

The Expression Analysis of Ghrelin and Growth Hormone Secretagogue Receptor in Mouse Thymic Epithelial Cells

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Abstract: The present study was to identify the expression of Ghrelin and GHS-R in mouse Medullary-Type Epithelial Cells (MTEC1). Whether the gene of ghrelin and GHS-R are expressing in MTEC1 cells or not was measured by using Quantitative real time RT-PCR (Q-PCR). Results identified that the mRNA of ghrelin and GHS-R can be detected in MTEC1 cells. The protein levels of GHS-R and ghrelin precursor were detected by the method of Western blot analysis. Moreover, the location of GHS-R in MTEC1 cells were detected by immunofluorescence and the quantity of ghrelin in the supernatant of MTEC1 cells were measured by ELISA. The results of Western blot analysis make sure that GHS-R and precursor of ghrelin are expressing in MTEC1 cells. GHS-R is a kind of transmembrane protein. As the results of immunofluorescence shown the quantity of ghrelin in the supernatant of MTEC1 cells measured by ELISA was 39.20 pg mL^{-1} while the control was 5.74 pg mL^{-1} .

Key words: Ghrelin, GHS-R, MTEC1 cells, expression, prokin, China

INTRODUCTION

Ghrelin is a peptide hormone and is firstly found in the rat stomach in 1999. The Open Reading Frame (ORF) of mouse ghrelin mRNA is 354 bp and the deduced protein contained 117 amino acid (aa) which is the precursor of ghrelin (Kojima *et al.*, 1999). After translation, precursor of Ghrelin was spliced and mature Ghrelin is a 28 aa peptide. Ghrelin is mainly produced by the stomach whereas substantially lower amounts are derived from the bowel, pituitary, kidney and placenta (Date *et al.*, 2000; Hosoda *et al.*, 2000; Ariyasu *et al.*, 2001; Korbonits *et al.*, 2001). The earlier studies have demonstrated that ghrelin plays a key role in response to negative energy balance and promotes Growth Hormone (GH) secretion from the pituitary via binding to its 7-Transmembrane G Protein-Coupled Receptor (7-TM GPCR), the GH Secretagogue Receptor (GHS-R) (Kojima *et al.*, 1999; Smith, 2005).

The GHS-R of human and swine including two distinctive cDNAs which are GHS-R type 1a and GHS-R type 1b (Howard *et al.*, 1996). GHS-R type 1a encodes a 7-TM GPCR with binding and functional properties consistent with its role as ghrelin's receptor while GHS-R type 1b encodes a 5-TM GPCR with no measurable function (Howard *et al.*, 1996). There is no reports

identified the GHS-R of mouse can be divide into any subtype until now on. The mRNA of GHS-R can be detected through RT-PCR in many peripheral organs including heart, lung, liver, kidney, pancreas, stomach, small and large intestines, adipose tissue and immune cells (Guan *et al.*, 1997; Hattori *et al.*, 2001; Gnanapavan *et al.*, 2002) which indicates that ghrelin may have multiple functions in these tissues.

Ghrelin has the ability to promote thymopoiesis during aging (Dixit *et al.*, 2007). In the thymus, thymocyte development is dependent upon the sequential contribution of morphologically and phenotypically distinct stromal cell compartments that comprise the thymic microenvironment (Anderson *et al.*, 2000; Anderson and Jenkinson, 2001; Kyewski *et al.*, 2002). The thymic epithelial cells are a major component of the stromal cells of the thymus and provide a crucial signal for intrathymic T cell development and selection (Rouse *et al.*, 1988; Ritter and Palmer, 1999). The earlier studies have demonstrated that ghrelin and its receptor played a key role in the regulation of thymic restore (Dixit *et al.*, 2007).

In order to research the mechanism of the role of ghrelin and its receptor in regulation of thymic development, researchers measured the expression levels of ghrelin and GHS-R in cultured MTEC1 cells.

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MATERIALS AND METHODS

Cell culture: MTEC1, a mouse (*Mus musculus*) thymic medullary-type epithelial cell line supporting the unctional maturation of CD4 single-positive thymocytes *in vitro* (Ge and Chen, 2000) was kindly provided by Dr. Yu Zhang (Department of Immunology, Peking University Health Science Center). These cells were cultured in Dulbecco’s Modified Eagle’s Medium (Gibco) containing 10% fetal bovine serum (Fbs; Gibco).

Primers and antibody: To investigate the expression level of ghrelin and Growth Hormone Secretagogue Receptor (GHS-R), the primers, GHS-R-F, GHS-R-R Ghrelin-F and Ghrelin-R were designed and synthesized based on the nucleotide sequences of GHS-R (NM_177330) and ghrelin (NM_021488) (TaKaRa Biotechnology (Dalian China) Co., Ltd.). The primers used in Q-PCR are shown in Table 1. The primary antibody, of GHS-R (sc-10359) used in Western blot and immunofluorescence was obtained from Santa Cruz Biotechnology (Santa Cruz Biotechnology, Inc. California USA). The antibody for Western blot analysis of Ghrelin was purchased from Phoenix Pharmaceuticals, Inc. USA.

