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Melatonin counteracts changes in hypothalamic gene expression of signals regulating feeding behavior in high-fat fed rats

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Short version of title: Melatonin and feeding behavior signals.

Abstract

Background: Previous studies indicate that the administration of melatonin caused body weight and abdominal visceral fat reductions in rodent models of hyperadiposity. The objective of the present study performed in high-fat fed rats was to evaluate the activity of melatonin on gene expression of some medial basal hypothalamus (MBH) signals involved in feeding behavior regulation, including neuropeptide Y (NPY), proopiomelanocortin (POMC), prolactin-releasing peptide (PrRP), leptin- and insulin-receptors (R) and insulin-R substrate (IRS)-1 and -2. Blood levels of leptin and adiponectin were also measured.

Methods: Adult Wistar male rats were divided into four groups (n= 16/group): (i) control diet (3 % fat); (ii) high-fat (35 %) diet; (iii) high-fat diet + melatonin; (iv) control diet + melatonin. Rats had free access to high-fat or control chow and one of the following drinking solutions: (a) tap water; (b) 25 µg/mL of melatonin.

Results: After 10 weeks, the high-fat fed rats showed augmented MBH mRNA levels of NPY, leptin-R, PrRP, insulin-R, IRS-1 and IRS-2. The concomitant administration of melatonin counteracted this increase. Feeding of rats with a high-fat diet augmented expression of MBH POMC gene through an effect insensitive to melatonin treatment. The augmented levels of circulating leptin and adiponectin seen in high-fat fed rats were counteracted by melatonin as was the augmented body weight: melatonin significantly attenuated body weight increase in high-fat fed rats without affecting chow or water consumption. Melatonin augmented plasma leptin and adiponectin in control rats.

Conclusions: The results indicate that an effect on gene expression of feeding behavior signals at the CNS may complement a peripheral rise of the energy expenditure produced by melatonin to decrease body weight in high-fat fed rats.

Key words: Melatonin; high-fat diet; food intake; gene expression; neuropeptide Y; proopiomelanocortin; prolactin-releasing peptide; leptin; adiponectin; medial basal hypothalamus.

Introduction

The possible involvement of melatonin in appetite regulation has been studied for years. Melatonin regulates food intake in rats [1], mice [2], hamsters [3], pigs [4], and in several submammalian species like goldfish [5], rainbow trout [6] and zebrafish [7,8]. In the rat the responses have been conflicting including reduction, increase or no effect on food consumption [9,10]. A decrease in fat mass and body weight has been reported in rats [10-13] whereas melatonin increases fat mass in gray mouse lemurs [14], Syrian hamster [3], raccoon dog [15] and garden dormouse [16].

Feeding behavior and regulation of nutrient intake are complex phenomena influenced by homeostatic, hedonic and peripheral signals [17,18]. The homeostatic pathway involves orexigenic neurons in the hypothalamic arcuate nucleus containing the orexigenic peptides neuropeptide Y (NPY) and agouti-gene related peptide (AgRP). These neurons project to second order neurons in the lateral hypothalamus which contain orexigenic peptides like orexin and melanin concentrating hormone. NPY and orexin increase appetite, arousal and spontaneous physical activity and therefore play an important role in energy balance providing a link among vigilance, energy balance and reward [17,18].

Another neuronal population in the arcuate nucleus encodes satiety and includes several anorexigenic peptides derived from the common precursor pro-opiomelanocortin (POMC), also called the “melanocortin” system, that co-exist with the cocaine and-amphetamine-regulated transcript (CART). Melanocortin and CART neurons project to satiety neurons located in the paraventricular nucleus which contain corticotropin-releasing hormone, a strong anorexigenic signal (for review see [19]).

This complex neuronal network assures a bidirectional homeostasis in response to peripheral signals reflecting the actual nutritional stage. Thus, satiety is produced by several factors like gastric dilatation, intestinal peptides liberated in postprandial state (cholecystokinin, glucagon-like peptide, peptide YY), metabolites as fatty acids and glucose and hormones as insulin and leptin [20]. Insulin is transported to the brain where it activates the anorexigenic neuronal pathways, causing a reduction in food intake. Leptin is another anorexigenic signal produced by the adipose tissue proportionally to the fat mass [21]. Plasma levels of insulin and leptin pass through the blood-brain barrier and interact with insulin receptors (insulin-R) and leptin receptors (leptin-R) that are expressed in first-order neurons of the arcuate nucleus, both in NPY/AgRP and POMC/CART neurons.

