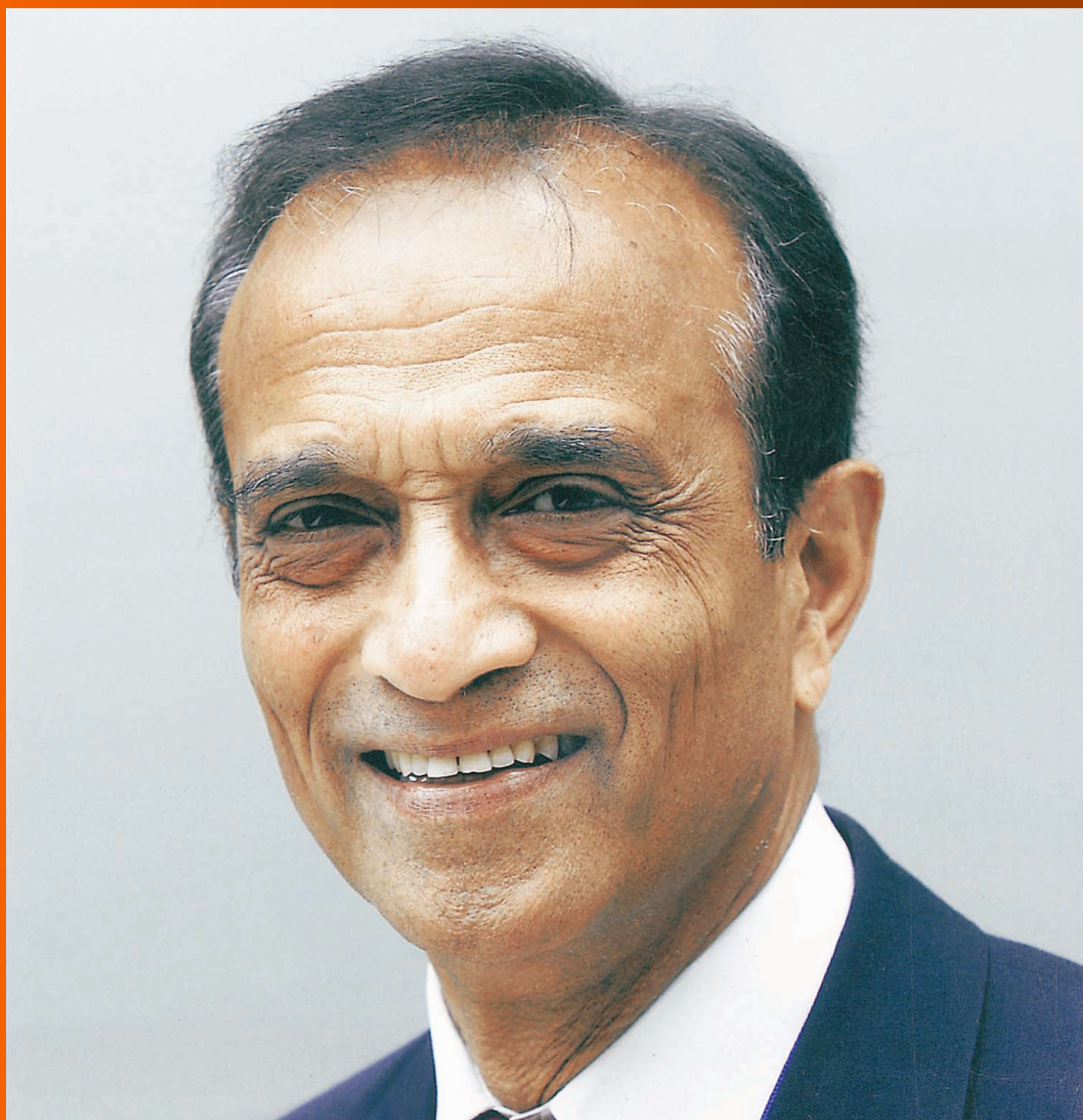


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Basic Study

High expression of type I inositol 1,4,5-trisphosphate receptor in the kidney of rats with hepatorenal syndrome

Jing-Bo Wang, Ye Gu, Ming-Xiang Zhang, Shun Yang, Yan Wang, Wei Wang, Xi-Ran Li, Yi-Tong Zhao, Hai-Tao Wang

Jing-Bo Wang, Ye Gu, Ming-Xiang Zhang, Yan Wang, Wei Wang, Xi-Ran Li, Yi-Tong Zhao, Liver Cirrhosis Ward, the Sixth People's Hospital of Shenyang, Shenyang 110006, Liaoning Province, China

