

# Neuropeptide Y Overexpression in the Preweanling Zucker (*fa/fa*) Rat

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Received 19 April 1999; Accepted 29 April 1999

KOWALSKI, T. J., T. A. HOUPPT, J. JAHNG, N. OKADA, S.-M. LIU, S. C. CHUA AND G. P. SMITH. *Neuropeptide Y overexpression in the preweanling Zucker (fa/fa) rat.* *PHYSIOL BEHAV* **67**(4) 521–525, 1999.—Hypothalamic preproNPY overexpression in the Zucker fatty (*fa/fa*) rat was examined. In situ hybridization was used to determine the relative level of preproNPY mRNA in the arcuate nucleus of *+/+*, *+/fa*, and *fa/fa* pups aged postnatal day 2 (P2), 5, 9, 12, or 25. The relative optical density (ROD) of probe hybridization in the arcuate, the area of hybridization (A), and the product of  $ROD \times A$  (a measure of total arcuate preproNPY mRNA hybridization) were measured. Values were normalized to the mean *+/fa* value within each litter. Initial analysis showed that preproNPY mRNA hybridization ( $ROD \times A$ ) in *fa/fa* pups was significantly higher than *+/fa* and *+/+* pups on P9, 12, and 25, and significantly higher than *+/fa* on P5. No significant difference between lean (*+/+* and *+/fa*) genotypes, however, were observed at any age tested. Values from the lean genotypes were, therefore, pooled, and data were normalized to the mean value of lean animals for analysis. This analysis revealed that preproNPY mRNA hybridization in *fa/fa* pups was higher than lean littermates as early as P2. © 1999 Elsevier Science Inc.

Genetic obesity    Arcuate nucleus    Hypothalamus    Development of obesity

THE Zucker fatty rat (*fa/fa*) is a model of genetic obesity with an autosomal recessive mode of inheritance (42). *fa/fa* rats display a juvenile-onset obesity with accompanying metabolic and behavioral abnormalities, such as hyperphagia, decreased energy expenditure, hyperinsulinemia, and hypercorticosteronemia (7). The *fa* mutation has recently been identified as an amino acid substitution in the extracellular domain of the receptor for leptin (10,11). All leptin receptor isoforms are affected by this mutation. As a consequence, the number of receptors at the cell surface (27), and their signal transduction efficiency (40), are decreased. Additionally, *fa/fa* rats have elevated plasma leptin levels (39), and are resistant to exogenous leptin administration (13).

Studies comparing adult *fa/fa* and *+/+* rats have shown that *fa/fa* rats have higher preproneuropeptide Y (preproNPY) mRNA in the arcuate nucleus of the hypothalamus (30), higher levels of NPY peptide in several hypothalamic nuclei (4), and higher rates of NPY release into the PVN (15). Hypo-

thalamic NPY activity is elevated in other models of obesity (37,38), and chronic intracerebroventricular administration of NPY mimics many of the physiological and behavioral symptoms present in obesity (41), suggesting that increased hypothalamic NPY activity is involved in the etiology of obesity and its associated metabolic and behavioral disturbances.

NPY is an orexigenic peptide when administered centrally, and is most potent when injected into the paraventricular nucleus of the hypothalamus (PVN) (34). The arcuate nucleus is the primary site of NPY synthesis in the hypothalamus (1), and an arcuo-paraventricular NPYergic projection exists (2). This arcuo-paraventricular NPYergic projection is believed to play a role in feeding behavior because its activity is correlated with deprivation state; arcuate preproNPY mRNA, PVN immunoreactive NPY, and release of NPY into the PVN all increase during periods of food deprivation and return to basal levels with refeeding (6,21,28). Additionally, exogenous central NPY administration decreases sympathetic drive to

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brown adipose tissue (decreasing thermogenesis) (5), increases plasma insulin levels (25), and stimulates white adipose tissue lipoprotein lipase activity (5).

