



Effect of the IGF-1/PTEN/Akt/FoxO signaling pathway in the duodenal mucosa of rats subjected to water immersion and restraint stress

P. Huang^{1,2}, Z.R. Zhou¹, M.Q. Zheng¹ and F.X. Shi¹

¹Laboratory of Animal Reproduction, College of Animal Science and Technology, Nanjing Agricultural University, Nanjing, China

²Department of Pathology, School of Medical Science and Laboratory Medicine, Jiangsu University, Zhenjiang, China

Corresponding author: F.X. Shi

E-mail: fxshi@njau.edu.cn / njauhp@gmail.com

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ABSTRACT. The insulin growth factor 1/phosphatase and tensin homologue deleted on chromosome 10/Akt/forkhead box (IGF-1/PTEN/Akt/FoxO) signaling pathway reportedly exhibits gastroprotective effects by reducing water immersion and restraint stress (WRS)-induced gastric mucosal cell apoptosis. We examined the expression and localization of IGF-1, PTEN, Akt, and FoxO proteins, caspase-3 activity, and the number of apoptotic cells in the duodenal mucosa of rats subjected to WRS to confirm whether the IGF-1/PTEN/Akt/FoxO signaling pathway has a role in the duodenal mucosa. The results indicated that WRS enhanced cell apoptosis in the duodenal mucosa. In addition, in normal rats, PTEN was found mainly in the cellular cytoplasm of the duodenal glands and lamina propria of villi. IGF-1 and total Akt were observed in the cellular cytoplasm of the duodenal glands. In addition, total Akt was found in the cellular cytoplasm of the myenteric plexus. FoxO3a and FoxO4 were primarily concentrated in the cellular cytoplasm of the lamina propria. Specifically, PTEN, FoxO3a and FoxO4 were also localized in the cellular cytoplasm of lamina propria of restituted villi in

the duodenal mucosa of rat subjected to WRS. In addition, messenger RNA transcript levels of *IGF-1*, *PTEN*, *Akt1*, *Akt2*, *FoxO3*, and *FoxO4* were upregulated in the duodenal mucosa, with a peak between the 4th and 8th day after 7 h of WRS. Furthermore, the results also suggested that *Akt3* messenger RNA transcript levels in the duodenal mucosa of rats after WRS showed no significant differences compared with those in the non-WRS group. Collectively, our results implied that the IGF-1/PTEN/Akt/FoxO signaling pathway was effective in regulating cellular apoptosis in the duodenal mucosa of rats after WRS.

Key words: IGF-1; FoxO; Signaling pathway; Duodenum; Rat

INTRODUCTION

The water immersion and restraint stress (WRS) rat has consistently been used as an animal model of gastroduodenal mucosal lesions (Wang and Johnson, 1991; Adachi et al., 2011). The insulin growth factor 1/phosphatase and tensin homologue deleted on chromosome 10/Akt/forkhead box (IGF-1/PTEN/Akt/FoxO) signaling pathway exerts important physiological impacts on many types of animal cells (Carnero et al., 2008; Reddy et al., 2008) and has been reported to exhibit gastroprotective effects by reducing WRS-induced gastric mucosal cell apoptosis (Nguyen et al., 2007; Zhao et al., 2009). The gastrointestinal tract has been identified as one of the most sensitive target tissues for IGF-1, which has critical biological functions, including the promotion of the differentiation of various cell types, gastrointestinal tract development, and potent anti-apoptotic activity (Warzecha et al., 2006; Karcher et al., 2009; Reyna et al., 2010). In particular, recent studies have demonstrated gastric ulceration-triggered upregulation of IGF-1 expression in gastrointestinal ulcer margins, suggesting that IGF-1 is involved in gastrointestinal repair processes (Cool et al., 2005; Ceranowicz et al., 2009).

The transduction of signals through the IGF-1 receptor triggers a series of multiple intracellular phosphorylation events that prevent cell death (Carroll, 2001). The phosphatidylinositol 3-kinase (PI3K)/Akt pathway predominantly activated by IGF-1 is a strong cell survival cascade. Akt [also called protein kinase B (PKB)] is a serine/threonine protein kinase downstream of PI3K (Song et al., 2005) and an important regulator of cell proliferation, growth, and survival (Cully et al., 2006). To date, 3 members of the Akt family - Akt1, Akt2, and Akt3 - have been isolated. Although they are products of different genes, they are closely related to one another, with up to 80% amino acid sequence identity. The 3 genes are expressed differentially, with a broader expression for Akt1 and Akt2 and a more restricted expression for Akt3 (Fresno Vara et al., 2004). Previous studies have demonstrated that PI3K/Akt plays critical roles in several gastrointestinal diseases (Coeffier et al., 2011; Tuo et al., 2011). Furthermore, IGF-1 has been reported to reduce WRS-induced gastric mucosal injury by inhibiting the gastric accumulation of neutrophils through the reduction of caspase-3 activation by PI3K/Akt signaling (Nguyen et al., 2007; Zhao et al., 2009).

