

Daily Profiles of Steroid Hormones and Their Metabolites Related to Food Intake

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Summary

The aim of this study was to look for changes in the daily profile of steroid hormones after standardized food intake. Eight young women not taking contraceptives were followed from 5:30 a.m. till 9:30 p.m. before and 1 and 2 h after eating breakfast, snack, lunch, the second snack and dinner. The differences in steroid levels before and after meals were evaluated. As expected, glucose, C-peptide and ghrelin levels changed postprandially. The steroid hormones cortisol, progesterone, pregnenolone and dehydroepiandrosterone showed a decrease after main meals, whereas testosterone and dihydrotestosterone showed no significant dependence on food intake. Estrogen levels did not exhibit a significant nycthemeral rhythm, but estradiol decreased after main meals. In our study the known nycthemeral rhythm of LH, FSH, cortisol, progesterone and pregnenolone after food intake were confirmed, but significant changes after meals were also observed in the levels of cortisol, dehydroepiandrosterone, estradiol and SHBG.

Key words

Daily pattern • Food intake • Steroid hormones • Ghrelin • Orexin
• SHBG

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Introduction

A large amount of information has been accumulated on the time structure of the endocrine system. Circadian rhythms of physiological actions including metabolism and behavior are generated by central and peripheral circadian oscillators, which are tuned up by periodic environmental or physiological

stimuli. A master circadian pacemaker in the hypothalamic suprachiasmatic nucleus (SCN) is directly controlled by daily changes of light-dark cycles, and coordinates the timing of other oscillators by direct and indirect neuronal, hormonal and behavioral signals. The daily rhythm of food intake creates stimuli that calibrate most peripheral and central oscillators. Some of these could be involved in the daily rhythm of food anticipatory activity (Patton and Mistlberger 2013).

The highly coordinated output of the hypothalamic biological clock not only governs the daily rhythm in sleep/wake (or feeding/fasting) behavior but also has direct control over many aspects of hormone release. In fact, a significant proportion of our current understanding of the circadian clock has its roots in studies of the intimate connections between the hypothalamic clock and multiple endocrine axes (Kalsbeek and Fliers 2013).

Until now, most attention was paid to the main hormones such as melatonin (Vaněček 1998, Trivedi and Kumar 2014), cortisol (Dickmeis 2009, Chung *et al.* 2011, Son *et al.* 2011), testosterone (Karatsoreos *et al.* 2007), estradiol (Christian and Moenter 2010), the neurohormones oxytocin and vasopressin (Forsling 2000) and enterohormones or hormones related to food intake (Konturek *et al.* 2011). Only rare reports have dealt with other steroid hormones. Some data were published on dehydroepiandrosterone (Stárka *et al.* 2015) and its hydroxymetabolites, no data could we find for allopregnanolone or pregnenolone. However, in many situations, particularly for neuroactive metabolites, these steroids can extensively modulate many brain functions like mood or behavior. Reports on the effects of food intake, which could change the basal course of the daily hormone profile, are scanty.

Materials and Methods

Subjects

Eight women were involved in this study. They were 25-35 years old with a mean of 29.48 ± 2.99 years. Their BMI was normal, with a mean BMI of 21.3 ± 1.3 . All subjects were premenopausal, non-smokers, healthy and used no medications or hormonal contraception. All were in the follicular phase (days 1-7 of the menstrual cycle) at the time of study. The volunteers were kept in standardized conditions for five days (8 h of sleeping and periodic eating according to a standardized menu, which did not differ significantly from their normal daily activities and habits).

The Ethical Committee of the Institute of Endocrinology in Prague approved the protocol of the study, and written informed consent was obtained from all participants.

During the trial, all of the volunteers consumed similar food with a standard amount of carbohydrates, proteins and lipids, with a total caloric value of about 8400 kJ. The participants ate only meals according to the prepared menu and kept a time table. Common physical activity was allowed during the trial. The study lasted 16 h, from 5:30 a.m. to 9:30 p.m.