Shun Yang, Liaoning Cancer Hospital & Institute, Shenyang 110042, Liaoning Province, China

Hai-Tao Wang, Department of General Surgery, the Second Affiliated Hospital of Shenyang Medical College, Shenyang 110002, Liaoning Province, China

ORCID number: Jing-Bo Wang (0000-0001-9207-6245); Ye Gu (0000-0002-3798-1119); Ming-Xiang Zhang (0000-0001-6519-3497); Shun Yang (0000-0003-4743-0588); Yan Wang (0000-0002-0427-3458); Wei Wang (0000-0002-4964-3280); Xi-Ran Li (0000-0002-3253-407X); Yi-Tong Zhao (0000-0002-5059-8608); Hai-Tao Wang (0000-0002-7385-0642).

Author contributions: Wang JB, Wang HT and Gu Y contributed equally to this work, and all of them were involved in the design and performing of the experiment, data analysis and drafting of the article; Zhang MX, Wang Y and Yang S participated in the study, hepatic and renal pathological examination and biochemical test; Li XR, Wang W and Zhao YT completed data analysis, Western blot analysis and real-time PCR and drafting of the paper.

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Correspondence to: Hai-Tao Wang, MSc, Surgeon, Department of General Surgery, the Second Affiliated Hospital of Shenyang Medical College, No. 20, Beijiuma Road, Heping District, Shenyang 110002, Liaoning Province, China. whszyypwk@163.com
Telephone: +86-18002452018
Fax: +86-24-31251510

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Abstract

AIM

To detect the expression of type I inositol 1,4,5-trisphosphate receptor (IP₃RI) in the kidney of rats with hepatorenal syndrome (HRS).

METHODS

One hundred and twenty-five Sprague-Dawley rats were randomly divided into four groups to receive an intravenous injection of D-galactosamine (D-GalN) plus lipopolysaccharide (LPS; group G/L, *n* = 50), D-GalN alone (group G, *n* = 25), LPS alone (group L, *n* = 25), and normal saline (group NS, *n* = 25), respectively.

At 3, 6, 9, 12, and 24 h after injection, blood, liver, and kidney samples were collected. Hematoxylin-eosin staining of liver tissue was performed to assess hepatocyte necrosis. Electron microscopy was used to observe ultrastructural changes in the kidney. Western blot analysis and real-time PCR were performed to detect the expression of IP₃RI protein and mRNA in the kidney, respectively.

RESULTS

Hepatocyte necrosis was aggravated gradually, which was most significant at 12 h after treatment with D-galactosamine/lipopolysaccharide, and was characterized by massive hepatocyte necrosis. At the same time, serum levels of biochemical indicators including liver and kidney function indexes were all significantly changed. The structure of the renal glomerulus and tubules was normal at all time points. Western blot analysis indicated that IP₃RI protein expression began to rise at 3 h ($P < 0.05$) and peaked at 12 h ($P < 0.01$). Real-time PCR demonstrated that IP₃RI mRNA expression began to rise at 3 h ($P < 0.05$) and peaked at 9 h ($P < 0.01$).

CONCLUSION

IP₃RI protein expression is increased in the kidney of HRS rats, and may be regulated at the transcriptional level.

Key words: Hepatorenal syndrome; Type I inositol 1,4,5-trisphosphate receptor; Glomerular mesangial cells; Vascular smooth muscle cells

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Core tip: Type I inositol 1,4,5-trisphosphate receptor (IP₃RI) protein expression is increased in the kidney of hepatorenal syndrome (HRS) rats, and IP₃RI protein expression may be regulated at the transcriptional level. Increased expression of IP₃RI may be closely associated with HRS development and progression through excessive renal vascular contraction resulting in insufficient renal blood perfusion.

Wang JB, Gu Y, Zhang MX, Yang S, Wang Y, Wang W, Li XR, Zhao YT, Wang HT. High expression of type I inositol 1,4,5-trisphosphate receptor in the kidney of rats with hepatorenal syndrome. *World J Gastroenterol* 2018; 24(29): 3273-3280 Available from: URL: <http://www.wjgnet.com/1007-9327/full/v24/i29/3273.htm> DOI: <http://dx.doi.org/10.3748/wjg.v24.i29.3273>

INTRODUCTION

Hepatorenal syndrome (HRS), one of the most severe complications of liver failure (LF) and the leading cause of death in LF^[1], is functional renal failure secondary to