A number of studies in the literature indicate that the administration of melatonin to obese rats caused body weight and abdominal visceral fat reductions that were accompanied by very slight or no decrease in gross food intake [13,22-30]. The same was observed by us in a recent study evaluating the efficacy of melatonin to affect mild inflammation in the metabolic syndrome induced by a high-fat diet in rats [31]. Although this may imply that the main effect of melatonin on body weight is exerted peripherally through a rise in the energy expenditure, the possibility of central changes in feeding regulatory mechanisms remains open. We hereby report the results of gene expression of NPY, leptin-R, POMC, prolactin-releasing peptide (PrRP), insulin-R, insulin-R substrate (IRS)-1 and IRS-2 in the medial basal hypothalamus (MBH) of the same animals employed in our previous study [31]. The changes observed were correlated with those in blood levels of leptin and adiponectin.

Materials and Methods

Animals and experimental design

The details on the animals and experimental design employed were given elsewhere [31]. Briefly, adult male Wistar rats were kept under standard conditions with controlled light (12:12 h light/dark schedule; lights on at 08:00 h) and temperature (22 ± 2 C). Rats had ad libitum access to a normal or a high-fat diet. Normal rat chow contained 3% fat, 16% protein and 60% carbohydrate (mainly as starch with less than 0.4% fructose) providing a total caloric content of 2.9 Kcal/g. The high (35%) fat chow contained 35% carbohydrates and 20% proteins, providing a total caloric content of 5.4 Kcal/g. Animals were randomly divided into four groups (n= 16/group) as follows: (i) control; (ii) high-fat diet (obese); (iii) obese + melatonin; (iv) control + melatonin. Rats had free access to high-fat or control chow and one of the following drinking solutions for 10 weeks: (a) tap water; (b) 25 μ g/mL of melatonin. Since ethanol was used as a melatonin's vehicle, drinking solutions in groups (i) and (iv) were added 0.015 % ethanol. Water bottles were changed every other day. Because rats drank about 30 mL/day, the daily melatonin dosage used provided approximately 2.3 mg/kg melatonin [31]. Because rats drank 90–95% of this total daily water taken up during the dark period, most melatonin was administered at the time of endogenous melatonin release. The human equivalence dose, calculated by using the body surface area normalization method [32] was about 0.35 mg/kg (about 25-30 mg/day for a 75 kg human adult).

The animals were weighed once a week for 10 weeks and were euthanized by decapitation under conditions of minimal stress at two time intervals: at the middle of the light period (13:00 h) and at the middle of the scotophase (01:00 h). All

experiments were conducted in accordance with the guidelines of the International Council for Laboratory Animal Science (ICLAS). The brains were rapidly removed and the medial basal hypothalamus (MBH) was dissected out following the landmarks of Szentagothai et al. [33]. Trunk blood was collected and plasma samples were obtained by centrifugation of blood at 1,500 x g for 15 min and were stored at -70 °C until further analysis.

Real-time quantitative polymerase chain reaction (qPCR)

Total RNA extraction was performed using the RNeasy protect mini kit and was analyzed using QuantiTec SYBR green kit (Qiagen, Hilden, Germany). The iScript™ cDNA Synthesis Kit (Bio-Rad Laboratories SA; Madrid) was used to synthesize cDNA from 1 µg of total RNA, according to the manufacturer's protocol. The reference gene β -actin was used as a constitutive control for normalization. Reactions were carried out in the presence of 200 nM of specific primers for genes of NPY, POMC, leptin receptor, insulin receptor (IR), insulin receptor substrate (IRS)-1, IRS-2 and prolactin releasing peptide (PrRP). Primers were designed using Primer3 software (The Whitehead Institute, http://frodo.wi.mit.edu/cgi-bin/primer3/primer3_www.cgi) and are shown in Table 1. PCR reactions were carried out in an Eppendorf Real Plex Mastercycler (Eppendorf AG, Hamburg, Germany). The real-time qPCR reaction program included a 94 °C enzyme activation step for 2 min followed by 40 cycles of 95 °C denaturation for 15 s, 60 °C annealing for 30 s and 72 °C extension for 30 s. Detection of fluorescent product was carried out at the end of the 72 °C extension period.

