Postnatal changes in ghrelin mRNA expression and in ghrelin-producing cells in the rat stomach

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Abstract

Ghrelin was recently isolated from the rat stomach as an endogenous ligand for the GH secretagogue receptor. Although it is well known that a large amount of ghrelin is produced in the gastrointestinal tract, developmental changes in ghrelin mRNA expression and differentiation of ghrelin-immunopositive (ghrelin-ip) and mRNA-expressing (ghrelin-ex) cells in the stomach have not been elucidated.

In this study, we therefore investigated the changes in ghrelin mRNA expression levels and in the numbers of ghrelin-ip and -ex cells in the stomachs of 1- to 8-week-old male and female rats by Northern blot analysis, immunohistochemistry and *in situ* hybridization. Northern blot analysis showed that the level of weak ghrelin mRNA expression was low in the postnatal period but then increased in a dimorphic pattern, i.e. transient stagnation at 4 weeks in the male rats and at 5 weeks in the

female rats. The number of ghrelin-ip and ghrelin-ex cells also increased after birth, and more numerous ghrelin cells were found in female rats than in male rats, and this finding was confirmed by Northern blot analysis. Ghrelin-ip and -ex cells first appeared in the glandular base of the fundic gland and then they were found in the glandular base and the glandular neck at 3 weeks of age, suggesting that the distribution of ghrelin cells is extended from the glandular base to the glandular neck during the postneonatal development period. This is the first report on detailed changes in postneonatal ghrelin expression level and in the number of ghrelin cells in the rat stomach. The sexual dimorphism of ghrelin expression and ghrelin cell differentiation suggest that ghrelin plays an important physiological role in the stomach.

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Introduction

Ghrelin was recently isolated from the rat stomach as an endogenous ligand for the growth hormone secretagogue receptor (GHS-R) and was found to consist of 28 amino acids, in which the third serine residue is *n*-octanoylated (Kojima *et al.* 1999). Ghrelin has been shown to regulate not only growth hormone (GH) release from the pituitary gland *in vivo* and *in vitro* (Kojima *et al.* 1999, Date *et al.* 2000, Seoane *et al.* 2000, Yamazaki *et al.* 2002) but also to have various physiological functions, such as food intake, gastric acid secretion and gastric motor activity (Masuda *et al.* 2000, Wren *et al.* 2000, Date *et al.* 2001, Nakazato *et al.* 2001). It has been reported that ghrelin produced in the stomach is a major source of plasma ghrelin (Ariyasu *et al.* 2001) and that ghrelin in the stomach may play an important role, either directly or indirectly, in these

functions (Asakawa *et al.* 2001a, Inui 2001). Rat ghrelin cells were found to be localized in the mucosa of the stomach, duodenum, ileum, cecum and colon but not in the myenteric plexus, and they can be classified into open- and closed-type cells. The largest number of ghrelin cells was found in the stomach (Sakata *et al.* 2002). It has also been reported that ghrelin and 27 amino acid des-Gln(14)-ghrelin (Hosoda *et al.* 2000b) exist as two molecular forms in the stomach, i.e. *n*-octanoyl ghrelin, a physiologically active form, and des-*n*-octanoyl ghrelin, an inactive form (Hosoda *et al.* 2000a), and that ghrelin mRNA in the stomach increases during fasting (Toshinai *et al.* 2001) and stress (Asakawa *et al.* 2001b).

GH secretory patterns change during development in both male and female rats, and GH secretion clearly shows a sexually dimorphic peak at the onset of puberty (Eden 1979). Although it has been reported that GH-releasing

hormone (GHRH) and somatostatin (SRIF) mRNA levels show age-dependent changes and sexual dimorphism (Argente *et al.* 1991), the developmental GH secretory pattern cannot be explained only by these neuropeptides. Meanwhile, the ghrelin gene has been reported in the pituitary gland and is highest in the fetus and declines during the postembryonic period, and pituitary GHS-R mRNA expression level has been found to fluctuate with increasing age in young rats (up to 70 days) (Kamegai *et al.* 1999), suggesting that pituitary ghrelin is an important hormone for weaning, body growth or onset of puberty during neonatal and postneonatal periods.