LF^[2-10]. At present, the exact pathogenesis of HRS is still unclear, and the reduction in renal blood flow induced by renal vasoconstriction is considered to play a central role in the development of HRS^[11,12]. Renal blood flow is regulated by the contraction and relaxation of vascular smooth muscle cells (VSMCs) of glomerular afferent arteries and glomerular mesangial cells (GMCs), while the contraction and relaxation of VSMCs and GMCs are regulated by intracellular Ca²⁺ concentrations^[13,14]. GMCs are in direct contact with glomerular endothelial cells. When GMCs contract, glomerular mesangial volume decreases by 20%-25%, glomerular capillary plexuses are reduced, and the area for glomerular filtration is reduced. Inositol 1,4,5-trisphosphate (IP₃) receptor (IP₃Rs) is the main Ca²⁺ release channel in cells. IP₃ is an intercellular second messenger mediating transmembrane signal transmission. When binding to IP₃Rs, IP₃ mediates intracellular calcium release and extracellular calcium influx^[15,16]. VSMCs and GMCs transmit extracellular signals into the cell via the IP₃-IP₃R pathway, increasing intracellular Ca²⁺ concentrations^[17,18]. Is high expression of renal IP₃Rs associated with HRS? To answer this question, in the present study we detected the expression of IP₃RI protein and mRNA in the kidney of HRS rats to determine the relationship between IP₃RI expression and HRS.

MATERIALS AND METHODS

Materials

Specific pathogen-free (SPF) Sprague-Dawley (SD) rats, weighing 220 g ± 20 g, were purchased from the Laboratory Animal Center of the Academy of Military Medical Sciences (Animal Certificate No. SCXK-2017-004; Beijing, China). Prior to experimentation, the rats were reared in separate cages at 23 °C ± 3 °C under a 12 h/12 h light/dark cycle, with free access to ordinary chow (purchased from the Laboratory Animal Center of China Medical University, Shenyang, China) and water. After one week of adaptation, the rats were used in the experiments.

D-galactosamine (D-GalN) and lipopolysaccharide (LPS) were purchased from Sigma (St. Louis, MO, United States). Anti-IP₃RI antibody was obtained from US Biological (St. Salem, OR, United States). An enhanced chemiluminescence (ECL) kit was purchased from Pierce, Dallas, TX, United States. RNAiso™ plus, Prime Script™ RT Reagent Kit, and SYBR® Premix EX Tag™ were purchased from TakaRa (Shiga, Japan).

Rat model of HRS

One hundred and twenty-five SD rats of SPF grade, weighing 220 ± 20 g, were randomly divided into four groups to receive an intravenous injection of D-GalN plus LPS (group G/L), D-GalN alone (group G), LPS alone (group L), and normal saline (group NS), respectively. Each group was further divided into five subgroups for testing at different time points (3, 6, 9,

12, and 24 h). Group G/L contained ten rats at each time point, and the other groups contained five rats at each time point. The rats were weighed and then injected with D-GalN (400 mg/kg body weight) and/or LPS (32 µg/kg) or NS (2 mL/kg) *via* the tail vein. Rats that died during the modeling process were excluded from the study. At 3, 6, 9, 12, and 24 h after modeling, the rats in groups G/L, G, and L were anesthetized with 0.8% pentobarbital sodium at 40 mg/kg *via* intraperitoneal injection and sacrificed to obtain liver and kidney tissues. A section of each tissue was fixed in formalin, and the remainder was preserved at -80 °C for Western blot and real-time PCR analysis of IP₃RI protein and mRNA expression, respectively.

Western blot analysis

For total protein preparation, renal tissue was lysed for 15 min in a lysis solution containing 50 mmol/L Tris-HCl (pH 7.4), 150 mmol/L NaCl, 1% Triton X-100, 1% sodium deoxycholate, 0.1% SDS, 1 mmol/L EDTA, 5 mg/mL leupeptin, sodium orthovanadate, sodium fluoride, and 1 mmol/L PMSF, and then centrifuged at 12000 rpm for 12 min. The supernatant was collected and preserved at -80 °C.

After total protein concentration was determined using the bicinchoninic acid (BCA) method, the protein samples were mixed with 5 × loading buffer at a ratio of 4:1 (v/v), boiled for 5 min, resolved by 8% SDS-polyacrylamide gel electrophoresis, and transferred to polyvinylidene difluoride membranes. The membranes were then blocked with 5% skimmed milk, Tris-buffered saline and Tween-20, and incubated with primary antibody against IP₃RI (dilution, 1:1000) at 4 °C overnight. This was followed by incubation with horseradish peroxidase (HRP)-conjugated goat anti-rabbit secondary antibody (dilution, 1:3000) for 2 h at room temperature. The immunoblots were visualized using an enhanced chemiluminescence system. The molecular weight of the target band was 230 kDa. β-actin (45 kDa) was used as an internal control. Digital imaging software was used for densitometry analyses, and the relative IP₃RI level was calculated as IP₃RI grey value divided by β-actin grey value.