Serial dilutions of cDNA from control MBH were used to perform calibration curves in order to determine amplification efficiencies. For the primers used there were no differences between transcription efficiencies, the amount of initial cDNA in each sample being calculated by the $2^{-\Delta\Delta Ct}$ method [34]. All samples were analyzed in triplicate and in three different measures. Fractional cycle at which the amount of amplified target becomes significant (Ct) was automatically calculated by the PCR device.

To estimate whether treatment or time of day modified the expression of MBH β -actin, PCR employing serial dilutions of this reference gene was performed. In this study Ct did not vary significantly as a function of treatment or of time of day, indicating the validity to employ β -actin as a reference gene.

Biochemical assays

Plasma concentrations of adiponectin and leptin were measured in a multianalyte profiling by using the Luminex-100 system and the XY Platform (Luminex Corporation, Oosterhout, The Netherlands) as described elsewhere [35]. Calibration microspheres for classification and reporter readings as well as sheath fluid were also purchased from Luminex Corporation. Acquired fluorescence data were analyzed by the MasterPlex™ QT software. All analyses were performed according to the manufacturer protocols.

Data analysis

After verifying normality of distribution of data, the statistical analysis of the results was performed by a one-way or a three-way factorial analysis of variance

(ANOVA) followed by Holm-Sidak multiple comparisons tests, as stated. The three-factor ANOVA was used to test for differences between individual data grouped according to the levels of each factor, i.e. treatment, diet, and time period, and for interactions between the factors. Four hypotheses were tested in the three-factor ANOVA: (a) there was no difference between the levels of treatment (melatonin or vehicle); (b) there was no difference between the levels of diet (normal or high-fat diet); (c) there was no difference between the levels of time period (middle of day or middle of night); (d) there was no interaction among the three factors. P values lower than 0.05 were taken as evidence of statistical significance.

Results

The detailed description of the changes in individual daily chow and water consumption and body weight in the four experimental groups are published elsewhere [31]. Body weight of high-fat fed rats receiving vehicle attained values that were 44 % higher than controls after 70 days of treatment. The concomitant administration of melatonin significantly attenuated body weight increase in high-fatfed rats without affecting chow or water consumption [31].

Figure 1 depicts the changes in MBH mRNA levels encoding NPY, POMC, leptin-R and PrRP at the two examined time intervals (13:00 and 01:00 h) in the four studied groups. For NPY, leptin-R and PrRP gene expression, the three way ANOVA indicated that all main factors were significant (F= 28.1, P< 0.001, F= 9.4, P< 0.003 and F= 28.5, P< 0.001 for treatment; F= 56.3, P< 0.001; F=20.4, P< 0.001 and F= 58.1 for diet; F= 5.5, P< 0.03; F= 15.2, P< 0.001 and F= 45.8, P< 0.001 for time of day, respectively) as well as were the interactions “treatment x diet” (F= 20.4, P< 0.001; F= 9.9, P< 0.003 and F=

13.7, $P < 0.001$, respectively) and “treatment x time of day” ($F = 5.9$, $P = 0.02$; $F = 4.1$, $P < 0.04$ and $F = 17.7$, $P < 0.001$, respectively). Collectively the statistical analysis supported the view that the high-fat diet augmented expression of MBH NPY, leptin-R and PrRP genes mainly at night hours and that melatonin decreased it in obese rats mainly at night hours. In the case of POMC gene expression, the three-way ANOVA yielded significance for diet ($F = 37.1$, $P < 0.001$) and time of day ($F = 18.6$, $P < 0.001$) with no significant interaction between them. Thus the high-fat diet augmented expression of MBH POMC mRNA mainly at night hours through an effect insensitive to melatonin treatment (Fig. 1).