PTEN is a tumor-suppressor gene that encodes a dually specific phosphatase that recognizes both lipid and peptide substrates, including phosphatidylinositol (3,4,5)-trisphosphate, a product of PI3K. Through its lipid phosphatase activity, PTEN controls Akt signaling and its downstream targets responsible for cell size, cell migration, cell cycle, cell death, and

focal adhesion formation (Leslie and Downes, 2004). A downstream target of IGF-1/PTEN/Akt signaling is the O subfamily of FoxO proteins, which is phosphorylated and thereby inhibited by activated Akt. Four members of the FoxO subfamily - FoxO3a, FoxO1, FoxO4, and FoxO6 - are reported to play key roles in mammalian cells. Phosphorylation of FoxO proteins by Akt results in cytoplasmic retention and inactivation and inhibits the expression of FoxO-regulated genes that control the cell cycle, cell death, and cell metabolism. The shuttling of FoxO between the cytoplasm and the nucleus is a key step in cell apoptosis (Burgering and Kops, 2002). Recently, we have demonstrated cell-specific and age-dependent expression patterns of FoxO4 and FoxO3a proteins in the duodenum and their involvement in the development and growth performance of rat duodenum (Huang et al., 2011). We have also found that FoxO4 is a primary transcriptional factor localized in the gastrointestinal tracts of pigs (Zhou et al., 2007). In particular, our previous studies have suggested that the IGF-1/PTEN/Akt/FoxO signaling pathway plays some role(s) in protecting against ulcers by regulating cellular apoptosis in the development and healing of rat gastric ulcers (Huang et al., 2012).

The aim of the present study was to determine whether the IGF-1/PTEN/PI3K/Akt/FoxO signaling pathway is involved in protecting against WRS-induced duodenum mucosal lesions. In the WRS rat model, we analyzed the expression and localization of IGF-1, PTEN, Akt, and FoxO with immunohistochemistry and real-time polymerase chain reaction (PCR), respectively. In addition, we investigated cell apoptosis in WRS-induced duodenal mucosal lesions using the terminal deoxynucleotidyl transferase 2'-deoxyuridine, 5'-triphosphate (dUTP) nick-end labeling (TUNEL) method and caspase-3 activity measurement.

MATERIAL AND METHODS

WRS-induced duodenum mucosal lesion formation in rats

All experiments were carried out in intact male Sprague-Dawley rats (9-11 weeks old; Qinglongshan Experimental Animal Breeding Farm, Nanjing, China) weighing 200-220 g. All procedures were designed in accordance with generally accepted ethical standards for animal experimentation and the guidelines established by the institutional animal care and use committee of Nanjing Agricultural University. The uniform commercial diets used in the experiment were also purchased from Qinglongshan Experimental Animal Breeding Farm. Regular rat chow and tap water were allowed *ad libitum*. Rats were housed individually at room temperature (25°C) with a 12-h:12-h light/dark cycle and humidity of 65-70%. Before each experiment, animals were deprived of food but not water for 24 h. The animals were then placed in a restraint cage and immersed in a water bath (20°-22°C) to the level of the xiphoid process as described elsewhere (Adachi et al., 2011). Some of the animals were killed after 3 and 7 h of WRS, and the rest were fed normally starting 1 h later and killed at various time points (4, 8, and 15 days) after the end of the 7-h WRS treatment. The animals were anesthetized with an intraperitoneal injection of ether. Their duodenums were removed and filled with 2 mL 1% formalin and immersed in 1% formalin for 24 h.

Assessment of apoptotic cell number

After transfer through a graded series of alcohol and xylene, the duodenum mucosal samples were embedded in paraffin and sectioned in 7- μ m thicknesses. The TUNEL method was per-

formed using an Apoptosis Kit Direct (Beyotime Institute of Biotechnology, Haimen, China) as described elsewhere (Kelly et al., 2003). Briefly, duodenum mucosal cells were counterstained with 2-(4-amidinophenyl)-1H-indole-6-carboxamide dihydrochloride (Beyotime Institute of Biotechnology) to label all nuclear DNA, and fragmented DNA was end-labeled with fluorescein isothiocyanate-labeled dUTP using terminal transferase. The duodenal mucosa was then observed by using a fluorescence microscope. Sections that were pretreated with DNase I to nick all DNA served as positive controls. For negative controls, dUTP was omitted, resulting in uniformly negative staining. Ten optical fields - approximately 500-1000 cells - were counted in each slide under high power (400X) microscopy, and the number of positive cells per field was expressed as the apoptotic index. These experiments were performed in triplicate with 6 rats per group per experiment.

Measurement of caspase-3 activity

The activity of caspase-3 was detected using a commercially available caspase-3 activity kit (Beyotime Institute of Biotechnology) with Ac-DEVD-pNA (acetyl-Asp-Glu-Val-Asp *p*-nitroanilide) as the colorimetrically specific substrate. In brief, duodenal mucosal samples (N = 6 for each treatment) were weighed and homogenized in lysis buffer containing 10 mM HEPES/KOH, pH 7.2, 2 mM ethylenediaminetetraacetic acid, 0.1% 3-([3-cholamidopropyl]dimethylammonio)-1-propanesulfonate, 5 mM dithiothreitol, 1 mM phenylmethyl-sulfonylfluoride, 10 µg/mL aprotinin, and 20 µg/mL leupeptin. The lysate was centrifuged at 20,000 *g* for 10 min at 4°C, and supernatants were incubated for 7 h at 37°C with 10 µL caspase-3 substrate (Ac-DEVD-pNA, 2 mM). Substrate cleavage was measured with a spectrofluorometer at 405 nm and was corrected as protein content in the lysate. The activity of caspase-3 was expressed as a value of enzyme activity compared with that in the control (Wang et al., 2010).