Before the first blood sampling the subjects were given an intravenous cannula (in the cubital or forearm vein). The first blood withdrawal was at 6 a.m. (30 min after awaking) after overnight fasting. The next withdrawals were always one and two hours after eating according to the following schedule (7:15, 8:15, 10:15, 11:15 a.m.; 1:15, 2:15, 4:15, 5:15, 7:15, 8:15 p.m.). The last blood withdrawal was 30 min before sleep, at 9:30 p.m. Breakfast (2 slice of bread, ham, cheese and tea) was served at 6:15, a snack (yogurt) at 9:15, lunch (beef broth soup, turkey, potato dumplings, and sauerkraut) at 12:15, an afternoon snack (apple) at 15:15, and dinner (tomato, 2 slices of bread, ham and cheese) at 18:15.

Analytical methods

Each sample was collected into a cooled plastic tube containing 100 µl of 5%EDTA. The plasma was obtained after centrifugation for 5 min at 2000 rpm at 4 °C. Plasma was separated and frozen within half an hour of being drawn from the subject, and was stored at -20 °C until analyzed.

C-peptide was measured in serum using ECLIA (electrochemiluminescence immunoassay, Modular E

170 analyser, Roche). The measuring range of the kit (defined by the lower detection limit and the maximum of the master curve) was 0.003-13.3 nmol/l or 0.01-40.0 ng/ml for plasma. Intra- and inter-assay coefficients of variation were 1.5 % and 2.3 %, respectively.

Blood glucose was measured using the enzymatic reference method with hexokinase (Cobas Integra 400 plus analyser, Roche). The measuring range of the kit was 0.12-40 mmol/l (2.16-720 mg/dl). Intra-set and inter-set reproducibility were 1.7 % and 2.6 %, respectively.

Total plasma ghrelin was determined by a commercially available RIA kit (Linco Research, Inc., St. Charles, Missouri, USA, detection range: 93-6000 pg/ml), and orexin A was also measured by a RIA kit (Phoenix Pharmaceuticals, Inc., detection range 10-1280 pg/ml).

Kits from Immunotech (France) were used to measure LH and FSH (IRMA kit). Cortisol was measured using RIA kit from Immunotech (France). Sex hormones binding globulin (SHBG) were measured by the IRMA kit Immunotech (France).

Steroids measured by GC/MS method

The levels of unconjugated steroids were measured in cubital vein blood using the GC-MS method (Hill *et al.* 2010). In brief, free steroids were extracted from plasma by diethyl-ether; steroid conjugates were hydrolyzed and extracted. The resulting residues were derivatized by methoxyamine hydrochloride and analyzed by GC/MS as follows.

Steroids were purchased from Steraloids (Newport, RI, USA), Sylon B from Supelco (Bellefonte, PA, USA), methoxylamine hydrochloride from Sigma (St. Louis, MO, USA) and solvents from Merck (Darmstadt, Germany).

Instruments

Measurements of steroid levels were done on a GCMS-QP2010 Plus system by Shimadzu (Kyoto, Japan) consisting of a gas chromatograph equipped with automatic flow control, an AOC-20s autosampler, and a single quadrupole detector with an adjustable electron voltage of 10-195 V. A capillary column with a medium polarity RESTEK RxI phase (diameter 0.25 mm, length 15 m, film thickness 0.1 µm) was used for analyses. Electron impact ionization with electron voltage fixed at 70 V and emission current set to 160 µA was used. The

temperatures of the injection port, ion source and interface were maintained at 220 °C, 300 °C, and 310 °C, respectively. Analyses were carried out in the splitless mode with a constant linear velocity of the carrier gas (He), which was maintained at 60 cm/s. The septum purge flow was set at 3 ml/min. The samples were injected using the high pressure mode (200 kPa), which was maintained for 1 min. The detector voltage was set to 1.4 kV.

Statistical data analysis

The changes of steroid levels were evaluated using a repeated measures ANOVA model consisting of a Subject factor, explaining differences between subjects, and a Stage factor. Due to the non-Gaussian data distribution and non-constant variance, the original data were transformed by a power transformation to attain a symmetric distribution of the data and residuals as well as homoscedasticity (Meloun *et al.* 2000). The homogeneity of the transformed data was checked by residual analysis as described elsewhere (Meloun *et al.* 2002, 2004). Then, the significance of the values before and after food intake was evaluated by least significant differences multiple comparisons.

Results

The majority of the hormones showed significant differences of their levels during the day between hours and especially between subjects (Table 1). Non-significant differences between hours were only found for orexin, testosterone, and dihydrotestosterone (Table 1, H-F), and non-significant differences between subjects (Table 1, S-F) were only found for orexin, glucose and C-peptide.