Quantitative real time RT-PCR (Q-PCR) assays: MTEC1 cells (1.0×10^6) were harvested at the indicated time points and washed twice with ice-cold PBS. Then, the total RNA (1 μ g) was isolated with the TRIZOL reagent (Invitrogen). Reaction mixture (20 mL) containing 1 μ g of total RNA was reverse transcribed to cDNA using PrimeScript RT-polymerase (Takara). Q-PCR was performed on the cDNA using primers specific for GHS-R and ghrelin, specifically. RNA input was normalized to the level of GAPDH for genes analysis. All reaction were carried out using SYBR Green Mix (Takara) and the PCR conditions for Q-PCR were as follows: activation of enzyme at 94°C for 1 min, 40 cycles of denaturation at 94°C for 5 sec, annealing at 60°C for 30 sec and extension at 72°C for 30 sec. Q-PCR was carried out using a ABI PRISM® every cycle. The fluorescence of samples was continuously traced during this period. Relative expression levels were calculated as ratios

normalized against those of GAPDH. These experiments were repeated three times. All results are expressed as the means \pm SD of three independent experiments.

ELISA assays: The supernatant of MTEC1 cells is prepared ELISA was carried out according to the manufacturer’s instruction (Uscn Life Science, USA. E90991 Mu). Finally, measurement was conduct at 450 nm immediately. The control group was culture medium.

Western blot analysis: MTEC1 cells (3.0×10^6) were harvested, washed twice with ice-cold PBS and lysed in 0.5 mL of lysis buffer. Equal amounts (60 mg) of proteins were loaded and resolved by Sodium Dodecylsulfate polyacrylamide Gel Electrophoresis (SDS-PAGE) and then transferred onto immobilon membrane (Millipore). The membrane was blocked with 5% non-fat milk in Tris-Buffered Saline (TBS, pH 7.4) for 1 h and then incubated with antibodies (GHS-R: 1:1000, Ghrelin: 1:500) in 5% non-fat milk in TBS. After the incubation of primary antibody, the membrane was washed with TBS including 5% Tween-20 (TBST) and incubated with secondary antibodies conjugated with horseradish peroxidase in 5% non-fat milk in TBS. After washing with TBST, the bound antibodies were visualized by immobilon Western chemilluminiscent HRP substrate (Millipore) and recorded on X-ray films.

Immunofluorescence: MTEC1 cells were grown on cover glasses in six-well plates and were fixed with 4% formaldehyde for 30 min, permeabilized with 0.2% Triton X-100 for 5 min. Then cells blocked with 3% BSA at 37°C for 2 h. Following, primary antibody for GHS-R (Phoenix Pharmaceuticals, Inc. USA) was deliquation according directions and incubated cells at 4°C overnight. The secondary antibody (1:2,000, Alexa 594-conjugated anti-mouse IgG) applied for 1 h at room temperature. And the cells was incubated in 0.1-1 mg mL⁻¹ DAPI (DNA stain) for 1 min. Triple washing with PBS was performed between each step. Finally, the coverslips were mounted in PPD-mounting medium (90% glycerol). Images were taken using Fluorescence microscope (Leica., Germany).

Table 1: The primers used in this study

Abbreviation	Primer sequence (5'-3')	Description
GHS-R-F	ATCACCTCTGGGCTCTTGTTGCTG	Quantitative RT-PCR primer, forward
GHS-R-R	GCTGAATGGCTCATTGTAGTCCTG	Quantitative RT-PCR primer, reverse
Ghrelin-F	TTGGCATCAAGCTGTCAAGGAG	Quantitative RT-PCR primer, forward
Ghrelin-R	GTCAATGGCCTGTCCGTGGTTA	Quantitative RT-PCR primer, reverse
Gapdh-F	GTGTCCGTCGTGGATCTGA	Quantitative RT-PCR primer, forward
Gapdh-R	TTGCTGTTGAAATCGCAGGAG	Quantitative RT-PCR primer, reverse

RESULTS AND DISCUSSION

mRNA expression level: Q-PCR was carried out as described in material and method. As shown in Fig. 1, the mRNA of the ghrelin and GHS-R were both detected in MTEC1 cells. The average threshold cycle of ghrelin mRNA is 25.06 and the average threshold cycle of GHS-R mRNA is 25.18. Comparatively, the control gene, *GAPDH* whose average threshold cycle is 14.46 was also been tested.

Protein expression profile: To investigate the expression of ghrelin and its precursor in MTEC1 cells *in vitro*, researchers make use of ELISA to detect the quantity of ghrelin and Western blot was performed to detect the precursor of ghrelin. ELISA was carried out according to the manufacturer’s instruction. The concentration of ghrelin in the supernatant of MTEC1 cells is 39.20 pg mL⁻¹ while the control is 5.74 pg mL⁻¹ according to the standard curve shown in Fig. 2. And the protein of ghrelin precursor was also been detected in MTEC1 cells (Fig. 3).