Immunohistochemical analysis

Antibodies for FoxO3A (Cat. No. 9467, Lot 4), FoxO4 (Cat. No. 9472, Lot 1), and total PKB/Akt (Cat. No. 9292, Lot 1) were purchased from Cell Signaling Technology (Beverly, MA, USA). Antibodies for PTEN (Cat. No. sc-9145, lot C0707) were obtained from Santa Cruz Biotechnology (Santa Cruz, CA, USA). Antibodies for IGF-1 (BA0939) were obtained from Boster Bio-Engineering (Wuhan, China). Streptavidin-biotin complex kits were obtained from BioGenex (San Ramon, CA, USA), and 3,3'-diaminobenzidine tetrachloride was purchased from Sigma Chemical Co. (St. Louis, MO, USA). All other chemicals were purchased commercially and were reagent grade.

After transfer through a graded series of alcohol and xylene, duodenum mucosal samples were embedded in paraffin and sectioned at 7-µm thicknesses. The sample sections were mounted on slides and processed for immunohistochemical analysis, which was conducted using a protocol similar to that described in our previous reports (Ding et al., 2010). Briefly, sections were incubated overnight at room temperature with a polyclonal rabbit immunofluorescence-purified antiserum directed against IGF-1 (1:200), PTEN (1:400), total Akt (1:400), FoxO3a (1:400), and FoxO4 (1:500). The specific protein immunoreactivity was visualized with an Elite ABC Kit and 0.05% 3,3'-diaminobenzidine tetrachloride in 10 mM phosphate-buffered saline containing 0.01% H₂O₂ for 5 min. The specificity of the antibody was examined using normal rabbit serum instead of the primary antibody. To identify structural components and cell morphology, we counterstained the sections with hematoxylin and mounted them with

coverslips. Relative levels of immunostaining between animals and cell types were repeated at least 4 times and evaluated by 3 independent observers.

Total RNA isolation and reverse transcription (RT)

Duodenal mucosa samples were collected and stored in liquid nitrogen until their use for RNA isolation. Total RNA was isolated after homogenizing duodenal mucosa in the Trizol[®] reagent (Invitrogen, CA, USA) following the manufacturer protocol. RNA quality was evaluated by examining a portion on an RNA gel. Bands of 18S and 28S were clear, and slight smearing occurred, indicating that the quality was acceptable. However, the 18S band was not as dark as expected, suggesting that some slight RNA degradation had occurred.

RT reactions were performed using RT reagent kits with gDNA Eraser (Takara, Dalian, China). The total reaction volume of 20 μ L contained 2 μ L 5X gDNA Eraser buffer, 1 μ L gDNA Eraser, 1 μ g total RNA, 4 μ L 5X RT buffer, 1 μ L RT enzyme mix, 1 μ L RT primer mix and sufficient nuclease-free H₂O. The RT reaction was carried out at 42°C for 2 min and 37°C for 15 min and a denaturation step was performed at 85°C for 15 s before the reaction was cooled on ice.

RT-PCR analysis of gene expression

Quantification of all transcripts was performed with RT quantitative PCR using the ABI 7300 PRISM system (Applied Biosystems, CA, USA). PCR products for 9 genes (*IGF-1*, *PTEN*, *Akt1*, *Akt2*, *Akt3*, *FoxO3a*, *FoxO4*, and *HPRT*) were detected with SYBR Green chemistry (Takara). The sequences and GenBank accession Nos. of the primer sets used to amplify the target genes are presented in Table 1. PCRs were run in triplicate in a total volume of 20 μ L (consisting of SYBR Premix Ex Taq, ROX Reference Dye, 200 nM each of sequence-specific primers, and 100 ng equivalent of cDNA). The amplification conditions were as follows: DNA polymerase activation at 95°C for 30 s, followed by 40 amplification cycles at 95°C for 5 s and at 60°C for 31 s. At the end of the amplification cycles, a melting curve analysis was performed to verify specific amplification.

Table 1. Primers used for real-time PCR analysis.

Genes and sequence reference (GenBank No.)	Primer sequence	Size of PCR product (bp)	Annealing temperature (°C)
HPRT (X62085)	F: 5'-AGTGATGATGAACCAGGTTA-3' R: 5'-ATTATAGTCAAGGGCATATC-3'	556	58.0
IGF-1 (BC086374)	F: 5'-TGGTGGACGCTCTTCAGTTC-3' R: 5'-GCTTCAGCGGAGCACAGTAC-3'	168	58.0
PTEN (NM031606)	F: 5'-AGCGTGCGGATAATGACAAG-3' R: 5'-GGATTGATGGCTCCTCTACTG-3'	151	56.0
Akt1 (NM033230)	F: 5'-TAGGCATCCCTTCCCTTACAG-3' R: 5'-GCCCGAAGTCCGTTATCT-3'	269	58.0
Akt2 (NM017093)	F: 5'-GAGCCGAGTCTACAGAATACC-3' R: 5'-GGCCATCTTTGTCCAGCATA-3'	263	58.0
Akt3 (NM031575)	F: 5'-AACGACCAAAGCCAAATACA-3' R: 5'-CCCCATTAACATATTCATCAC-3'	498	58.0
FoxO3a (NM001106395)	F: 5'-TTCGCAACGACCCAATGA-3' R: 5'-TCCAAGCTCCCATGAACAT-3'	331	57.4
FoxO4 (NM001106943)	F: 5'-GGTGCCTACTTCAAGGACAA-3' R: 5'-ATCGGGGTTTCAGCATCCA-3'	148	58.0

The comparative CT method was used for relative quantification of target gene expression levels (ABI Prism Sequence Detection System, Applied Biosystems). The quantity of each measured cDNA sample was normalized to the endogenous gene *HPRT* (a housekeeping gene), and all samples were measured in triplicate. The mean values of the replicate wells for each sample were calculated and divided by the value of *HPRT* to obtain a normalized value for each transcript.