Figure 2 shows the changes in MBH mRNA levels encoding insulin-R, IRS-1 and IRS-2. A three-way ANOVA indicated statistical significance for the three factors examined and for the three parameters tested (treatment: $F = 77.8$, $P < 0.001$, $F = 32.2$, $P < 0.001$; $F = 54.3$, $P < 0.001$; diet: $F = 131.1$, $P < 0.001$; $F = 25.4$, $P < 0.001$; $F = 102.5$, $P < 0.001$; time of day: $F = 33.1$, $P < 0.001$; $F = 6.4$, $P < 0.02$; $F = 5.6$, $P < 0.02$, for insulin-R, IRS-1 and IRS-2, respectively). A significant interaction “treatment x diet” was found ($F = 37.1$, $P < 0.001$ for insulin-R; $F = 16.6$, $P < 0.001$ for IRS-1; $F = 16.9$, $P < 0.001$ for IRS-2). Therefore, the statistical analysis supported the conclusion that that the high-fat diet augmented expression of MBH insulin-R, IRS-1 and IRS-2 mRNA, an effect reduced by melatonin administration.

Figure 3 summarizes the changes in plasma levels of leptin and adiponectin. The three-way ANOVA indicated that treatment and diet were significant as main factors (treatment: $F = 11.1$, $P = 0.001$ and $F = 4.4$, $P < 0.04$; diet: $F = 164.9$, $P < 0.001$ and $F = 15.8$, $P < 0.001$ for leptin and adiponectin, respectively). In addition the interaction “treatment x diet” was significant ($F = 38.8$, $P < 0.001$ and $F = 13.8$, $P < 0.001$,

respectively). This supported the conclusion that the augmented levels of leptin and adiponectin seen in high-fat fed rats were counteracted by melatonin. In the All Pairwise Multiple Comparison Procedure of the Holm-Sidak method, and with an overall significance level = 0.05, comparisons for factor: “treatment within normal” for leptin and adiponectin indicated a $t= 15.5$ and 5.27 , respectively $P < 0.001$. Thus melatonin augmented leptin and adiponectin levels significantly in the control group (Fig. 3).

Discussion

Information in the literature on the effect of melatonin on feeding behavior signals is scant. Adult pinealectomized rats administered with melatonin [36], or male rat offspring exposed to melatonin prenatally [37], showed augmented hypothalamic levels of NPY while in submammalian species increases [6], decreases [7,8] or absence of changes [5] in hypothalamic NPY were reported after melatonin. Melatonin reduced the increase in neuronal NPY of striatal neurons in a model of cerebral ischemic damage in rats [38].

In a recently study we examined whether melatonin affected mild inflammation in high-fat fed rats [31]. A number of plasma parameters were measured at two time intervals, i.e. the middle of daylight period and the middle of the scotophase. In obese rats melatonin decreased the augmented circulating levels of the pro-inflammatory signals interleukin (IL)-1 β , IL-6, tumor necrosis factor- α , interferon- γ and C-reactive protein seen in obese rats and restored the depressed levels of the anti-inflammatory cytokines IL-4 and IL-10. Rats fed with the high-fat diet showed significantly higher body weights and augmented systolic blood pressure from the 3rd and 4th week

onwards, respectively, melatonin effectively preventing these changes [31]. In high-fat fed rats circulating low-density lipoprotein-cholesterol, total cholesterol and triglyceride concentration augmented significantly, melatonin counteracting these changes. Melatonin-treated rats showed a decreased insulin resistance, the highest values of plasma high-density lipoprotein-cholesterol and the lowest values of plasma uric acid [31].

The foregoing results reported the changes in MBH expression of a number of feeding behavior signals and in plasma leptin and adiponectin levels in the same animals reported in the previously cited study [31]. The data obtained indicate that the concomitant administration of melatonin counteracted the augmented expression of MBH genes encoding NPY, PrRP, leptin-R, insulin-R, IRS-1 and IRS-2 found in high-fat fed rats. Feeding of rats with a high-fat diet augmented expression of MBH POMC gene through an effect insensitive to melatonin treatment. The augmented levels of circulating leptin and adiponectin seen in high-fat fed rats were decreased by melatonin whereas melatonin augmented plasma leptin and adiponectin in control rats. All these effects of melatonin were seen in the absence of significant modifications of daily food intake suggesting that the satiety signals triggered were compensated by an increase in the central orexinergic pathways to attain a new set point to equal food intake.