On the other hand, it is thought that ghrelin produced in the stomach is the major source of circulating ghrelin (Ariyasu *et al.* 2001) and this plasma ghrelin also may exert a powerful influence on the development of the rat. It is thus important to clarify the developmental expression of ghrelin mRNA and the differentiation of ghrelin cells in the stomach. In this study, we therefore investigated the developmental change in ghrelin in the stomach of male and female rats by Northern blot analysis, immunohistochemistry (IHC) and *in situ* hybridization (ISH).

Materials and Methods

Animals

Male and female Wistar rats (1–8 weeks of age) were purchased from Japan Laboratory Animals, Inc. (Tokyo, Japan) and were housed under controlled conditions (temperature of 23 °C and light from 0800 to 2000 h). The rats were also provided with standard rat chow and water *ad libitum*. All procedures used in this study were performed in accordance with institutional guidelines for animal care at Saitama University.

Tissue preparations

Animals were deeply anesthetized with sodium pentobarbital (50 mg/kg i.p.) and perfused first with 10 mM PBS, pH 7.5, and then with 4% paraformaldehyde in 50 mM phosphate buffer (PB), pH 7.4. Segments of the stomach, approximately 1 cm in length, were quickly removed and opened along their longitudinal axes. The tissue blocks were dehydrated with an ascending ethanol series and immersed in xylene and then embedded in Paraplast (Oxford Labware, MO, USA). Serial sections (6 µm thick) were made and mounted on silane (ShinEtsu Chemicals, Tokyo, Japan)-coated slides.

Immunohistochemistry

Immunohistochemical detection of ghrelin cells using rabbit anti-ghrelin serum (no. 603) was carried out by the

avidin–biotin–peroxidase complex (ABC) method. The production and specificity of the anti-ghrelin serum used in this study have been reported previously (Kojima *et al.* 1999), and this antiserum recognizes the *n*-octanoylated side chain of the N-terminal region of ghrelin (Date *et al.* 2000, Hosoda *et al.* 2000a). Immunohistochemical staining was performed basically according to the previously reported procedure (Sakai *et al.* 1994). Briefly, the sections were deparaffinized with xylene and rehydrated through descending concentrations of ethanol. Next, the sections were treated with 0.5% sodium metaperiodate to block endogenous peroxidase for 15 min at room temperature and then incubated with 1% normal horse serum and 0.4% Triton X-100 in PBS for 1 h. After washing with PBS for 15 min, the sections were incubated with anti-ghrelin serum diluted 1:150 000 in PBS for 15 h in a humid chamber. After washing with PBS, a second incubation with biotin-conjugated anti-rabbit IgG serum (Vectastain ABC kit; Vector, Burlingame, CA, USA), diluted 1:300 with PBS, was carried out for 1 h, and this was followed by further washing with PBS. Finally, the sections were incubated for 30 min with an avidin–biotin–peroxidase complex (Vectastain ABC kit) prepared according to the manufacturer's instructions. After washing with PBS for 15 min, the sections were reacted in 0.02% 3,3'-diaminobenzidine-tetrachloride mixed with 0.006% H₂O₂ in 0.05 M Tris-HCl, pH 7.6, for 1–3 min to detect immunostaining. After washing with distilled water, the sections were dehydrated with a graded ethanol series, cleared in xylene, mounted with Entellan (Merck, Darmstadt, Germany), and viewed under a light microscope (BX60; Olympus, Tokyo, Japan). All the incubations were carried out in a humidity chamber at room temperature.

In situ hybridization

The sections were deparaffinized with xylene, rehydrated through descending concentrations of ethanol, and washed in diethylpyrocarbonate (DEPC)-treated PBS for 15 s twice. Next, the sections were washed in DEPC-treated PBS, treated with 8.0 µg/ml proteinase K for 30 min at 37 °C, and fixed with 4% paraformaldehyde in 50 mM DEPC-treated PB, pH 7.4. After washing in PBS for 1 min, the sections were incubated with 0.2 M HCl in DEPC water and then washed in PBS for 1 min. The sections were treated with 0.1 M triethanolamine-HCl, pH 8.0, for 1 min and 0.25% acetic anhydride in 0.1 M triethanolamine for 10 min, washed in PBS for 1 min, and then immersed in a graded ethanol series (70, 80 and 90%) for 15 s each. Next, the sections were treated with 100% ethanol for 15 s twice and dried for 20 min. Finally, 2 ng/µl Digoxigenin (DIG)-labeled anti-sense and sense RNA probes (position 112–458; GenBank no. AB029433), synthesized from stomach cDNA using a labeling kit (Boehringer Mannheim GmbH, Mannheim,