Recent work has shown that central leptin administration to deprived rats or to *ob/ob* mice (that lack leptin) decreases arcuate preproNPY mRNA expression (31,32). This has led to the hypothesis that the inability of *fa/fa* rats to respond appropriately to circulating leptin results in increased arcuate preproNPY expression with subsequent hyperphagia and decreased energy expenditure (36). It is implicit within this hypothesis that the elevation in hypothalamic NPY expression occurs prior to or simultaneously with the emergence of hyperphagia and decreased energy expenditure. Decreased energy expenditure in *fa/fa* rats has been observed as early as P2 (26) and hyperphagia at P12 (23). The onset of arcuate preproNPY overexpression in *fa/fa* rats, however, has been less clearly defined, varying from P6 to after week 11 (12,20). The purpose of this study was to identify the age that preproNPY overexpression occurs in *fa/fa* rats more precisely. Our results indicate that arcuate nucleus preproNPY mRNA levels in *fa/fa* pups are higher than lean pups as early as P2.

#### MATERIALS AND METHODS

##### Animals

Lean (+/+ and +/*fa*) and obese (*fa/fa*) Zucker rats aged P2, 5, 9, 12, and 25 were used in the experiments. Animals were the progeny of primiparous and multiparous +/*fa* females mated with +/*fa* males. All breeders were derived from the Vassar College colony (Poughkeepsie, NY). Mating pairs and pregnant females were housed in Plexiglas containers with wood shavings as bedding. Rats received pelleted chow and water ad lib, and were maintained on a 12:12 h light:dark cycle (0700–1900 H) at  $22 \pm 2^\circ\text{C}$ . Pregnant females were checked daily for pups, and the day pups were first seen was termed P0. On P3 or P4, animals assigned for study on P9, 12 or, 25 were earclipped for identification and tissue collection (for genotyping; see below). Pups studied on P2 or P5 were marked for identification using a dorsal subcutaneous injection of ink, and spleen was collected for genotyping upon death. A total of 116 pups (+/+  $n = 24$ ; +/*fa*  $n = 56$ ; *fa/fa*  $n = 36$ ) were used, with 27 pups tested on P2, 23 on P5, 25 on P9, 17 on P12, and 24 on P25. Four to seven litters consisting of 7 to 12 pups were used at each age (not all pups within each litter were used for this experiment). All of the animals from each litter tested were sacrificed on the same day. Data from 45 lean Zucker pups aged P2, 5, 9, and 12 used in the calculations has been previously reported (22).

##### Tissue Preparation

Between 0700 to 1200 h, animals were individually removed from the dam, weighed, and given an overdose of sodium pentobarbital i.p. When unresponsive, animals were transcardially perfused with 0.9% NaCl, 0.5% NaNO<sub>2</sub>, 100 U/mL heparin followed by ice-cold 4% buffered paraformaldehyde (4% paraformaldehyde in 0.1 M phosphate buffer, pH 7.4). Brains were removed, postfixed for 24 h in 4% buffered paraformaldehyde and cryoprotected in 30% sucrose at  $4^\circ\text{C}$  for 48 h before sectioning.

##### In Situ Hybridization

Forty-micron frontal sections were cut using a sliding microtome and placed into ice-cold  $2\times$  SSC (SSC = 0.15 M NaCl/0.015 M Na Citrate). Six to 10 sections through the mid-

region of the arcuate were taken. Tissues were prehybridized in glass vials in 1 mL of 60% formamide, 0.02 M Tris pH 7.4, 1 mM EDTA, 10% dextran sulfate, 0.8% Ficoll, 0.8% PVP, 0.8% BSA,  $2\times$  SSC, 0.1 M dithiothreitol, and 1.6 mg/mL herring sperm DNA for 2 h at  $48^\circ\text{C}$ . After 2 h, radiolabeled probe was added ( $\sim 1.0 \times 10^7$  cpm/vial) and incubated for 16–20 h at  $48^\circ\text{C}$ . Hybridization was performed using a 511 base pair (bp)  $\alpha$ -[<sup>35</sup>S]dATP random prime labeled cDNA encoding preproNPY (19). Sections were then sequentially rinsed in  $2\times$ ,  $1\times$ ,  $0.5\times$ ,  $0.25\times$ , and  $0.125\times$  SSC for 15 min at  $48^\circ\text{C}$  and placed into 0.1 M phosphate buffer for mounting. The tissue sections were mounted onto gelatin-coated slides, air dried, and apposed to X-ray film ( $\beta$ -max, Amersham) for 12–24 h. Each litter was processed together, and slides were exposed to the same film. Slides were then dipped into photoemulsion (Kodak) and exposed at  $4^\circ\text{C}$  for 1 week for histological verification of probe labeling.