Real-time PCR

Total RNA was prepared from renal tissue using Trizol according to the manufacturer's instructions. After reverse transcription to cDNA in a 10-µL system containing 2.0 µL of 5 × Prime Script™ Buffer (for real time), 0.5 µL of Prime Script™ RT Enzyme Mi, 0.5 µL of Oligo dT Primer (50 µmol/L), 0.5 µL of Random 6-mers (100 µmol/L), 5.5 µL of RNase Free dH₂O, and 1.0 µL of RNA (500 ng/µL), real-time PCR was performed in a 25-µL system containing 12.5 µL of 2 × SYBR Premix Ex Tag™, 0.5 µL of PCR Forward Primer (10 µmol/L), 0.5 µL of PCR Reverse Primer (10 µmol/L), 9.5 µL of RNase Free dH₂O, and 2.0 µL of cDNA. Cycling parameters were 95 °C for 30 s and 45 cycles of 95 °C for 5 s, 57 °C

for 20 s, and 72 °C for 30 s. Glyceraldehyde 3-phosphate dehydrogenase (GAPDH) was used as an endogenous reference. The primers used were as follows: forward, 5'-TCTGGCCAGCTGTCAGAACTAAAG-3' and reverse, 5'-GTGGGTTGACATTCATGTGAGGA-3' for IP₃RI, and forward, 5'-GACAACTTTGGCATCGTGGA-3' and reverse, 5'-GACAACTTTGGCATCGTGGA-3' for GAPDH. The double-standard curve method was used to determine the relative IP₃RI mRNA expression.

Statistical analysis

All statistical analyses were performed using SPSS 13.0 software. Numerical data, expressed as mean ± standard error of the mean, were compared using analysis of variance. *P* values < 0.05 were considered statistically significant.

RESULTS

Successful induction of HRS in rats with D-GalN/LPS

Following intravenous injection of D-GalN at 400 mg/kg body weight combined with LPS at 32 µg/kg in male SD rats, HRS was successfully induced. Twelve hours after injection, glomerular filtration rate (GFR) significantly decreased, liver and kidney function were severely impaired, and serum biochemical indices, such as alanine aminotransferase (ALT), blood urea nitrogen (BUN), and creatinine (Cr), exhibited significant changes. Hematoxylin-eosin staining showed massive hepatocyte necrosis with severe hemorrhage (Figure 1), while renal tissue had a normal morphology at the various time points (Figure 2). These changes were consistent with the clinical features of HRS.

Western blot analysis of IP₃RI protein expression

IP₃RI (230 kDa) and β-actin (45 kDa) were detected in all groups. Densitometry analyses showed that IP₃RI protein expression was significantly elevated in group G/L compared with group NS. This elevation began at 3 h (1.46 ± 0.07 vs 1.00 ± 0.05, *P* = 0.011), became obvious at 9 h, and reached a peak at 12 h (2.89 ± 0.14 vs 1.00 ± 0.05, *P* = 0.000) (Figure 3A).

At 12 h, the expression of IP₃RI protein in the kidney was significantly higher in group G/L than in groups G (1.17 ± 0.08) and L (1.02 ± 0.09) (*P* = 0.000 for both), thus excluding the impact of D-GalN or LPS on the expression of IP₃RI protein. There was no significant difference in IP₃RI protein expression between groups G and L (*P* = 0.245) or between group G or L and group NS (*P* > 0.05 for both) (Figure 3B).

RT-PCR analysis of IP₃RI mRNA expression

IP₃RI mRNA expression was significantly elevated in group G/L compared with group NS. This elevation began at 3 h (2.89 ± 0.51 vs 1.00 ± 0.00, *P* = 0.05), became obvious at 6 h (5.01 ± 0.38, *P* = 0.000), and reached a peak at 9 h (9.96 ± 0.63, *P* = 0.000). IP₃RI mRNA expression began to decline at 12 h, and at 24