Germany), were diluted in hybridization buffer and dropped onto the tissue sections. The sense RNA probe was used as negative control. The sections were covered with Parafilm (American Can Company, CT, USA) and incubated for 16 h at 42 °C in a humidity chamber. The covers were removed by soaking the slides in $5 \times$ SSC and immersed in $2 \times$ SSC containing 50% formamide for 30 min. The sections were then treated with TNE (10 mM Tris-HCl, pH 7.6, 500 mM NaCl, 1 mM EDTA, pH 8.0) for 10 min and next with RNase A (50 µg/ml in TNE) for 30 min at 37 °C. The sections were immersed in TNE for 10 min at 37 °C and washed with $2 \times$ SSC for 20 min at 55 °C and then with $0.2 \times$ SSC for 20 min, twice each. The sections were incubated for 5 min in buffer-1 (100 mM Tris-HCl, pH 7.5, 150 mM NaCl, 0.01% Tween 20), immersed in 1.5% blocking reagent (Boehringer Mannheim GmbH) in buffer-1 for 1 h at 37 °C, and then washed in buffer-1 for 5 min. After washing, the sections were incubated with an alkaline phosphatase-conjugated anti-DIG antibody (Roche Diagnostics Corporation, Indianapolis, IN, USA) diluted 1:2000 in buffer-1. The sections were washed in buffer-1 for 15 min twice and in buffer-2 (100 mM Tris-HCl, pH 9.5, 100 mM NaCl, 50 mM MgCl₂) for 3 min. A chromagen solution (337 µg/ml 4-Nitroblue tetrazolium chloride, 175 µg/ml 5-Bromo-4-chloro-3-indoyl-phosphate in buffer-2) was added, and the sections were incubated until a visible signal was detected. The reaction was stopped by adding a reaction stop solution (10 mM Tris-HCl, pH 7.6, 1 mM EDTA, pH 8.0), and the sections were immersed in 4% paraformaldehyde in 0.05 M PB for 120 min. The sections were then washed with distilled water, dehydrated with a methanol series, cleared in xylene, mounted with Entellan, and viewed under a light microscope (BX60).

Northern blot analysis

Total RNA was extracted from the stomach body by the acid guanidium thiocyanate-phenol-chloroform method. Electrophoresis of 10 µg total RNA was carried out in 1.0% agarose formalin gel. RNA in the gel was transferred onto a sheet of nylon membrane (Gene Screen; NEN, Boston, MA, USA) by capillary action. The membrane was prehybridized for 1 h at 65 °C in a hybridization buffer. After hybridization for 16 h at 65 °C with 1 ng/µl of an antisense DIG-labeled rat ghrelin cRNA probe, the same probe as that used in ISH, and 1 ng/µl of an antisense DIG-labeled glyceraldehyde 3-phosphatase dehydrogenase (GAPDH) RNA probe (position 242–679; GenBank no. NM 017008), the membrane was washed twice with 0.1% SDS in $2 \times$ SSC ($1 \times$ SSC: 16.6 mM sodium chloride, 16 mM sodium citrate) buffer at room temperature for 5 min and immersed twice in 0.1% SDS in $0.2 \times$ SSC buffer at 65 °C for 15 min. The membrane was incubated in 1.5% blocking reagent in buffer-1 for

30 min at 37 °C. After washing, the membrane was incubated with an alkaline phosphatase-conjugated anti-DIG antibody (Roche Diagnostics Corporation) diluted 1:10 000 in buffer-1. A chemiluminescent assay was used to visualize the probe-target hybrids. The membrane was washed in buffer-1 for 15 min twice and then in detection buffer (100 mM Tris-HCl, pH 9.5, 100 mM NaCl) for 5 min. The membrane was placed inside a plastic sheet and CDP-star (Roche Diagnostics Corporation, Indianapolis, USA) was applied and incubated for 5 min. Finally, the membrane in the sealed envelope was exposed to X-ray film (Eastman Kodak Company, Rochester, NY, USA). After exposing, the X-ray films were developed with Kodak developer D-19 (Eastman Kodak Company), and the signals obtained were analyzed using Scion Image software (Scion Corporation, MD, USA). In the analysis, values were normalized by the amount of GAPDH mRNA.