Statistical analyses

Statistical analyses were performed using SPSS 17.0. Values are reported as means \pm standard error of the mean. The data were analyzed using one-way analysis of variance and the Fisher protected least significant difference test. A value of $P < 0.05$ was considered to be statistically significant. All experiments were repeated at least 3 times, and representative data are shown.

RESULTS

Effect of WRS on apoptosis in duodenal mucosa

Duodenal tissue sections were stained using the TUNEL method to determine the quantity and distribution of apoptotic cells and examine nuclear condensation and fragmentation. Sections exposed to DNase I, which causes DNA fragmentation, showed intense staining of all nuclei and were used as positive controls (data not shown). Sections stained using the described procedure but without the terminal deoxynucleotidyl transferase enzyme showed no staining and were used as negative controls (see Figure 1A-C). As shown in Figure 1, WRS induced an absence of several duodenal villi (see Figure 1D-I). In addition, apoptotic cells were observed in the lamina propria (see Figure 1D-R) after WRS. However, in the duodenal mucosa of non-WRS rats, a small amount of labeling was found, mainly concentrated in the lamina propria of villi (see Figure 1S-U). At higher magnification and resolution, the stained cells showed the characteristic morphology of apoptotic cells. The number of apoptotic cells in the duodenal mucosa of the rats subjected to 3 and 7 h of WRS were increased by 13 and 32%, respectively, compared with that in the control group (see Figure 2A; $P > 0.05$, $N = 6$). Subsequently, from the 4th to the 15th day after the 7-h WRS treatment, the number of apoptotic cells in the duodenal mucosa decreased gradually (see Figure 2A).

To reconfirm cell apoptosis in the duodenal mucosa of the WRS rat, caspase-3 activity in the duodenal mucosa was detected using colorimetric analysis. As shown in Figure 2B, caspase-3 activity in the duodenal mucosa after 3 and 7 h of WRS was also time-dependently enhanced by 23 and 31% compared with that in the control group. However, on the 4th, 8th, and 15th day after the 7-h WRS treatment, caspase-3 activity decreased gradually compared with that in the 7-h WRS group.

Immunohistochemical localization of IGF-1, PTEN, total Akt, FoxO3a, and FoxO4 in the duodenal mucosa of rats after WRS

To assess the localization of IGF-1, PTEN, total Akt, FoxO3a, and FoxO4 in rat duodenal mucosa, we stained sections from non-WRS and WRS rat duodenal mucosa with specific antibodies against these proteins. In normal rat duodenal mucosa, PTEN (see

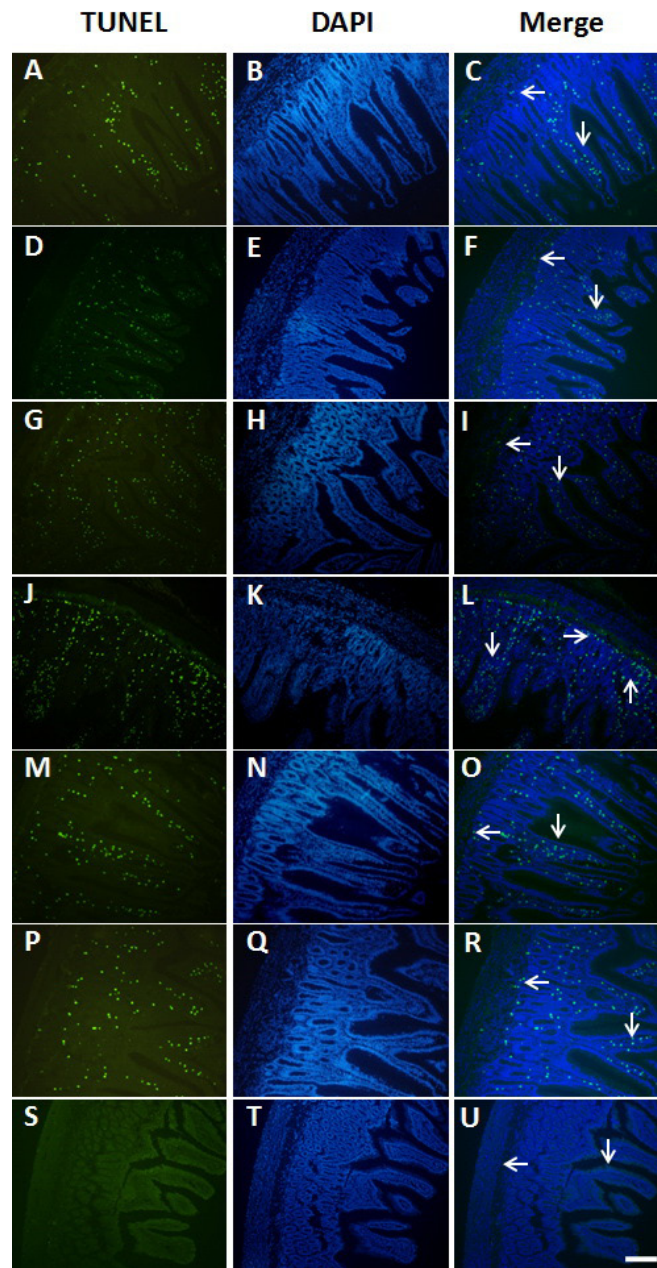


Figure 1. Effect of water immersion and restraint stress (WRS) on apoptotic cells in the duodenal mucosa. First, sections of rat duodenal mucosa were stained using the TUNEL method, and sections were then counterstained with DAPI to label all nuclei. Blue color = DAPI-stained nuclei. Green color = TUNEL-positive cells. The figures indicate non-WRS (A-C), 3 h of WRS (D-F), 7 h of WRS (G-I), 4 days after WRS (J-L), 8 days after WRS (M-O), and 15 days after WRS (P-R). Sections stained by the previous procedure but without use of the TdT enzyme showed no staining and were used as negative controls (S - U). → = muscularis mucosa. ↓ = duodenal villi. Bar = 50 μ m.