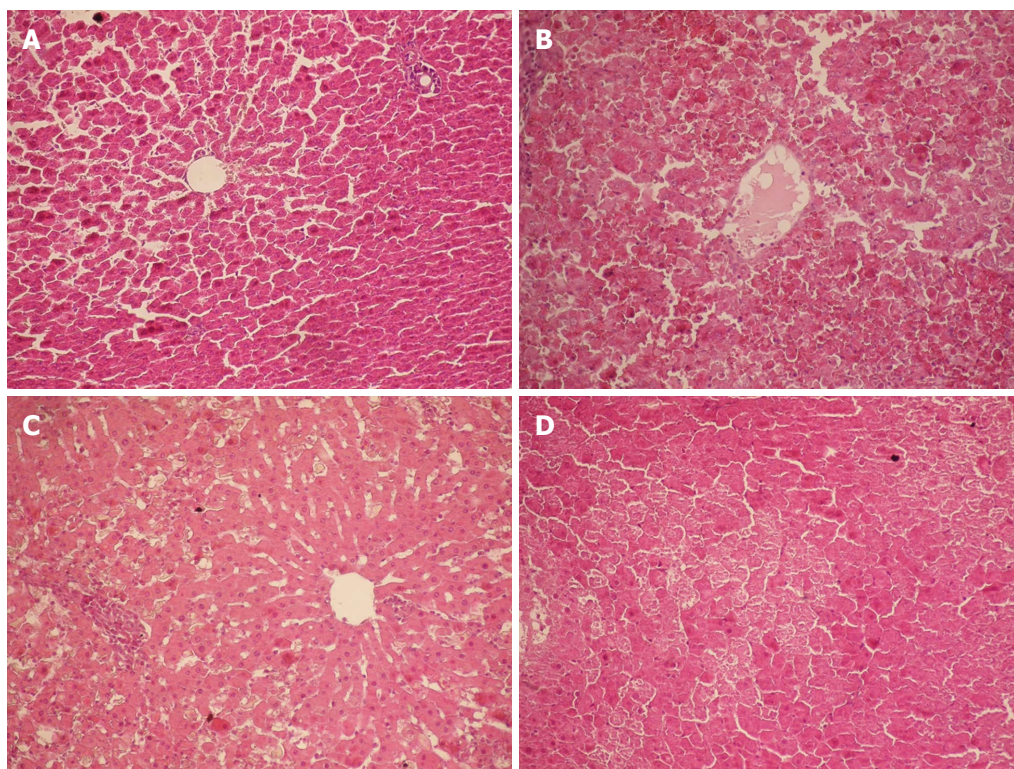


Figure 1 Histopathology of the liver (HE staining, × 200). A: Group Normal Saline (NS). Normal hepatocytes were arranged in cords; B: Group D-galactosamine (D-GalN) plus lipopolysaccharide (LPS) (G/L). At 12 h, massive hepatocyte necrosis with severe hemorrhage developed; C: Group D-GalN (G). At 12 h, spotty hepatocyte necrosis was observed; D: Group LPS (L). At 12 h, hepatocytes began to develop necrosis, with incomplete necrosis visible.

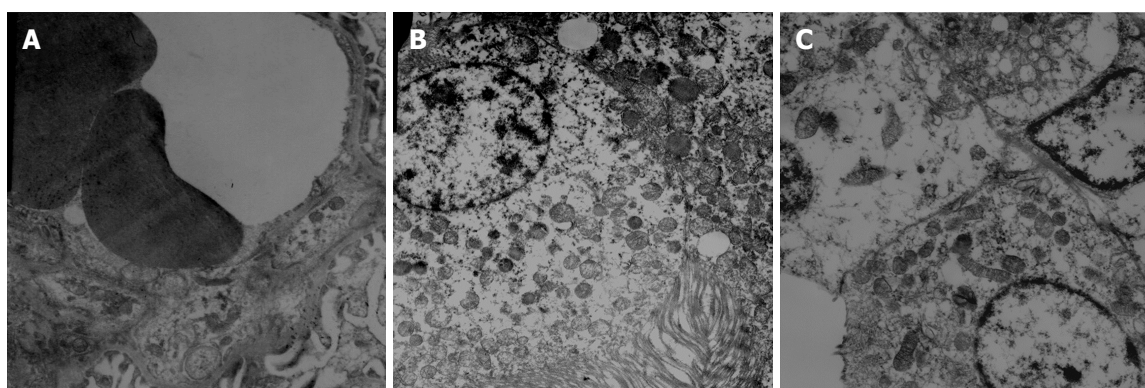


Figure 2 Histopathology of the kidney. A: The glomerular basement membrane of the kidney was intact, and the foot processes of podocytes and fenestra of endothelial cells were clearly visible; B: The basal part of proximal tubule cubical epithelial cells had abundant plasma membrane infolding, which was rich in longitudinally arranged mitochondria with intact cristae. On the free surface of proximal tubule cubical epithelial cells, microvilli were long and dense; C: The basal part of distal tubule cubical epithelial cells also had abundant plasma membrane infolding, which was rich in mitochondria. On the free surface of distal tubule cubical epithelial cells, microvilli were short and sparse.