Morphometric analysis

The densities of ghrelin cells in the stomach were estimated. After taking digital photographs under a light microscope (BX60) with a digital camera (Coolpix950; Nikon), the number of ghrelin cells in each section was counted and the area of the mucosal layer in each section was measured using a computerized image analysis program, Scion Image. The ghrelin cell density was calculated as the number of immunopositive mucosal cells per unit area. All the data are expressed as means $\pm$ s.e.

Results

Northern blot analysis

Ghrelin mRNA expressions in the stomach body were detected just after birth in both male and female rats, and then expression levels gradually increased with transient stagnation until 4 weeks of age in the male rats (Fig. 1A) and until 5 weeks of age in the female rats (Fig. 1B). The largest increase (about 200%) in ghrelin mRNA in the stomach was found between 1 and 2 weeks of age in both sexes. In the male rats, the stomach ghrelin mRNA level was highest at 8 weeks of age (Fig. 1A) but, in the female rats, the highest level was found at 7 weeks of age, and the mRNA level had significantly declined at 8 weeks of age (Fig. 1B).

Distribution of ghrelin-immunopositive and ghrelin mRNA-expressing cells

Ghrelin-immunopositive cells (ghrelin-ip cells) and ghrelin mRNA-expressing cells (ghrelin-ex cells) were observed to be closed-type cells as previously reported (Sakata *et al.* 2002) (Fig. 2), and these cells showed the

Figure 1 Changes in ghrelin mRNA levels during the postneonatal period in the rat stomach as determined by Northern blot analysis. The top panel shows representative results of Northern blot analysis of ghrelin and GAPDH mRNA in the rat stomach. The data in (A) and (B) are shown as the % of 1-week-old (1W) rats (100%) and each bar represents the mean $\pm$ S.E. ($n=3$). (A) Ghrelin mRNA levels in male rats. Stomach ghrelin mRNA levels increased overall with age but showed transient stagnation at 4 weeks of age, and the amount of stomach ghrelin mRNA was highest at 8 weeks of age. (B) Ghrelin mRNA levels in female rats. As in the male rats, stomach ghrelin mRNA levels increased with increasing age but showed transient stagnation at 5 weeks of age. The amount of female stomach ghrelin mRNA peaked at 7 weeks of age and was lower at 8 weeks of age.

same localization in the stomach during all stages studied (up to 8 weeks of age). Ghrelin-ip and -ex cells were also found in the mucosal layer but not in the myenteric plexus. Ghrelin-ip and -ex cells were found just after birth in both sexes. The distributions of ghrelin-ip and -ex cells in the stomach changed; ghrelin cells were observed in the glandular base until 2 weeks of age (Figs 2A and D and 3A and D) but were localized in the glandular base to the body of the fundic gland at 3 weeks of age (Fig. 2B and E and Fig. 3B and E). It is therefore thought that ghrelin-ip and -ex cells had spread with increasing age from the glandular base to the neck of the fundic gland (Fig. 2C and F and Fig. 3C and F). The ghrelin cell density in the male fundic gland showed an age-dependent increase after birth (Fig. 4), whereas the cell density in the female fundic gland remained almost constant during the period from 1 to 8 weeks after birth (Fig. 5). Moreover, cell densities in female rats were 2- to 2.5-fold higher at 1

week after birth, and the densities of ghrelin-ip and -ex cells were found to be significantly higher in the female rats than in the male rats at least until 3 weeks of age (data not shown).