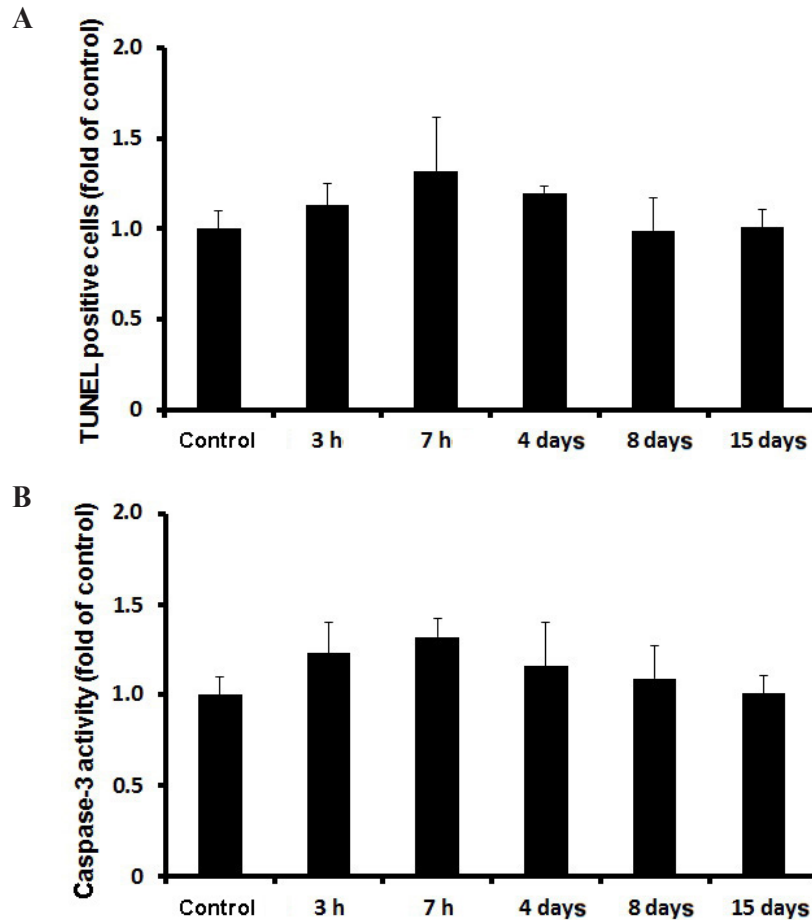


Figure 2. Effect of water immersion and restraint stress on number of TUNEL-positive cells in duodenal mucosa (A) and caspase-3 activity in duodenal mucosa (B) in rats. Data (fold of control) represent means \pm SEM.

Figure 3B) was found mainly in the cellular cytoplasm of the duodenal glands (see Figure 3C) and the lamina propria of villi (see Figure 3D). In addition, IGF-1 (see Figure 3E and F) and total Akt (see Figure 3G and H) were observed in the cellular cytoplasm of the duodenal glands. Total Akt (see Figure 3G and H) was also found in the cellular cytoplasm of the myenteric plexus. FoxO3a (see Figure 3I and J) and FoxO4 (see Figure 3K and L) were primarily concentrated in the cellular cytoplasm of the lamina propria. Notably, in duodenal mucosa of rats subjected to WRS, PTEN (see Figure 3B and D), FoxO3a (see Figure 3I and J), and FoxO4 (see Figure 3K and L) were also localized in the cellular cytoplasm of the lamina propria of restituted villi. However, the expression patterns of IGF-1 (see Figure 3E and F) and total Akt (see Figure 3G and H) were unchanged in the duodenal mucosa of rats after WRS compared with those of normal rats.

Relative expression of *IGF-1*, *PTEN*, *Akt1*, *Akt2*, *Akt3*, *FoxO3a*, and *FoxO4* in the gastric mucosa of rats after WRS

Expression levels of selected genes were analyzed using RT-PCR. Amplification products were identified through melting curve profile analysis and confirmed with gel electrophoresis and sequencing. The data showed the relative transcript of each target gene normalized to *HPRT*. The real-time RT-PCR analyses of transcripts of 168 bp *IGF-1*, 151 bp *PTEN*, 269 bp *Akt1*, 263 bp *Akt2*, 498 bp *Akt3*, 331 bp *FoxO3a*, 148 bp *FoxO4*, and 556 bp *HPRT* are shown in Figure 4. All selected genes were transcriptionally active. After 3 and 7 h of WRS, messenger RNA (mRNA) transcript levels of *IGF-1* (see Figure 4A), *PTEN* (see Figure 4A), *FoxO3a* (see Figure 4C), and *FoxO4* (see Figure 4C) in rats subjected to WRS showed no significant difference compared with those in the non-WRS group ($P > 0.05$, $N = 6$). In addition, no significant difference was found in *PTEN* (see Figure 4A) mRNA transcript levels among the 4th, 8th, and 15th days after 7 h of WRS ($P > 0.05$, $N = 6$). However, *PTEN* (see Figure 4A) mRNA transcript levels at these stages were greater than those of the non-WRS group ($P < 0.05$, $N = 6$).