The activity of the orexigenic pathway driven by NPY appeared to be effective to counteract food intake inhibition even in the presence of the increased circulating levels of the anorexigenic signals leptin (Fig. 3) and insulin [31], as well as of the augmented expression of MBH POMC, PrRP, leptin-R, insulin-R, IRS-1 and IRS-2 genes. NPY is synthesized in the cell bodies of the arcuate nucleus of the hypothalamus and is

transported to the paraventricular nucleus by axonal flow to achieve high local concentrations [39]. Previous studies in rats showed that intra-hypothalamic injection of NPY stimulates food intake and produced hyperinsulinemia and insulin resistance [40,41]. Moreover, energy expenditure decreases with the simultaneous induction of lipid synthesis in both the liver and the white adipose tissue [41,42]. NPY overactivity results in expression of neuronal insulin-R and leptin-R, thus generating an anorexigenic response that counterbalances the NPY effect on appetite [20,21].

The foregoing results suggest that melatonin administration is able to set up a new equilibrium among feeding signals in the MBH. Remarkably, melatonin reduced both gene expression of the strong orexinergic signal NPY and of the anorexigenic signal PrRP, as well as that of receptors for anorexigenic signals like leptin and insulin and of insulin intracellular signaling (IRS-1, IRS-2).

Blood levels of leptin and adiponectin (Fig. 3) and of insulin [31] were also significantly reduced by melatonin in obese rats while melatonin augmented plasma leptin and adiponectin in control rats. Our present and previous results [26,35] have indicated an increase in mean levels of adiponectin in high fat-fed rats as well as a significant modification in its daily pattern in circulation. In contrast most studies have showed a reduction in plasma concentration of adiponectin in obesity (see for ref.[43]) with very few exceptions [44]. The possibility deserves to be considered that some fat components of the diet employed could be stimulus for adiponectin secretion in view that several nutraceutical products like fish oil [45], safflower oil [46] or linoleic acid [47] increased adiponectin production in either animals or humans.

Data in the literature on the effect of melatonin on feeding behavior signals are scant. By immunohistochemical staining of NPY neurons in the hypothalamic arcuate

nucleus of Sprague-Dawley rats a depressive effect of pinealectomy and a reversal after melatonin administration were found [36]. In that study, and resembling the results of Figure 1 (upper left panel), melatonin administration to sham-operated rats did not modify NPY immunopositive cell number in the arcuate nucleus significantly. One of us reported that prenatal melatonin exposure of the mother augmented hypothalamic NPY content in the male rat offspring [37]. In zebrafish, melatonin reduces food intake and increases genes codifying for molecules involved in feeding inhibition such as leptin, with a significant reduction in the major orexigenic signals including NPY [7,8]. In the goldfish, a species in which melatonin reduces body weight, plasma leptin and hypothalamic NPY remained unaltered after melatonin, suggesting that these feeding regulatory signals are not involved in the effects of melatonin on energy homeostasis [5]. In the hypothalamus of the rainbow trout melatonin altered glucosensing capacity and induced increased mRNA expression levels of NPY and decreased mRNA levels of POMC and CART [6]. In the study reported herein melatonin was unable to modify POMC gene expression.

To our knowledge the present study is the first reporting an effect of melatonin on the expression of the gene encoding PrRP. This peptide has been linked to the regulation of appetite, being one of the more recently described anorexigenic signals. The gene encoding PrRP is located mainly in the hypothalamus and the medulla in areas involved in the regulation of appetite which also express PrRP receptors [48]. PrRP effects on appetite are presumably mediated by changes in circulating levels of leptin as well as for its involvement in the regulation of anorexigenic peptides in the melanocortin system [49].

It must be noted that regardless of the universality of the changes following melatonin treatment of high-fat fed rats, melatonin effectiveness to reduce body weight appeared to be independent on the central mechanisms regulating food intake since a new set point to equal food intake was rapidly attained. The data indirectly support the view that the main effect of melatonin on body weight may be exerted peripherally through a rise in the energy expenditure rather than centrally on feeding behavior mechanisms. Results in the literature tend to support this conclusion. The nocturnal locomotor activity in rats increased after melatonin [10,50] and the core body temperature also rise [10], a finding compatible with a rise in energy expenditure. Tan and co-workers put forth the hypothesis that the brown adipose tissue (BAT) is a factor through which animals lose weight in response to melatonin administration (and gain weight when there is a deficiency of melatonin) [51]. BAT is present in adult humans [52], has a high metabolic activity and is responsible for nonshivering thermogenesis [53]. As a result, BAT burns calories for the purpose of heat production, thereby consuming glucose and fatty acids and limiting fat deposition [54,55]. BAT is also of crucial importance in the regulation of glycemia, lipidemia and insulin sensitivity [56]. To what extent the weight-loss-promoting effect of melatonin is attributable to an increase in energy expenditure by BAT deserves further exploration [53].