Two kinds of stained ghrelin-ex cells, weakly or strongly stained cells, were found in the female rats at the early stage of development (Fig. 3D and E). Interestingly, the ratios of these cells in the male and female rats were markedly different, and numerous weakly stained cells were prominent in the female rats (Fig. 3D and E). The existence ratio of these cells in the female rats decreased with growth, and few weakly stained ghrelin-ex cells were found at 7 weeks of age (Fig. 3F).

Discussion

It is generally accepted that GH release from somatotropes in the anterior pituitary gland is controlled by GHRH and

Figure 2 Microphotographs of ghrelin-ip and -ex cells in the male stomach. (A–C) Ghrelin-ip cells detected by IHC. (D–F) Ghrelin-ex cells detected by ISH. Ghrelin cells were found to be restricted to gastric mucosa. Almost all of the ghrelin cells in the stomach were round-shaped cells, so-called closed-type cells. The distribution of ghrelin-ip cells determined by IHC corresponds well to the distribution of ghrelin-ex cells. (A and D) A few ghrelin cells were found at 1 week (W) of age and were localized in the glandular base. (B and E) Many ghrelin cells were found at 3 weeks in the glandular base to the body of the fundic gland. (C and F) Many ghrelin cells were found at 7 weeks, and they had become scattered from the glandular base to the glandular neck. Although a few weakly stained cells (arrowheads) were found in the early stage (D and E), the staining signals in the cells remained almost constant throughout all stages of development. MU, mucosa; SL, smooth muscle layer. Bars=100 μ m.

Figure 3 Microphotographs of ghrelin-ip and -ex cells in the female stomach. (A–C) Ghrelin-ip cells detected by IHC. (D–F) Ghrelin-ex cells detected by ISH. Distribution pattern of ghrelin cells and cell types are the same as those in the male, but the ghrelin-ip and -ex cells in the female stomach were more numerous than those in the male stomach at all stages. (A and D) Ghrelin -ip and -ex cells were found at 1 week (W) of age and were localized in the glandular base. Note that ghrelin -ip and -ex cells were more numerous than those in the male. (B and E) Ghrelin-ip and -ex cells were found in the glandular base to the body of the fundic gland at 3 weeks. (C and F) A large number of ghrelin cells were found at 7 weeks, and they were scattered through the glandular base to glandular neck. In the female, weakly stained cells (arrowheads) and strongly stained cells (arrows) were observed by ISH (D, E and F). The ISH signals at 1 and 3 weeks of age were shown to be a mixture of different staining intensities, but the staining in most of the cells at 7 weeks was strong (arrows). The ratio of weakly stained cells was higher in the female rats than in the male rats. MU, mucosa; SL, smooth muscle layer. Bars=100 μ m.

Figure 4 (A) Histogram showing the densities of ghrelin-ip cells (cells/mm²) in the male stomach detected by IHC. The number of ghrelin-ip cells increased with age. (B) Histogram showing the densities of ghrelin-ex cells (cells/mm²) in the male stomach detected by ISH. The number of ghrelin gene-expressing cells increased with increasing age but showed transient stagnation at 4 weeks as shown by Northern blot analysis.

SRIF, and these neuropeptides are secreted from the hypothalamus and act on the anterior pituitary gland via the hypophysial portal vein. After the discovery of ghrelin, the mechanisms of GH release should be considered together with the inter-relationship between these peptides. However, among these peptides, ghrelin may have a complex effect on the pituitary gland because, although its content is significantly different from those of other peptides, sources of ghrelin have been reported in various regions, including the stomach, hypothalamus and the pituitary gland itself (Horvath *et al.* 2001). Ghrelin from these different sources may have an effect not only on GH release in adults but also on pituitary gland development. However, the source of ghrelin that is involved in GH release during the period of development has not been determined. Interestingly, the pattern of gastric ghrelin mRNA expression and differentiation of ghrelin-producing cells found in this study are significantly different from the previously reported pattern of pituitary ghrelin mRNA expression, i.e. ghrelin mRNA level in the pituitary gland being highest in the fetal stage then decreasing with increasing age (Kamegai *et al.* 2001). The gastric ghrelin mRNA levels in the present study showed

a completely opposite pattern; lowest just after birth and increasing with increasing age. On the other hand, it has been reported that the pituitary GHS-R mRNA expression level was significantly higher just after birth and at 4–5 weeks of age (Kamegai *et al.* 1999), and the periods of these two peaks are consistent with the increases in pituitary and gastric ghrelin expression levels respectively. Taken together, these results suggest that GH secretion from somatotropes is controlled by ghrelin from different origins in the various stages of development.