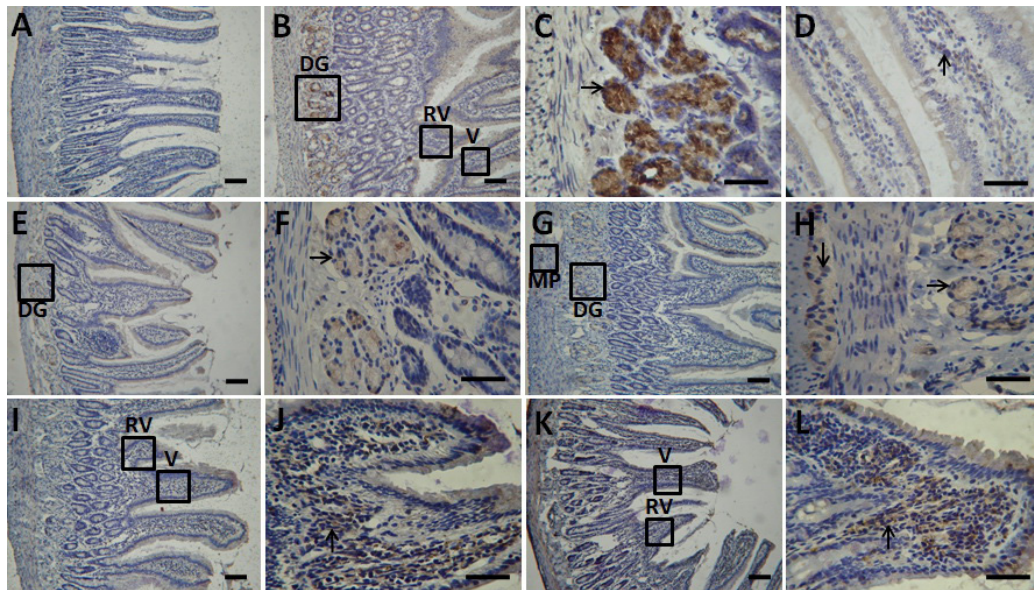


Figure 3. Immunohistochemical localization of IGF-1, PTEN, total Akt, FoxO3a, and FoxO4 in the duodenal mucosa of rats after water immersion and restraint stress. The immunohistochemical signals appear brown and the counterstained background appears blue in color. The figures indicate immunohistochemical localization of PTEN (B-D); immunohistochemical localization of IGF-1 (E and F); immunohistochemical localization of total Akt (G and H); immunohistochemical localization of FoxO3a (I and J), and immunohistochemical localization of FoxO4 (K and L). In control sections, normal albumin bovine was used instead of primary antibody (A). DG = duodenal gland; V = villi; RV = regenerated villi; MP = myenteric plexus; ↓ = myenteric plexus; → = duodenal gland; ↑ = lamina propria. Bar = 50 μ m.