A significant increase of glucose and C-peptide was observed 1 and 2 h following the main meals lunch and dinner ($P=0.001$, Figs 1A,B; means and 95 % confidence intervals are given in all figures). The most striking marker of food intake was the increase of C-peptide (Fig. 1B), which was significant after breakfast, lunch and dinner.

Ghrelin levels had a significant minimum after lunch and dinner (Fig. 1C), whereas the differences in orexin levels over the course of the day were insignificant (Fig. 1D, Table 1). LH and FSH (Fig. 2A,B) showed a continuous decrease during the day, with no significant additional decrease after main

meals. Androgen levels (testosterone Fig. 3A, dihydrotestosterone Fig. 3B) did not show significant relationships to food intake, with the exception of an increase in dihydrotestosterone 2 h after lunch. Dehydroepiandrosterone (Fig. 3C) significantly decreased 1 and 2 h after lunch and breakfast respectively. SHBG (Fig. 5C) showed a small decrease after meals.

Table 1. Statistical characteristic of diurnal profiles of steroid hormones and other hormones/substances using repeated measures ANOVA with the factors Hour and Subject.

Substance	Hour		Subject	
	H-F	p	S-F	p
Ghrelin	3	0.0059	30.2	<0.0001
Orexin	0.6	0.7789	2.3	0.0769
Glucose	4	0.0006	1.9	0.1235
C-Peptide	12.4	<0.0001	2.7	0.0463
LH	3.1	0.0037	25.1	<0.0001
FSH	3.9	0.0007	19.1	<0.0001
SHBG	3.2	0.003	17.9	<0.0001
Cortisol	16	<0.0001	8.5	<0.0001
Progesterone	7.3	<0.0001	25.6	<0.0001
Pregnenolone	5.7	<0.0001	22.7	<0.0001
Testosterone	1.7	0.1236	47.8	<0.0001
DHT	1.9	0.0669	15.3	<0.0001
DHEA	11.42	<0.0001	12.4	<0.0001
Estrone	10	<0.0001	122.1	<0.0001
Estradiol	2.8	0.0101	41.5	<0.0001

H-F – Hour factor showing the differences between time periods, S-F – Subject factor showing the differences between individual subjects, p – significance.

All three C21 steroids (cortisol Fig. 4A, progesterone Fig. 4B and pregnenolone Fig. 4C) showed a significant decrease one hour after awakening and then a continuous decrease during the day. Cortisolemia decreased continuously throughout the day, and an additional significant decrease of cortisol related to food intake was only observed 2 h after lunch (Fig. 4A).

Estrogens (estrone Fig. 5A and estradiol Fig. 5B) did not show any striking trend in daily profiles; estradiol decreased significantly after lunch and dinner.