h, it returned to the level observed at 6 h. IP₃RI mRNA expression at 9 h was significantly higher in group G/L than in groups G (1.43 ± 0.18) and L (1.29 ± 0.17) ($P = 0.000$ for both; Figure 4). IP₃RI mRNA expression did not differ significantly between group G or L and group NS ($P > 0.05$ for both).

DISCUSSION

HRS is one of the most common and severe complications of fulminant liver failure (FHF) and an advanced

liver disease, with approximately 55% of FHF patients developing HRS^[19,20]. The pathogenesis of HRS is still not completely clear, although it is believed to be associated with excessive renal vascular contraction, insufficient renal blood perfusion, sympathetic nervous system activation, and increased synthesis of vasoactive substances, all of which make the kidneys more sensitive to low perfusion^[21,22]. Renal blood flow and GFR decrease significantly in HRS due to renal vasoconstriction, and many factors are involved in this process. A significant increase in vasoconstricting factors

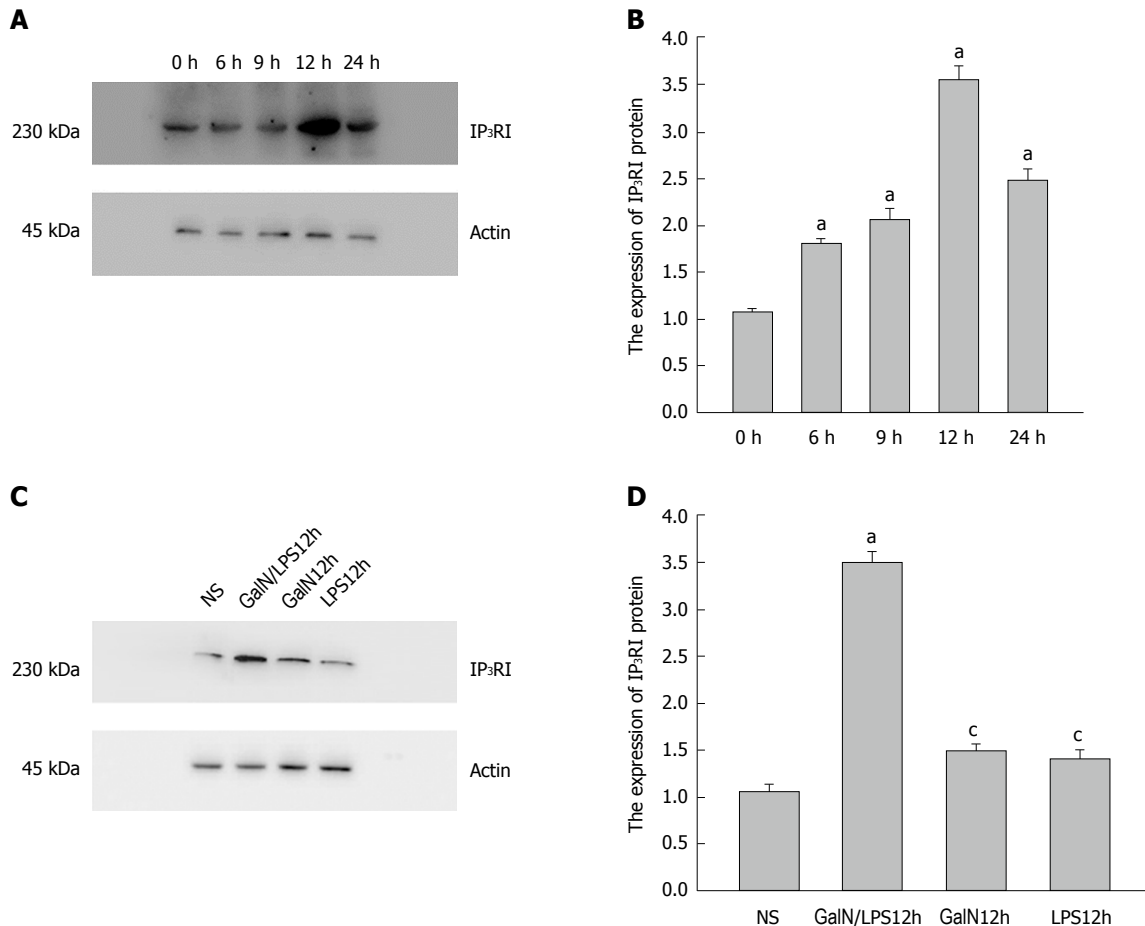


Figure 3 Expression of type I inositol 1,4,5-trisphosphate receptor (IP₃RI) protein in the kidney of rats in each group. A: The expression of IP₃RI protein in the kidney significantly increased in group D-galactosamine (D-GalN) plus lipopolysaccharide (LPS) (G/L), and was especially prominent at 12 h [^a*P* < 0.05 vs group Normal Saline (NS)]. B: The expression of IP₃RI protein in the kidney significantly increased in group G/L compared with the other groups (^a*P* < 0.05 vs group NS, ^c*P* < 0.05 vs group G/L).

[e.g., endothelin (ET) and angiotensin II] in the blood not only leads to renal vascular contraction, but also decreases the glomerular filtration coefficient (*K_f*) and GFR^[23,24]. The contraction of VSMCs results in reduced renal blood flow, while GMC contraction reduces the glomerular filtration fraction and coefficient. As both VSMCs and GMCs are extremely sensitive to vasoactive substances, GFR is significantly decreased in HRS. ET and angiotensin II are important renal vasoconstricting factors, and they activate Ca²⁺ channels *via* the IP₃-IP₃Rs pathway. IP₃Rs is the intracellular calcium reservoir, which is present mainly in the endoplasmic reticulum