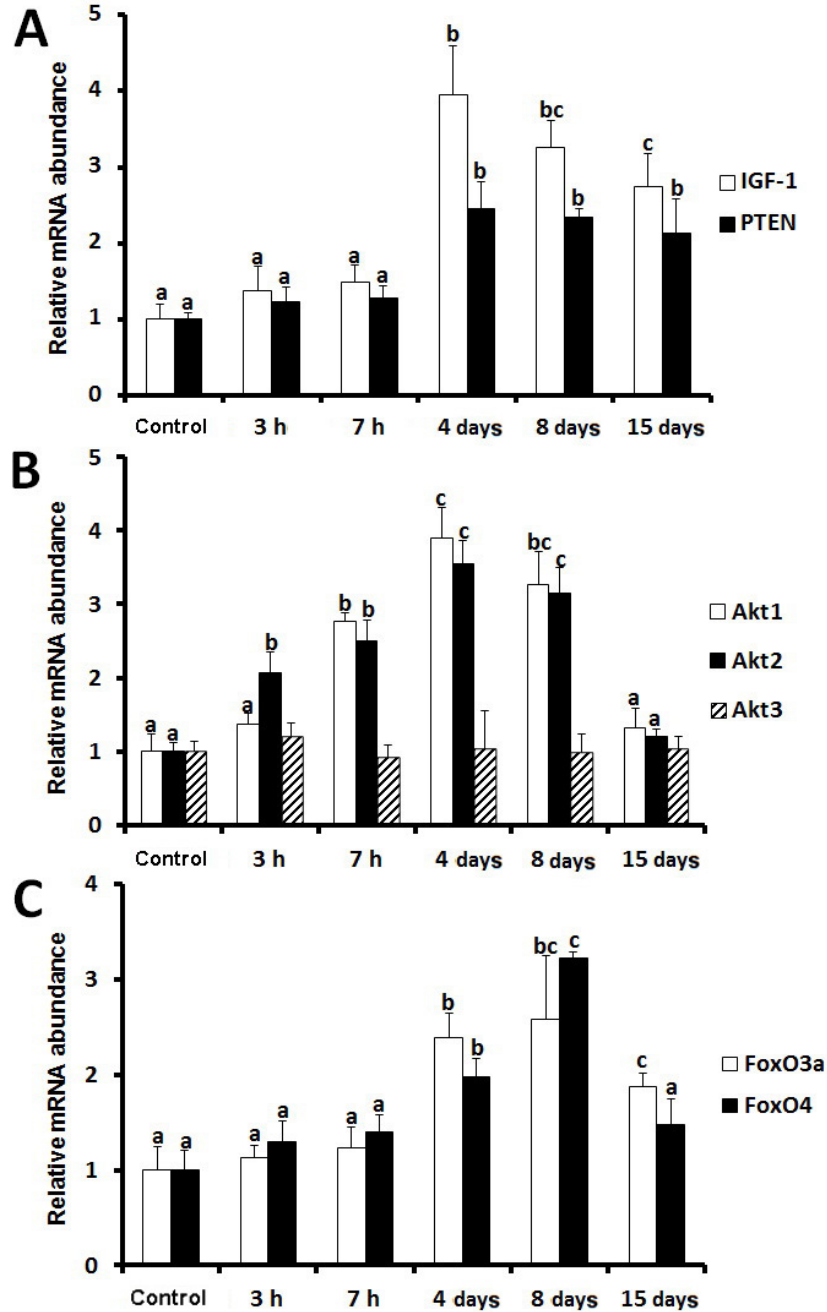


Figure 4. Relative expressions of *IGF-1*, *PTEN*, *Akt1*, *Akt2*, *Akt3*, *FoxO3a*, and *FoxO4* in the duodenal mucosa of rats after water immersion and restraint stress. The figures indicate *IGF-1* and *PTEN* (A); *Akt1*, *Akt2* and *Akt3* (B); *FoxO3a* and *FoxO4* (C). Different letters above the bars indicate a statistically significant difference in the same gene ($P < 0.05$, $N = 5$) between treatment groups.

In duodenal mucosa of rats after WRS, mRNA transcript levels of *IGF-1* (see Figure 4A), *Akt1* (see Figure 4B), and *Akt2* (see Figure 4B) were upregulated, with a peak 4 days after 7 h of WRS. The mRNA transcript levels of *Akt1* and *Akt2* returned to near baseline 15 days after 7 h of WRS. However, the mRNA transcript levels of *IGF-1* (see Figure 4A) in the WRS groups were still greater than that in the non-WRS group ($P < 0.05$, $N = 6$). Similarly, mRNA transcript levels of *FoxO3a* (see Figure 4C) and *FoxO4* (see Figure 4C) were also upregulated after WRS, peaking 8 days after 7 h of WRS. The *FoxO3a* mRNA transcript level (see Figure 4C) was still greater than that in the non-WRS group 15 days after 7 h of WRS ($P < 0.05$, $N = 6$). The *FoxO4* mRNA transcript level (see Figure 4C) returned to near baseline levels, and *Akt3* (see Figure 4B) mRNA transcript levels in rats subjected to WRS were unchanged compared with that of the non-WRS group ($P > 0.05$, $N = 6$).

DISCUSSION

The WRS rat has long been used as a model of gastroduodenal mucosal lesions (Wang and Johnson, 1992; Bogdaran et al., 2005; Adachi et al., 2011). As shown in previous studies, WRS results in microscopic damage evidenced by a nearly complete absence of villi. Significant macroscopic lesions are absent after stress. Mucosal repair is evident 12 h after stress and is almost complete at 24 h, although the restituted villi are short (Wang and Johnson, 1992). In addition, apoptosis plays an important role in the maintenance of normal gastrointestinal homeostasis and mucosal integrity (Sun et al., 1998). Recent studies have demonstrated that apoptosis is critically involved in gastric ulceration 24 h after ulcer induction (Konturek et al., 2001). Caspases are causative enzymes that induce apoptosis and are always present in intact cells, playing key roles in the pathogenesis of tissue injury by activating neutrophils. All known stimuli that induce apoptosis initiate events that culminate in caspase activation (Creagh et al., 2003). Previous studies have reported that IGF-1 and capsaicin administration to rats markedly reduces WRS-induced gastric mucosal injury by limiting the gastric accumulation of neutrophils through inhibition of caspase-3 activation (Zhao et al., 2009). In our present study, we found that TUNEL-positive cells are concentrated mainly in the lamina propria of the duodenal mucosa. In addition, WRS increased both caspase-3 activity and the number of TUNEL-positive cells after 3 and 7 h of WRS. Subsequently, caspase-3 activity and the number of TUNEL-positive cells gradually returned to near baseline levels, suggesting that apoptosis played a role in the duodenal mucosa of rats after WRS.

The IGF-1/PTEN/Akt/FoxO signaling pathway plays some critical roles in the regulation of survival, growth, differentiation, and migration in many cell types and tissues (Reddy et al., 2008). In particular, our previous study has demonstrated that the IGF-1/PTEN/Akt/FoxO signaling pathway is involved in protecting against ulcer through regulation of cellular apoptosis in the development and healing of rat gastric ulcers. To confirm whether the IGF-1/PTEN/Akt/FoxO signaling pathway is involved in the responses of the duodenal mucosa of rats subjected to WRS, we detected the localization and expression of IGF-1, PTEN, Akt, and FoxO.