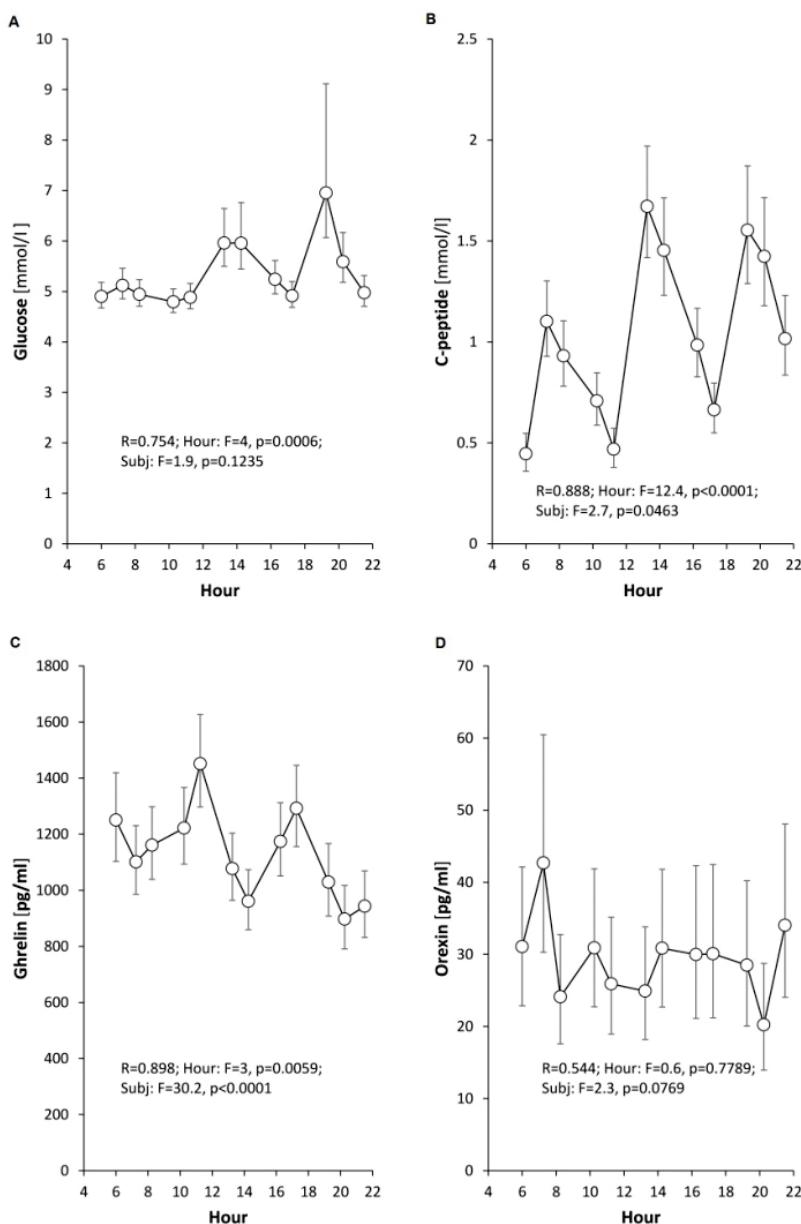


Fig. 1. Daily profiles of glucose (**A**), C-peptide (**B**), and the food-related hormones ghrelin (**C**) and orexin (**D**).

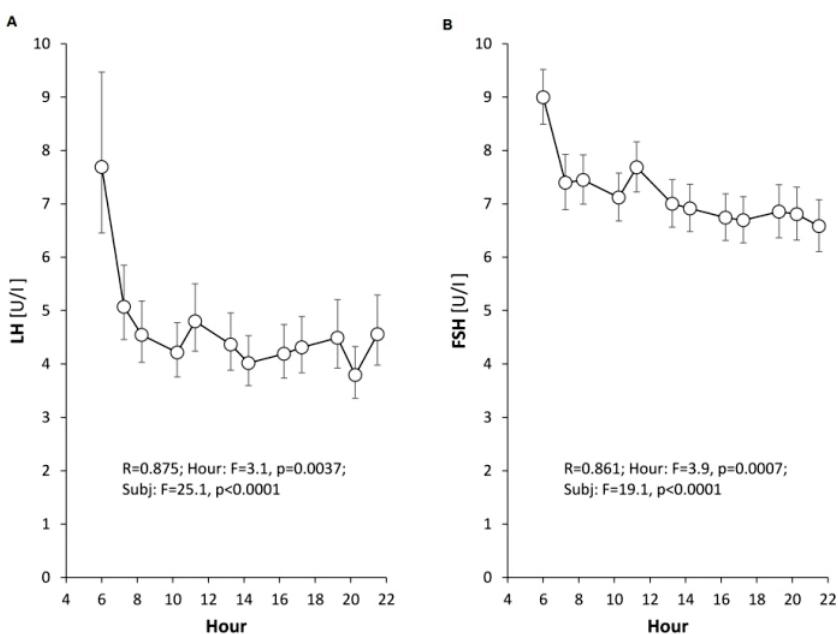


Fig. 2. Daily profiles of the gonadotropins LH (**A**) and FSH (**B**).