Probe hybridization in the arcuate was quantified by densitometry of autoradiograms using the MCID system (Imaging Research, Inc.; St. Catherines, Canada). The relative optical density of hybridization signal in the arcuate (ROD), the area of hybridization in the arcuate (A), and the product of ROD and A (ROD  $\times$  A; a measure of total hybridization in the arcuate) were determined from 6–10 sections/animal. The mean ROD, A, and ROD  $\times$  A values from each animal were used in the calculations (see below).

##### Determination of Genotype at *Lepr*

Genotypes were determined according to a previously described method (11). Briefly, the A to C mutation at nucleotide 880 of *Lepr*<sup>fa</sup> introduces an MspI restriction site, which can be used to detect the number of copies of the *Lepr*<sup>fa</sup> allele. Tissue was digested with proteinase K and genomic DNA was extracted using the Quiagen QIAamp Tissue Kit. Primers: (5' TGAAGCCCGATCCACCGCTGG 3' and 5' CTCTCT-TACGATTGTAGAATTCTC 3') were used to generate a 143 base pair polymerase chain reaction (PCR) fragment. PCR was performed on a Perkin-Elmer DNA Thermal Cycler using the following conditions:  $92^\circ\text{C}$  (2 min), 1 cycle;  $92^\circ\text{C}$  (30 s),  $55^\circ\text{C}$  (30 s),  $72^\circ\text{C}$  (1 min), 35 cycles; and  $72^\circ\text{C}$  (5 min), 1 cycle. The Advantage Genomic PCR Kit (Clontech) was used for PCR reactions. The PCR fragments were digested with MspI (80 mU/ $\mu\text{L}$  final concentration) for 1.5 h at  $37^\circ\text{C}$  and electrophoresed on a 2% agarose–2% low melting point agarose gel. Digestion yields an uncut 143 bp product for the wild-type allele, while the mutant allele (*fa*) yields both a 106 bp and a 37 bp product.

##### Statistical Analysis

Results are presented as means  $\pm$  SEM. The ROD, A, and ROD  $\times$  A measures were normalized to the mean of heterozygotes within each litter. Heterozygotes were chosen because they were the most numerous. Significant differences in preproNPY mRNA levels among genotypes were initially assessed by a Kruskal-Wallis *H*-test. Post hoc analyses were performed using a Mann-Whitney *U*-test where appropriate. No significant differences between +/+ and +/*fa* (lean) pups were observed at any age tested. In a subsequent analysis, therefore, lean animals were pooled and the ROD, A, and ROD  $\times$  A of all animals were normalized to lean animals within each litter. Significant differences between *fa/fa* and lean animals were assessed at each age using a *U*-test. An *H*-test was used to evaluate the magnitude of *fa/fa* overexpression among ages. A *p*-value  $<0.05$  was considered significant.

TABLE 1  
 BODY WEIGHT AND NORMALIZED RELATIVE OPTICAL DENSITY (ROD), AREA (A), AND  
 ROD × A IN PREWEANLING LEAN (+/+, +/fa) AND OBESE (*fa/fa*) ZUCKER RATS