In this study, Western blot and immunofluorescence was used to detect the expression of GHS-R. The result

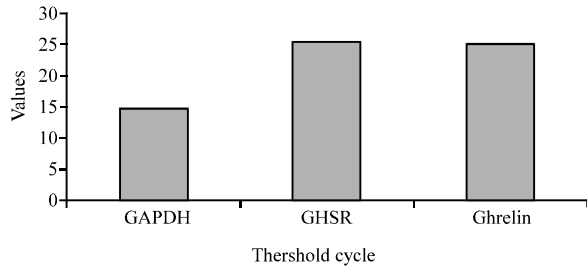


Fig. 1: Ghrelin and GHS-R expression levels in mouse MTEC1 cells were detected by Q-PCR analysis. GAPDH serves as internal control. The bars indicate mean±SD from three different experiments

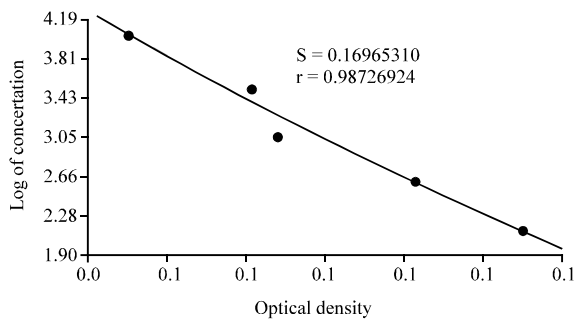


Fig. 2: Typical standard curve for mouse ghrelin ELISA analysis

indicated that the expression of GHS-R can be measured in MTEC1 cells and it is a kind of transmembranes protein as shown in Fig. 3. In addition, the expression of GHS-R was detected by Western blot analysis as it’s shown in Fig. 4.

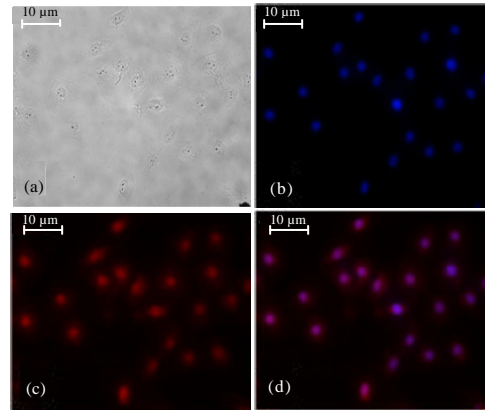


Fig. 3: Immunofluorescence analyses of expression levels of GHS-R in mouse MTEC1 cells. a) photos of MTEC1 taken on natural light; b) the nuclei stained blue fluorescence; c) GHS-R stained red fluorescence and d) merge the nuclei and GHS-R

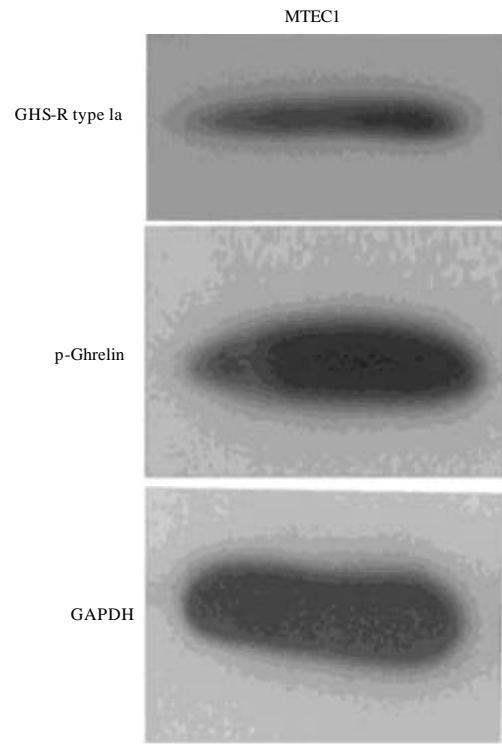


Fig. 4: Western blot analysis of GHS-R and precursor Ghrelin (p-Ghrelin) in mouse MTEC1 cells

The thymus is critical for the development, selection and maintenance of the peripheral T cell pool possessing a broad spectrum of TCR specificities (Dixit *et al.*, 2007). Moreover, thymic epithelial cells provide a crucial signal for intrathymic T cell development and selection. And ghrelin has been reported to have the function of activating human T cells and mediating anti-inflammatory signals in several cells types.

CONCLUSION

By this study, researchers firstly investigated the expression of ghrelin and GHS-R in the MTEC1 cells *in vitro*. The results indicated that the mRNA and protein of Ghrelin and GHS-R can both be detected in the MTEC1 cells. Based on the results of present study, researchers can investigate the role of ghrelin and GHS-R in the MTEC1 in further based on this model.

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