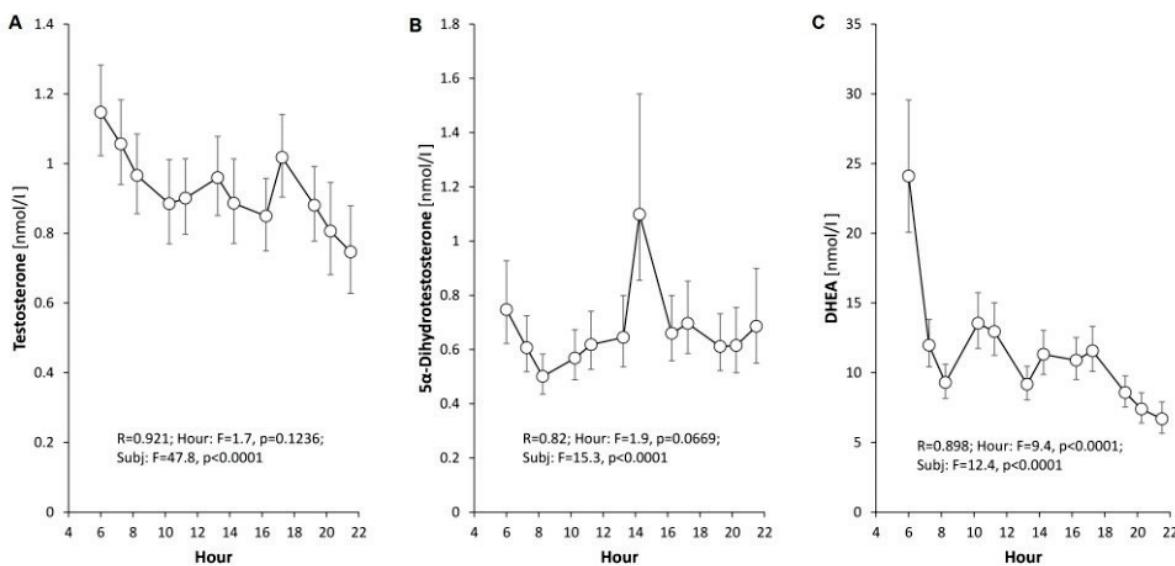


Fig. 3. Daily profiles of the androgens testosterone (A), dihydrotestosterone (B) and dehydroepiandrosterone (C).

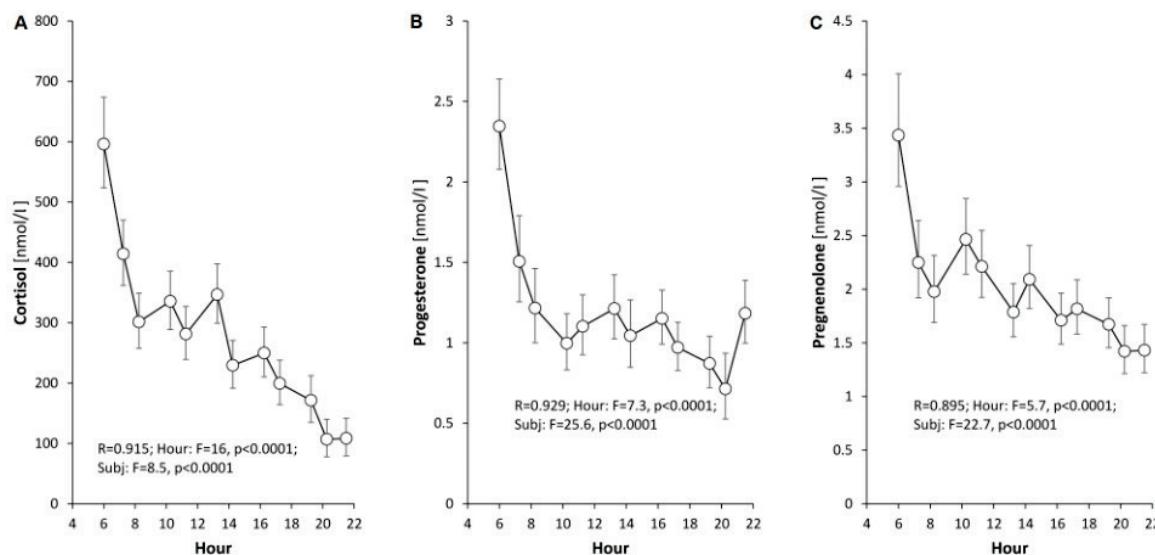


Fig. 4. Daily profiles of the C21 steroids cortisol (A), progesterone (B) and pregnenolone (C).