It has been reported that plasma acylated ghrelin is very unstable (Ariyasu *et al.* 2001). In addition, ghrelin from the stomach may be degraded rapidly by passing from the portal vein to the liver. Therefore, it is questionable whether or not gastric ghrelin directly stimulates GH release from the pituitary gland via physiologically active plasma ghrelin. Moreover, the results of our study using a perfusion system demonstrated that GHRH stimulates GH release from the anterior pituitary gland more strongly (about ten times more strongly) than does ghrelin (Yamazaki *et al.* 2002). These results suggest that gastric ghrelin in the adult does not directly stimulate somatotropes but has an effect on GH release by an

Figure 5 (A) Histogram showing the densities of ghrelin-ip cells (cells/mm²) in the female stomach detected by IHC. The number of ghrelin-ip cells increased rapidly from 1 to 2 weeks of age and then became almost constant. (B) Histogram showing the densities of ghrelin-ex cells (cells/mm²) in the female stomach detected by ISH. The density of ghrelin cells remained almost constant throughout the period of development. Note that the densities of ghrelin cells in the female rat were higher than those in the male rats at all stages (compare with Fig. 4).

indirect route. Recently, many studies have demonstrated that ghrelin also regulates food intake, gastric acid secretion and gastric motor activity via vagal control systems (Masuda *et al.* 2000, Asakawa *et al.* 2001a, Date *et al.* 2001). Considering these findings, it is not surprising that plasma ghrelin stimulates GH secretion via the vagal nerve and has indirect effects through neural pathways. Elucidation of the sites of action of stomach ghrelin and changes in stomach ghrelin during development may be needed to understand the physiological role of ghrelin.

Recently, several studies have demonstrated ontogeny of stomach ghrelin gene expression. For example, Tanaka *et al.* (2001) reported that no significant changes in the amount of ghrelin mRNA in the mouse stomach was observed during the period 2–8 weeks after birth. On the other hand, Lee *et al.* (2002) reported that rat gastric ghrelin mRNA increased progressively through the second and third weeks after birth and that the ghrelin mRNA level in the stomach thereafter remained constant until the end of the nursing period. In addition, Gualillo *et al.* (2001) reported that gastric ghrelin gene expression in the rat increased with increasing age (up to 90 days). In

this study, we found that ghrelin gene expression increased in a stepwise manner, a finding that is almost in agreement with the results of the latter two reports mentioned above. Notably, although their study was not continued after 23 days of age, Lee *et al.* (2002) reported that rat ghrelin gene expression peaked at 18 days of age and then decreased until 22 days of age. This change in gene expression is in good accordance with that of the first peak of our results. Although further studies are needed to elucidate the physiological significance of ghrelin mRNA fluctuation, it should be pointed out that the first increase in the expression level is closely related to the weaning period or rapid body growth period. Moreover, this period is almost consistent with the onset of puberty.

In the male rats, the densities of ghrelin-ip and -ex cells increased gradually, and the increase paralleled that in ghrelin mRNA expression level. On the other hand, in the female rat stomach, although numerous ghrelin-ex cells were found in neonates and cell densities remained almost constant throughout development, ghrelin mRNA expression showed a stepwise increase, suggesting that changes in ghrelin gene expression did not reflect the

morphological results. For this reason, since we counted ghrelin-ex cells without taking into consideration the staining intensity in this study, even weakly stained cells were considered as ghrelin gene-expressing cells. However, as pointed out in the Results section, many weak signals were found by ISH in the female rats, and these cells were replaced with strongly stained cells with the advance of the process of development, indicating that the ghrelin mRNA expression level increases after birth. These results suggest that ghrelin-producing cells in female rats differentiate at an earlier stage of development than they do in male rats. The difference between the stages of development at which ghrelin-producing cells differentiate in male and female rat stomachs is a very interesting issue, and this difference suggests that ghrelin has distinct physiological effects on male and female development.

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