| Age | Genotype (n)     | Body Weight (g) | Normalized* ROD | Normalized* Area | Normalized* ROD × A |
|-----|------------------|-----------------|-----------------|------------------|---------------------|
| P2  | <i>fa/fa</i> (8) | 5.58 ± 0.23     | 1.10 ± 0.05     | 1.18 ± 0.04†     | 1.27 ± 0.07†        |
|     | +/fa (13)        | 5.84 ± 0.15     | 1.00 ± 0.04     | 1.00 ± 0.05      | 1.00 ± 0.08         |
|     | +/+ (6)          | 5.66 ± 0.15     | 1.04 ± 0.08     | 1.10 ± 0.09      | 1.16 ± 0.17         |
| P5  | <i>fa/fa</i> (8) | 9.26 ± 0.63     | 1.09 ± 0.05     | 1.22 ± 0.08†     | 1.32 ± 0.07†        |
|     | +/fa (9)         | 9.16 ± 0.47     | 1.00 ± 0.03     | 1.00 ± 0.04      | 1.00 ± 0.07         |
|     | +/+ (6)          | 9.00 ± 0.52     | 1.04 ± 0.02     | 1.13 ± 0.12      | 1.18 ± 0.13         |
| P9  | <i>fa/fa</i> (7) | 17.05 ± 0.70    | 1.18 ± 0.04‡    | 1.09 ± 0.08      | 1.29 ± 0.12‡        |
|     | +/fa (12)        | 16.90 ± 0.65    | 1.00 ± 0.01     | 1.00 ± 0.02      | 1.00 ± 0.03         |
|     | +/+ (6)          | 18.10 ± 0.79    | 0.95 ± 0.03     | 0.95 ± 0.08      | 0.91 ± 0.09         |
| P12 | <i>fa/fa</i> (5) | 26.6 ± 4.0      | 1.28 ± 0.09†    | 1.22 ± 0.09§     | 1.60 ± 0.15†        |
|     | +/fa (9)         | 24.0 ± 1.2      | 1.00 ± 0.04     | 1.00 ± 0.07      | 1.00 ± 0.10         |
|     | +/+ (3)          | 28.4 ± 2.8      | 1.10 ± 0.10     | 0.91 ± 0.05      | 0.99 ± 0.13         |
| P25 | <i>fa/fa</i> (8) | 51.9 ± 5.6      | 1.37 ± 0.05‡    | 1.30 ± 0.06‡     | 1.78 ± 0.13‡        |
|     | +/fa (13)        | 49.8 ± 3.7      | 1.00 ± 0.01     | 1.00 ± 0.03      | 1.00 ± 0.03         |
|     | +/+ (3)          | 43.7 ± 1.7      | 0.98 ± 0.03     | 0.91 ± 0.09      | 0.89 ± 0.09         |

\*Relative optical density (ROD) and area of hybridization (A) values were normalized to +/fa littermates.

†Significantly different from +/fa (P < 0.05).

‡Significantly different from +/+ and +/fa (p < 0.05).

§Significantly different from +/+ (p < 0.05).

RESULTS

PreproNPY mRNA levels in the arcuate nucleus (expressed as ROD × A) of *fa/fa* pups were higher than those in +/+ and +/fa pups as early as P9 (Table 1). On P5, *fa/fa* pup preproNPY mRNA levels were higher than their +/fa littermates but not +/+. No differences between lean pups were observed at any age. When the data from lean animals within each age group were pooled for comparison to *fa/fa* pups, preproNPY mRNA levels (ROD × A) of *fa/fa* pups were higher on P2, P9, 12, and 25, with a trend towards overexpression on P5 (p = 0.057) (Fig. 1).

On P2, the significantly higher ROD × A observed in *fa/fa* animals was due to a significantly larger A, whereas on P9 it was due to a higher ROD. On P12 and 25, the higher ROD × A was due to an increase in both ROD and A. There was a significant effect of age on the magnitude of overexpression

(ROD × A normalized to lean pups) in *fa/fa* pups. The normalized ROD × A of *fa/fa* rats on P25 was significantly higher than on P2 and 5. The same results were observed in the normalized A, however, the ROD of P25 pups was higher than on P2, 5, and 9. No differences in body weight were observed at any age (Table 1).

The preproNPY mRNA hybridization was observed in small cells within the medial portion of the arcuate nucleus, as previously reported (6). PreproNPY mRNA hybridization was also observed in the cortex, the reticular thalamus, and the hippocampus. Similar levels of preproNPY mRNA hybridization were observed in the cortex and reticular thalamus of *fa/fa* and lean pups on P12 (cortex: 1.00 ± 0.03 versus 0.98 ± 0.12 and reticular n.: 1.00 ± 0.03 versus 0.94 ± 0.10; normalized ROD in lean versus *fa/fa* pups, respectively), indicating that the elevated NPY expression in *fa/fa* animals was specific to the arcuate nucleus.

DISCUSSION

This study shows that the levels of arcuate preproNPY mRNA in *fa/fa* pups are higher than those in lean (+/+ and +/fa) pups as early as P2. The presence of preproNPY overexpression on P2, 9, 12, and 25, with a trend towards overexpression on P5, is interpreted as a consistent elevation in arcuate preproNPY mRNA in the *fa/fa* rat from P2. The magnitude of preproNPY overexpression in *fa/fa* rats increases from approximately 1.2-fold higher during the first week of life to a level similar to that previously reported in 5-week-old *fa/fa* rats (29) by P25. Although effects of heterozygosity in Zucker pups have been demonstrated for both fat mass and fat-free dry mass (33,35), we did not detect differences in preproNPY expression between +/+ and +/fa genotypes in this study.