One important limitation of the present descriptive study is that gene expression needs to be completed with Western blotting analysis of the proteins formed in order to obtain a complete understanding of melatonin effect on feeding behavior signals. In the present study the results are presented as relative changes of mRNA levels and while statistically significant differences were detected, the presentation of data may

distort the actual physiological importance of such changes since a similar relative change in NPY and insulin-R mRNA could well represent a several fold difference in the actual amount of the specific protein synthesized. In addition, the gross picture of daily food intake reported [31] must be completed with the analysis of feeding frequency and meal size at both daylight and scotophase periods during the whole observation period to further substantiate subtle changes in feeding behavior after melatonin.

In summary, an adequate supplementation with melatonin lowers body weight and body weight gain in a rodent model of hyperadiposity [31]. This might be the result of the recovery of feeding behavior signaling, of the disappearance of insulin resistance and glucose intolerance and of the accentuation of the energy expenditure over the energy intake, resulting in weight loss and stabilization of weight gain.

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Conflict of Interest

The authors declare no conflict of interest.

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FIGURE LEGENDS

Figure 1. Expression of mRNA for NPY, POMC, leptin-R and PrRP in MBH of rats fed a normal or a high-fat diet and melatonin (25 µg/mL) or vehicle in drinking water for 10 weeks. Groups of rats were euthanized at the middle of the light period (13:00 h) or at the middle of the scotophase (01:00 h). MBH mRNA levels encoding NPY, POMC, leptin-R and PrRP were measured as described in the text. Shown are the means ± SEM (n= 7-8 per group). Letters indicate significant differences in a one-way ANOVA followed by a Holm-Sidak multiple comparisons test at a given time interval, as follows: (a) P< 0.01 as compared to the remaining groups; (b) P< 0.05 as compared to obese and obese + melatonin groups; (c) P< 0.05 as compared to the remaining groups. For further statistical analysis, see text.

Figure 2. Expression of mRNA for insulin-R, IRS-1 and IRS-2 in MBH of rats fed a normal or a high-fat diet and melatonin (25 µg/mL) or vehicle in drinking water for 10 weeks. Groups of rats were euthanized at the middle of the light period (13:00 h) or at the middle of the scotophase (01:00 h). MBH mRNA levels encoding insulin-R, IRS-1 and IRS-2 were measured as described in the text. Shown are the means ± SEM (n= 7-8 per group). Letters indicate significant differences in a one-way ANOVA followed by a Holm-Sidak multiple comparisons test at a given time interval, as follows: (a) P< 0.05 as compared to the remaining groups; (b) P< 0.05 as compared to obese and obese + melatonin groups. For further statistical analysis, see text.

Figure 3. Plasma levels of leptin and adiponectin in rats fed a normal or a high-fat diet and melatonin (25 µg/mL) or vehicle in drinking water for 10 weeks. Groups of rats

were euthanized at the middle of the light period (13:00 h) or at the middle of the scotophase (01:00 h). Shown are the means \pm SEM (n= 7-8 per group). Letters indicate significant differences in a one-way ANOVA followed by a Holm-Sidak multiple comparisons test at a given time interval, as follows: (a) $P < 0.05$ as compared to obese and obese + melatonin groups; (b) $P < 0.05$ as compared to the remaining groups. (*) $P < 0.001$ vs. control when compared for the factor: "treatment within normal" in the All Pairwise Multiple Comparison Procedure of the Holm-Sidak method, three-way ANOVA . For further statistical analysis, see text.