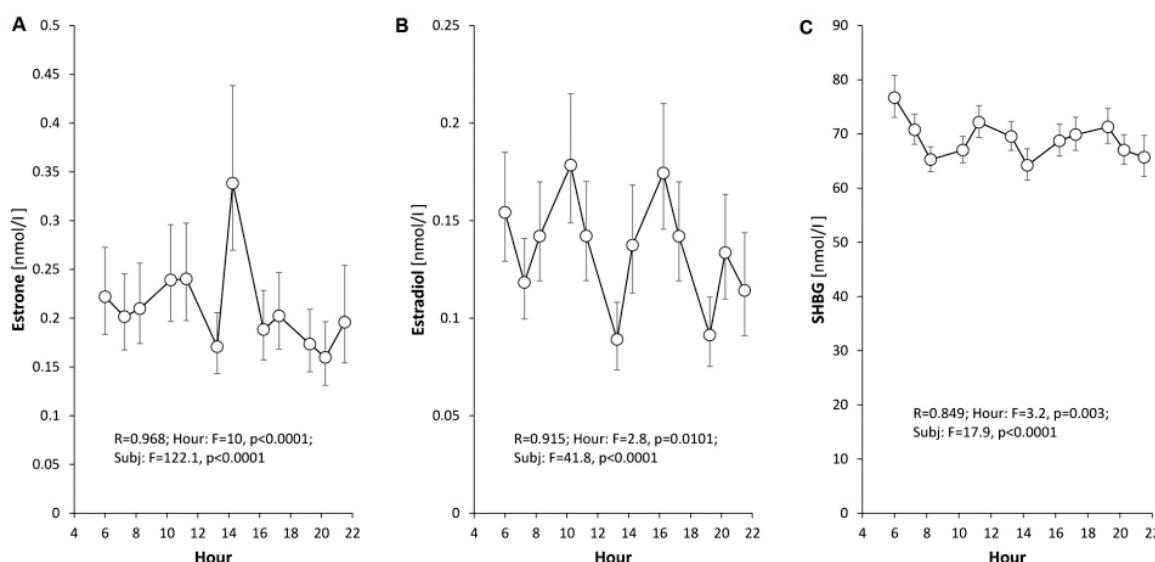


Fig. 5. Daily profiles of the estrogens estrone (A) and estradiol (B), and SHBG (C).

Discussion

Whereas the daily profiles of the main steroid hormones are well known, minor differences in the course of their levels related to defined and standardized food intake have not been studied systematically. In contrast to plenty of papers dealing with the effects of steroid hormones on eating behavior and disorders, the effect of food intake – with the exception of the effect of diet on endocrine homeostasis – has not yet received much attention in the literature. In animals, meal timing is a powerful synchronizer; however, there are no definitive and conclusive data to prove that meal timing is also a true synchronizer in humans, although there have been some reports suggesting this (Sensi *et al.* 1993). Meal timing was found to be an important factor determining the leptin diurnal rhythm, but other factors must contribute since the leptin response to a standard meal taken in the evening was greater than to the same meal taken in the morning (Elimam and Marcus 2002). The effects of two different feeding regimes on the 24-h profiles of maternal and fetal plasma cortisol and adrenocorticotropic hormone concentrations were studied in pregnant ewes (Simonetta *et al.* 1991). There were differences between the time of maximal glucose concentrations in maternal and fetal plasma in the once-daily-fed ewes in contrast to the multi-fed group. A significant diurnal variation in the plasma concentrations of cortisol was also present in the once-daily-fed ewes dependent on the stage of gravidity.

The timing of food intake is associated with

body weight regulation, insulin sensitivity and glucose tolerance, but the mechanisms are unknown. Bandín *et al.* (2015) found that eating late is associated not only with decreased resting-energy expenditures, decreased fasting carbohydrate oxidation and decreased glucose tolerance, but also smoothed out the daily profile of free cortisol concentrations.

In our study the known nycthemeral rhythm of LH, FSH, cortisol, dehydroepiandrosterone, progesterone and pregnenolone and decrease of ghrelin after food intake were confirmed, but significant changes after meals were also newly observed for the levels of cortisol, dehydroepiandrosterone, estradiol and SHBG. These effects were seen only after the main meals lunch and dinner. After breakfast, these effects are probably masked by the huge changes in the nycthemeral rhythm, and after snacks (yogurt and apples) the effects of food intake were either too small or the time period between eating the meal and the blood withdrawal was too long. It follows that for analytical determination of above mentioned hormones not only the time of blood withdrawals but also food consumption has an important influence upon the resulting values.

Conflict of Interest

There is no conflict of interest.

Acknowledgements

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