and on the membrane, directly or indirectly mediating the calcium influx^[25-27]. In addition, IP₃Rs is also present in the nucleus, participating in nuclear calcium release and regulating gene expression^[28,29]. When IP₃ binds to IP₃Rs, a conformational change in IP₃Rs occurs, the calcium channel is open, and the calcium reserve in the endoplasmic reticulum is released into the cytoplasm. As a result, cytoplasmic free Ca²⁺ concentration ([Ca²⁺]_i) increases, thus causing cell contraction^[30-32]. Therefore, IP₃Rs mediates an important Ca²⁺ signaling pathway in the cell, and the expression of IP₃Rs is closely related

to the sensitivity of the kidney to vasoconstrictors^[33-35]. IP₃Rs has four types of ligand binding sites associated with calcium channels^[36-39], and renal IP₃RI is mainly found in GMCs and VSMCs, and there is almost no IP₃RI on the surface of other renal cells^[40,41]. Therefore, the expression levels of IP₃RI in renal GMCs and VSMCs may be related to renal vasoconstriction. As the opening of IP₃-IP₃Rs channels can increase intracellular [Ca²⁺]_i, theoretically the expression level of IP₃RI is closely related to the intracellular [Ca²⁺]_i level. Wang *et al* observed increased expression of IP₃RI in the glomerular capillary loops and anterior artery of rats with liver cirrhosis by immunohistochemistry. However, it is unknown whether the expression of IP₃RI increases in FHF. To answer this question, we detected IP₃RI expression in the renal tissue of a rat model of FHF at different time points at both the protein and mRNA levels using Western blot and real-time quantitative PCR, respectively.

Semi-quantitative Western blot analysis demonstrated that IP₃RI protein expression was low in normal kidney tissue. Following treatment with D-GalN plus LPS, IP₃RI protein expression began to rise at 3 h and

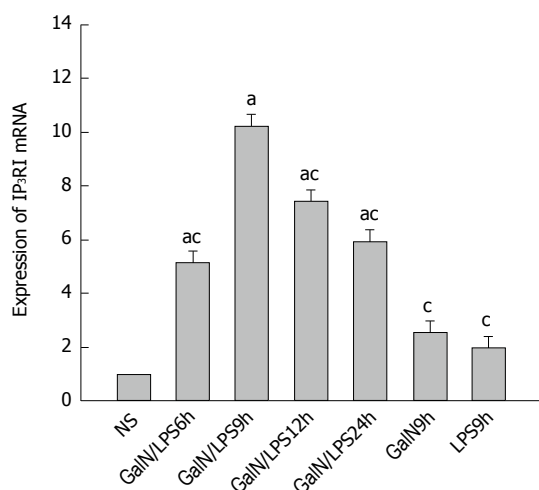


Figure 4 Expression of type I inositol 1,4,5-trisphosphate receptor (IP₃RI) mRNA in the kidney of rats in group D-galactosamine (D-GalN) plus lipopolysaccharide (LPS) (G/L). The expression of IP₃RI mRNA in the kidney significantly increased in group G/L, and was especially prominent at 12 h (^a*P* < 0.05 vs group Normal Saline (NS), ^c*P* < 0.05 vs group G/L).