Table 1. Sequence of the primers used for real-time PCR

Gene	Primers		Product Size (bp)
β- Actin	Forward	5'-ccagatcatgtttgagaccttcaa-3'	91
	Backward	5'-ccagaggcgtacagggatagc-3'	
NPY	Forward	5'-tactccgctctgcgacacta-3'	110
	Backward	5'-tcatcagcaggaggagtgc-3'	
POMC	Forward	5'-cgtgtgtccttctgactcc-3'	108
	Backward	5'-gactggctttccaagatac-3'	
Lep-R	Forward	5'-cgtgtgtccttctgactcc-3'	107
	Backward	5'-gactggctttccaagatac-3'	
Insulin-R	Forward	5'-tcttcaggcaatggtgct-3'	119
	Backward	5'gtggaggagatgttgggaaa-3'	
IRS-1	Forward	5'-ggcaccatctcaacaatcct-3'	104
	Backward	5'-tttcccaccaccatactg-3'	
IRS-2	Forward	5'-ctaccactgagccaagag-3'	151
	Backward	5'-ccagggatgaagcaggacta-3'	
PrRP	Forward	5'-caccactcaaggacagagg-3'	115
	Backward	5'-cagcagcaagcacagaagc-3'	

Figure 1

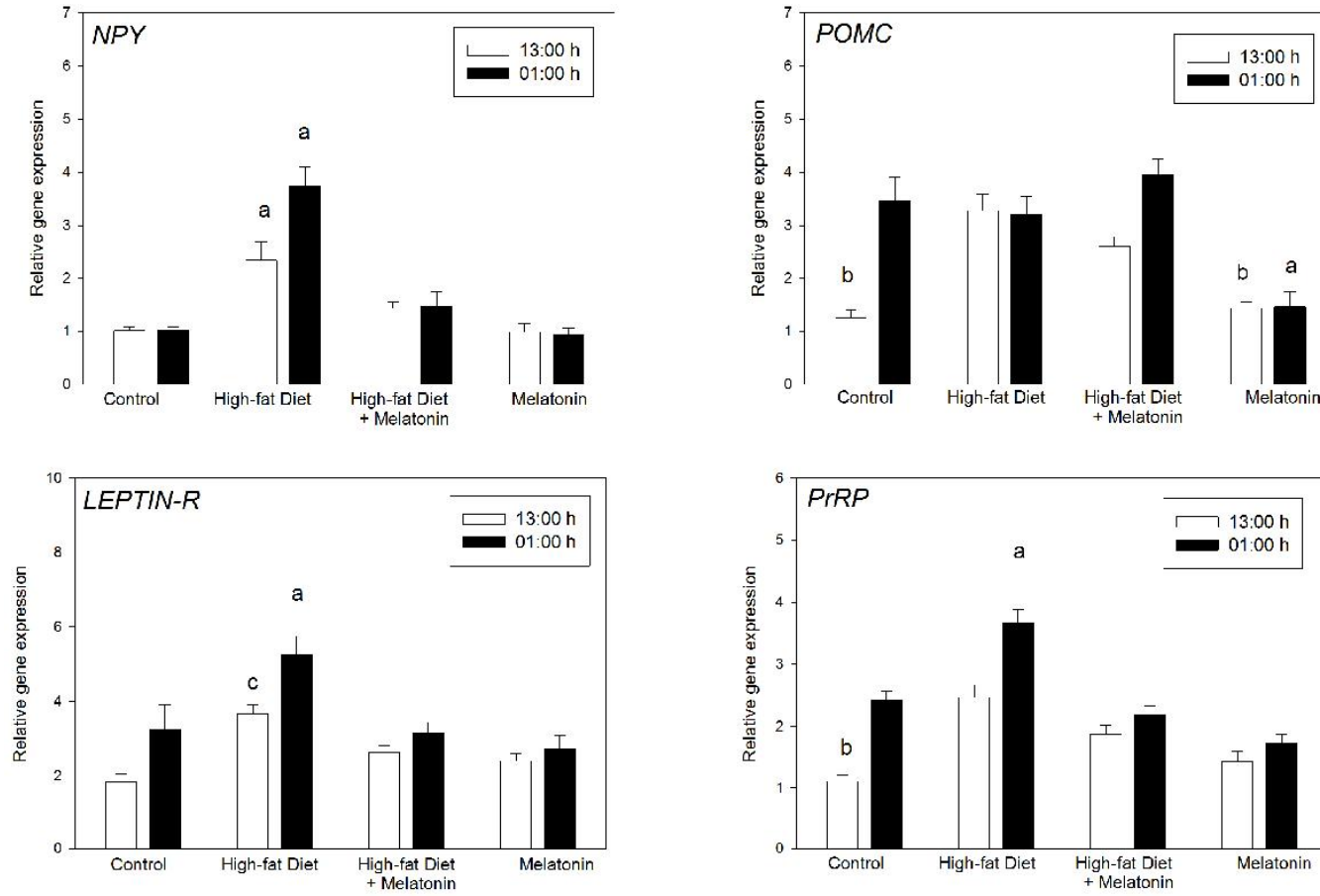


Figure 2

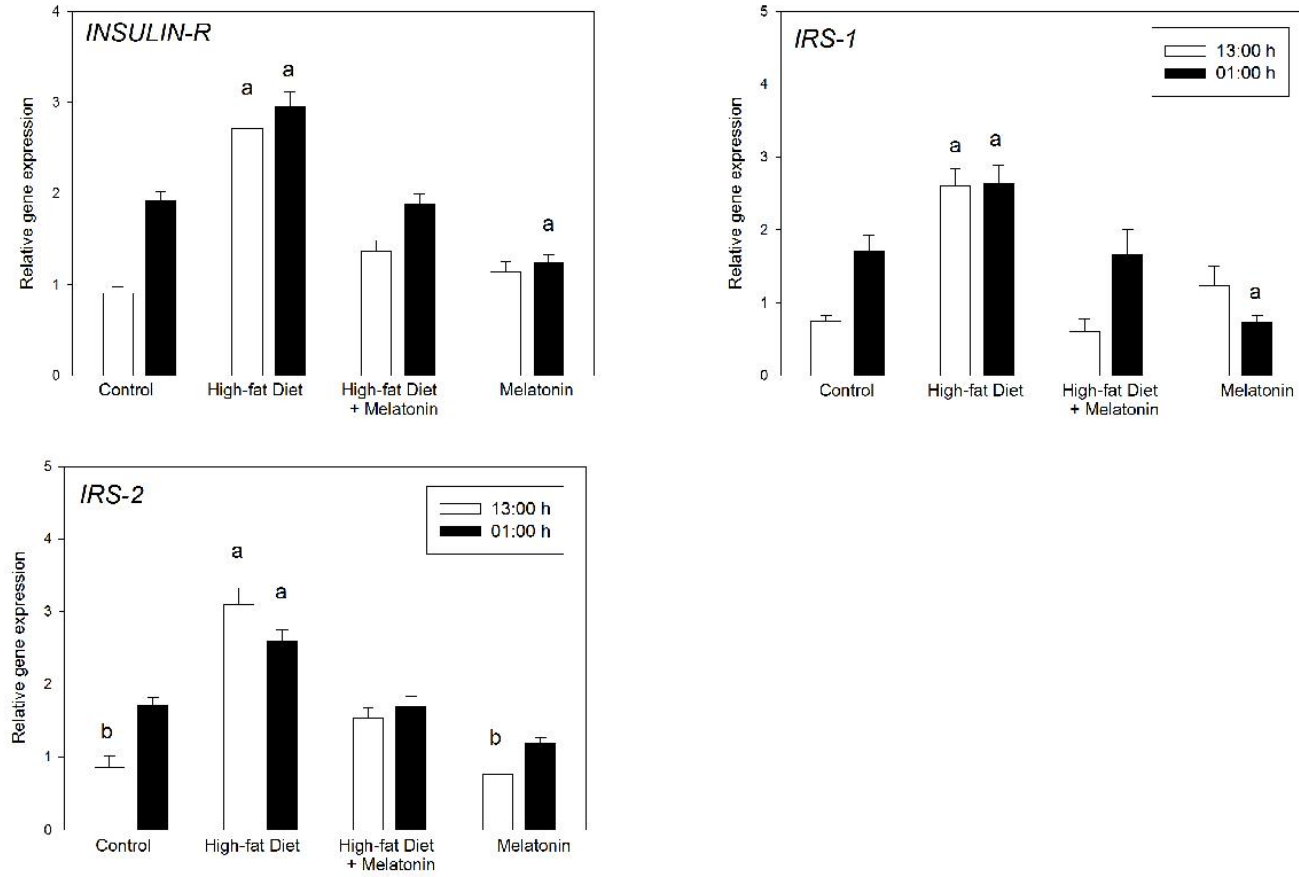


Figure 3

