0.95	21.95	53.45	6.73
9-HODE [†]	38.38	60.56	1.07	0.97	0.96	0.98	32.76	41.16	5.46
13-HPODE ^{*,†}	33.49	54.16	12.35	0.97	0.96	0.98	31.92	40.59	19.38
9-HPODE [†]	34.14	64.39	1.47	0.96	0.93	0.97	56.14	77.10	11.67
Linoleic acid*	88.60	5.45	5.95	0.94	0.90	0.96	0.13	0.03	0.03
Plasma AOX enzymes									
PGPX	70.38	26.23	3.39	0.95	0.93	0.97	11.45	6.99	2.51
Serum AOX enzymes									
PONa*	82.80	15.35	1.85	0.98	0.95	0.98	33.61	14.47	5.03
PONp	94.44	4.56	1.00	0.99	0.98	0.99	9.33	2.05	0.96
Erythrocyte AOX enzymes									
eSOD	7.49	91.82	0.69	0.92	0.87	0.95	0.61	2.14	0.19
EGSHR*	45.48	48.43	6.09	0.88	0.82	0.92	6.22	6.42	2.27
eGPX	80.05	16.85	3.10	0.96	0.98	0.98	5.25	2.41	1.03

* Log-transformed to improve the approximation to a normal.

[†] Values reported below the limit of quantitation. For definitions see footnote for Table II.

Table V

The calculated parameters of biological variation including the analytical acceptability index (A_I), index of individuality (I_I), index of heterogeneity (I_H) and the critical difference between serial results (CD).

Analyte	A_I	I_I	I_H	Analytical performance	CD	k
Vitamins and micronutrients						
Retinol	0.69	0.32	0.26	Minimum	18.565	1.3
δ -Tocopherol	0.26	2.25	3.90	Desirable	15.315	2.9
γ -Tocopherol	0.16	0.69	0.60	Optimum	52.878	9.3
α -Tocopherol	0.31	0.60	0.14	Desirable	18.565	2.5
Lutein	0.40	0.49	0.39	Desirable	30.272	2.0
β -Cryptoxanthine *	0.25	0.28	0.18	Desirable	35.263	1.2
β -Carotene	0.74	1.92	0.49	Minimum	2.459	7.9
Lycopene	0.38	0.69	1.17	Desirable	91.174	2.8
Vitamin C	0.32	0.97	0.87	Desirable	69.49	4.0
Lipids and lipid peroxides						
TBARS	0.54	1.00	1.12	Minimum	83.907	4.1
13-HODE	0.13	2.45	1.65	Optimum	149.230	3.7
9-HODE [†]	0.13	1.27	1.28	Optimum	115.008	5.2
13-HPODE ^{*,†}	0.48	1.41	1.64	Desirable	124.603	5.8
9-HPODE [†]	0.15	1.39	2.43	Optimum	216.004	5.7
Linoleic acid *	1.04	0.36	0.00	Insufficient	0.132	1.5
Plasma AOX enzymes						
pGPX	0.36	0.65	0.26	Desirable	20.573	2.7
Serum AOX enzymes						
PONa *	0.69	0.32	0.53	Minimum	42.437	1.9
PONp	0.47	0.24	0.08	Desirable	6.27	1.0
Erythrocyte AOX enzymes						
eSOD *	0.09	3.52	0.06	Optimum	5.961	14.5
eGSHR *	0.35	1.09	0.24	Desirable	18.854	4.5
eGPX *	0.43	0.50	0.09	Desirable	7.256	2.1

* Log-transformed to improve the approximation to a normal.

[†] Values reported below the limit of quantitation. For definitions see footnote for Table II.