The second postnatal day is the earliest age at which a defect in the expression of a hypothalamic neuropeptide in the *fa/fa* rat has been reported. The onset of preproNPY overexpression in *fa/fa* animals precedes the earliest detection of an

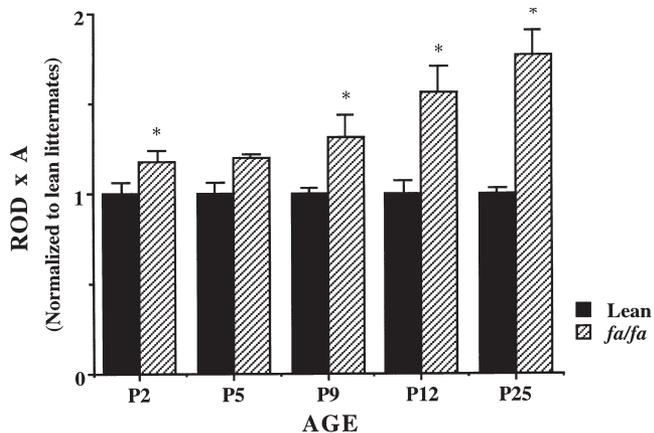


FIG. 1. ROD × A of Zucker rat pups normalized to lean littermates. \*Significantly different from lean pups (p < 0.05).

increase in fat mass on P7 (24), and of hyperphagia during independent ingestion on P12 (23). At the time of preproNPY overexpression in *fa/fa* pups at P2, however, decreased energy expenditure is apparent (26). These observations are consistent with the hypothesis that increased hypothalamic NPY activity in the *fa/fa* rat contributes to the decreased energy expenditure and increased food intake with subsequent obesity (36).

Deprivation (4 to 24 h) of rat pups as early as P2 increases the intake of milk or sucrose in ingestive tests independent of the dam (17,18). Additionally, increased NPY expression in response to a 24-h deprivation is observed as early as P2 in lean Zucker rats (22), and exogenous NPY administration to rats aged P2 elicits an increase in water and milk intake in an independent ingestive test (9). These data suggest that shortly after birth, the hypothalamic NPY system is responsive to deprivation, and may be driving independent ingestive behavior as described in the adult. Although *fa/fa* rats overexpress NPY as early as P2, increased suckling has not been observed in the *fa/fa* rat (8,16), while the earliest age that hyperphagia during independent ingestion is observed is P12 (23). Furthermore, others have reported that an increase in hypothalamic NPY content in *fa/fa* rats was observed on P30 but not P16 (3), after hyperphagia is evident. Studies of *fa/fa* pups examining NPY release in the PVN, and the response to exogenous NPY, are required to better define the role of hypothalamic NPY and arcuate preproNPY overexpression in the initiation of hyperphagia.

The increased arcuate preproNPY expression seen with 24 to 72 h of deprivation in adult rats has been attributed to a decrease in circulating leptin associated with deprivation. This is supported by the fact that an absence of leptin or of a functional leptin receptor, as seen in the *ob/ob* mouse and *fa/fa* rat, respectively, results in the overexpression of arcuate preproNPY (30,37). Additionally, central leptin administration decreases arcuate preproNPY expression in deprived rats (32) and in *ob/ob* mice (31). Because *fa/fa* rats overexpress NPY on P2, it is likely that NPY-expressing neurons in the arcuate are sensitive to circulating leptin at this early age. Consistent with this, deprivation of rat pups aged P0 to P1 decreases brown adipose tissue leptin expression and plasma leptin levels (14), while a 24-h deprivation in lean Zucker rats increases arcuate preproNPY expression as early as P2 (22).

In summary, overexpression of arcuate preproNPY is observed in *fa/fa* pups as early as P2. The role of increased preproNPY expression in the emergence of hyperphagia and metabolic disorders in the *fa/fa* rat remains to be determined.

#### ACKNOWLEDGEMENTS

This work was funded by NIH Grants MH 00149, T32 MH 18390, and MH 15455. We thank Drs. Aron Weller and James Gibbs for constructive criticism of this article.

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