Our results indicated that the localization of IGF-1 was unchanged in the duodenal mucosa of rats after WRS compared with that in non-WRS rats. IGF-1 was mainly observed in the cellular cytoplasm of duodenal glands. IGF-1 has also reportedly been discernible in the muscularis externa of the duodenum in the one-humped camel (Al Haj Ali et al., 2003). We suspected species-specific expression of IGF-1 in the duodenum. mRNA transcript levels of *IGF-1* were

upregulated, peaking at 4 days after 7 h of WRS. In the gastrointestinal tract, IGF-1 is secreted by salivary and other exocrine glands. Gastric ulceration reportedly triggers a several-fold increase in IGF-1 expression in ulcer margins (Coerper et al., 2001; Nguyen et al., 2007). Related findings have also indicated that IGF-1 is upregulated in injured skin, bone, and brain (Nguyen et al., 2007). In particular, IGF-1 reduces tissue injury by preventing cell death in animal models of renal ischemia/reperfusion. Among the various actions of IGF-1, its anti-apoptotic activity plays an important role in the reduction of ischemia/reperfusion-induced tissue injury by attenuating inflammatory responses (Harada et al., 2007). In addition, previous studies have demonstrated that IGF-1 has an essential role in gastrointestinal tract growth and development (Georgiev et al., 2003; Warzecha et al., 2006; Karcher et al., 2009; Mullen et al., 2011) and is an important growth factor in gut maintenance (Ryan and Costigan, 1993). Accordingly, our results suggested that IGF-1 was critically involved in the duodenal mucosa of rats after WRS.

As an important downstream effector of IGF-1, the PI3K/Akt signaling pathway is associated with pharmacotherapy through upregulation of cell proliferation in duodenal mucosa (Coeffier et al., 2011; Tuo et al., 2011). In addition, Nguyen et al. (2007) have found that upregulation of IGF-1 in gastric ulcer margins enhanced gastric ulcer healing by promoting cell re-epithelization and proliferation via the PI3K pathway (Nguyen et al., 2007). Furthermore, IGF-1 has been reported to reduce WRS-induced gastric mucosal injury by reducing gastric accumulation of neutrophils through inhibition of caspase-3 activation by PI3K/Akt signaling (Zhao et al., 2009). In the present study, total Akt was observed in the cellular cytoplasm of the duodenal glands and myenteric plexus. WRS triggered increases in *Akt1* and *Akt2* mRNA expression in rat duodenal mucosa, with a peak at 4 days after 7 h of WRS. mRNA transcript levels of *Akt3* were unchanged in the duodenal mucosa of rats after WRS compared with that of the non-WRS group. These results were consistent with our previous studies on WRS-induced gastric ulcers in rats (Huang et al., 2012), suggesting that Akt1 and Akt2 might participate in the regulation of cell proliferation in the duodenal mucosa of rats after WRS.

PTEN is an important tumor suppressor that controls Akt signaling. Its downstream targets are responsible for regulating many physiologically and pathologically significant processes, such as cellular proliferation, survival, growth, and motility (Leslie and Downes, 2004). Many researchers have concentrated on *PTEN* in gastric cancer (Liu et al., 2011; Wang et al., 2011; Xiong et al., 2011). The present study indicated that *PTEN* was found mainly in the cellular cytoplasm of the duodenal glands and lamina propria of villi. In the duodenal mucosa of rat subjected to WRS, *PTEN* was also detected in the cellular cytoplasm of the lamina propria of restituted villi. In addition, *PTEN* mRNA transcript levels of rats 4, 8, and 15 days after 7 h of WRS were greater than those of rats in the non-WRS group. Wang et al. (2011) have revealed that caspase-3 activity is related to the upregulation of *PTEN* in human gastric cancer MGC-803 cells. Our previous studies also demonstrated that *PTEN* is upregulated in WRS-induced gastric ulcers (Huang et al., 2012). Therefore, it was reasonable to suspect that *PTEN* may be involved in the regulation of cell apoptosis through the PI3K/Akt pathway in the duodenal mucosa of rats after WRS.

FoxO1, FoxO3a, and FoxO4 are all targets of the IGF-1/*PTEN*/Akt pathway and participate in several physiologic processes including cell proliferation, apoptosis, stress resistance, differentiation, and metabolism (Nakae et al., 2002). Our previous studies have found that FoxO3a and FoxO4 are the primary forkhead transcriptional factors localized to the gastrointestinal tract (Zhou et al., 2007; Huang et al., 2011). Liu et al. (2008) have shown that

FoxO1 mRNA is expressed at lower levels in duodenal subcutaneous adipose tissue than in other tissues in pigs. Therefore, in the present study, we detected the localization and expression of FoxO3a and FoxO4 in the duodenal mucosa of rats after WRS. The results indicated that FoxO3a and FoxO4 were primarily concentrated in the cellular cytoplasm of the lamina propria. PTEN was also detected in the cellular cytoplasm of the lamina propria of restituted villi in rats after WRS. At the same time, mRNA transcript levels of *FoxO3a* and *FoxO4* were upregulated, peaking 8 days after 7 h of WRS. This result was similar to that in rats with WRS-induced gastric ulcers (Huang et al., 2012). Phosphorylation of FoxO proteins by Akt results in cytoplasmic retention and inactivation, which in turn inhibits the expression of FoxO-regulated genes that control the cell cycle, cell death, and cell metabolism. The shuttling of FoxOs between the cytoplasm and the nucleus is a key step in apoptosis (Burgering and Kops, 2002). Therefore, FoxO3a and FoxO4 may also play some roles related to cell apoptosis regulation in the duodenal mucosa of rats after WRS. In conclusion, these observations raised the possibility that the IGF-1/PTEN/Akt/FoxO signaling pathway regulates cellular apoptosis in the duodenal mucosa of rats after WRS.

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