reached a peak at 12 h. Interestingly, liver and kidney dysfunction and hepatocyte necrosis were most severe and blood TNF- α and ET-1 levels were highest at 12 h, which were concomitant with the elevation of IP₃RI protein expression in the kidney^[42,43]. By treating the animals with D-GalN or LPS alone, we excluded the effect of these drugs on IP₃RI protein expression. As HRS developed 12 h after D-GalN/LPS administration, we compared the IP₃RI protein expression at this time point among the groups. The results showed that IP₃RI protein expression was high in group G/L and low in groups G, L, and NS.

In order to understand whether IP₃RI protein expression in the kidney is regulated at the transcriptional level, real-time quantitative PCR was performed. The fluorescent dye SYBR Green I^[44,45] added to the PCR reaction system can be incorporated into double-stranded DNA with PCR amplification and markedly enhance fluorescence^[46-48]. The relative expression levels of these two parameters were calculated by the housekeeping gene GAPDH. It was found that the relative expression of IP₃RI mRNA to GAPDH mRNA began to rise 3 h after D-GalN and LPS administration, but the protein level did not rise at this time point. At 9 h, the expression level of IP₃RI mRNA reached the highest level. Although the protein level was also high at this time point, it was lower than that at 12 h. The expression of IP₃RI mRNA began to decrease, but it was still significantly higher than that in the control group. These changes can be explained from two aspects. On the one hand, IP₃RI mRNA expression may be prior to protein expression, which is associated not only with the translation efficiency and the speed of mRNA degradation, but also with the rate of protein degradation. On the other hand, the protein synthesis process also includes the assembly and translocation of proteins in ribosomes, which may affect the final expression of IP₃RI protein.

In conclusion, joint D-GalN/LPS administration can induce HRS in SD rats at 12 h, which is concomitant with peak IP₃RI protein expression in the kidney. Increased IP₃RI protein expression may be regulated at the transcriptional level. Thus, increased expression of IP₃RI may be closely associated with HRS development and progression.

ARTICLE HIGHLIGHTS

Research background

Hepatorenal syndrome (HRS) is one of the common and severe complications of liver failure and advanced liver disease, with approximately 55% of these patients developing this severe complication. At present, HRS has unclear pathogenesis, limited treatment options, and poor therapeutic efficacy. Once renal dysfunction aggravates rapidly, 60%-80% of patients with HRS will die. Therefore, elucidating the mechanism underlying the development and progression of HRS and taking effective preventive and therapeutic measures may improve the success rate of rescue, the incidence rate, and the mortality rate of HRS.

Research motivation

To detect the protein and mRNA expression of type I inositol 1,4,5-trisphosphate receptor (IP₃RI) in the kidney of rats with HRS by Western blot and real-time PCR.

Research objectives

To explore whether high expression of renal IP₃RI is associated with Ca²⁺ influx in vascular smooth muscle cells of glomerular afferent arteries and glomerular mesangial cells in rats with HRS.

Research methods

D-galactosamine (D-GalN) and/or lipopolysaccharide (LPS) were used to treat male Sprague-Dawley (SD) rats via the tail vein. Twelve hours after injection, massive hepatocyte necrosis with severe hemorrhage occurred in the liver, while renal tissue had a normal morphology. In addition, liver and kidney function was impaired severely, and serum biochemical indexes exhibited significant changes. These changes were consistent with the clinical features of HRS. Western blot and real-time PCR were then used to detect the protein and mRNA expression of renal IP₃RI, respectively.

Research results

IP₃RI protein expression was significantly elevated in rats with HRS. The elevation began at 3 h and reached the peak at 12 h. IP₃RI mRNA expression was also significantly elevated in rats with HRS. The elevation began at 3 h and peaked at 9 h.

Research conclusions

Joint D-GalN/LPS administration can induce HRS in SD rats at 12 h, which is concomitant with peaked IP₃RI protein and mRNA expression in the kidney. Increased expression of IP₃RI may be closely associated with HRS development and progression.

Research perspectives

Our results suggest that IP₃RI may be a signal molecule involved in the reduction of renal blood flow induced by renal vasoconstriction in HRS, thus providing a theoretical basis for further research of the pathogenesis of HRS. Gene silencing technology may be adopted to further elucidate the role of IP₃RI in the pathogenesis of HRS.

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