



Wip1^{-/-} ameliorates hepatic ischemia/reperfusion injury via PI3K/Akt activation

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Background: Liver is the largest organ in the body. It constitutes 2.5% of the body weight and receives approximately 25% of the cardiac output via both the dual blood flow and the portal vein as well as the hepatic artery, the latter of which contributes to 75–80% of the total flow and >50% of the total oxygen supply. Warm ischemia/reperfusion (I/R) injury is a common acute liver injury in clinical scenario. The Pringle manoeuvre in many liver surgeries is one of the many causes. By now researchers have found out that I/R can be mediated by many mechanisms including the pressure change in liver sinusoids mediated by sinusoidal endothelial cells (SEC) and nitric oxide (NO), innate immunity regulation by Kupffer cell, ATP-depletion-dependent liver cell necrosis and caspase-dependent apoptosis. Akt is a serine/threonine kinase which plays a critical role in regulating various biological processes including apoptosis, autophagy, cell growth, regeneration and protein synthesis. It has been revealed that Akt activates downstream proteins like Bad to regulate pathogenesis of liver I/R injury. Wip1 (wild-type p53 induced phosphatase 1), another serine/threonine phosphatase, plays a key role in immunity and inflammation. However, it still remains mystery whether wip1 is involved in pathogenesis and progression of liver I/R injury. In this study, we were aimed to discover the functional role of *wip1* gene in acute liver injury and the possible underlying mechanisms.

Methods: We used partial (2/3) liver warm I/R injury model and established wip1 knock-out mice to investigate the expression of wip1 and its impact on pathological changes after I/R injury. Moreover, we used terminal deoxynucleotidyl transferase dUTP nick end labeling (TUNEL) assay and western blot to study if PI3K/Akt and apoptotic pathway was affected by wip1 expression following liver injury.

Results: We have found that wip1 expression level was significantly reduced after I/R injury in wild-type mice, whereas in wip1 knock-out mice, the serum levels of alanine aminotransferase (ALT) were significantly decreased (851.3±270.9 U/L) compared with wild-type mice (1172.5±237.1 U/L) (P<0.01), indicating that wip1 down-regulation in return may protect liver tissues against I/R injury. Moreover, the Suzuki's scores of wip1^{-/-} mice (5.25±0.43) was dramatically decreased compared with WT mice (7.75±0.43), indicating less severe morphological damage of the liver following I/R injury by wip1 knock-out. The Terminal deoxynucleotidyl transferase dUTP nick end labeling (TUNEL) assay has confirmed that the rate of apoptosis was significantly reduced in wip1^{-/-} mice (30.4%±3.7%) compared to WT mice (62.3%±5.6%). Finally, Western Blot has showed that the expression levels of p-Akt, p-p70 S6K and p-S6 were markedly up-regulated in wip1^{-/-} mice following I/R injury, indicating their possible role in wip1^{-/-} mediated liver protection.

Conclusions: In conclusion, this study has demonstrated that wip1 expression was down-regulated following I/R injury and wip1 knock-out in return may protect liver tissues from programmed cell death partially through activating PI3K/Akt pathway. Thus, targeting wip1 expression and the downstream PI3K/Akt pathway may be beneficial for patients undergoing I/R injury and liver damage.

Keywords: Wip1; gene knockout mice; warm ischemia/reperfusion injury; Akt, apoptosis

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Introduction

As the largest organ of the body liver plays important roles in various biological and physiological functions including biosynthesis, secretion and transformation, etc. Ischemia/reperfusion (I/R) injury is a causal factor which contributes to both morbidity and mortality in several clinical settings, including hepatic sinusoidal obstruction syndrome, hemorrhagic shock, trauma and cardiac arrest (1). Moreover, during hepatectomy and vascular reconstruction, a major obstacle to transplantation surgery and liver resection where reperfusion after sustained ischemia is unavoidable is the vulnerability of the liver to I/R injury. Hepatic I/R injury occur in diverse pathological and clinical circumstances, including liver preservation for transplantation, liver surgery, hemorrhagic shock-resuscitation, veno-occlusive disease, and heart failure as well. The liver can present three different forms of ischemia, the cold ischemia (or hypothermic), the warm ischemia (or normothermic), and the rewarming ischemia. Cold ischemia occurs while the organ awaits implantation where it is intentionally applied to reduce metabolic activities of the graft, and thus it happens almost exclusively in transplant settings. Warm ischemia occurs when hepatic inflow occlusion (Pringle maneuver) or total vascular exclusion are induced to minimize blood loss when dividing the liver parenchyma. Therefore, unlike cold ischemia, warm ischemia occurs in diverse situations, including hepatic transplantation and trauma, hemorrhagic shock, and liver surgery. However, rewarming ischemia typically occurs during situations of graft manipulation or graft implantation or while performing the vascular reconstruction when the cold liver is subjected to room. Although ischemic stress by itself primes cells for damage and will ultimately induce cell death, ischemic injury usually does not manifest itself until after the ischemic liver is reperfused (2).

In the 1980s, it has already been shown that cold ischemia caused injury specifically to sinusoidal endothelial cells (SEC). Moreover, endothelial wall disruption induces leukocyte and platelet adhesion, the latter of which leads to SEC apoptosis after cold ischemic liver reperfusion (3-6). Unlike cold

ischemia, warm ischemia is poorly tolerated and leads to rapid death of hepatocytes (7). This severe injury of liver cells is most likely preceded by massive destruction of endothelial cells (8). There is a growing body of studies focused on understanding the underlying mechanisms of warm I/R injury. The most well studied mechanism is TNF- α mediated apoptosis of the Kupffer cells. It has been found that I/R activates the resident macrophages of the liver—the Kupffer cells, and Kupffer cell activation after reperfusion generate reactive oxygen species (ROS) and leads to release of both lysosomal enzymes and various pro-inflammatory cytokines like TNF- α (9,10), whereas functional inactivation of Kupffer cells, on the contrary, attenuates liver injury during both early and late reperfusion (9,11,12). Moreover, during early stages of reperfusion, the release of oxygen free radicals as well as further binding of these cytokines to respective receptors lead to the initiation of the complex apoptotic machinery and eventually programmed cell death of hepatocytes (10). It has been proved that several pro-apoptotic proteins including caspase-8 and caspase-3 are activated during the reperfusion phase, and cytochrome c is found to be released from the mitochondria into the cytoplasm. This cascade activates both extrinsic and intrinsic apoptotic pathway, finally resulting in cell death (13) (*Figure 1*). In addition, stress-activated protein kinases, especially c-Jun kinase, are found activated during hypoxia reoxygenation by extracellular stimuli and results in activation of pro-apoptotic transcription factors and initiation of apoptosis (14,15).

The loss of mitochondrial ATP production is the primary ischemic stress to liver and most other solid tissues. Another mechanism of I/R injury is membrane Na/K ATPase inactivation due to ATP depletion, which leads to cell rounding and swelling, as well as mitochondria swelling and ER dilatation, and bleb formation (16,17). Increasing evidence indicate apoptosis activation as a key mechanism of hepatic I/R injury (6,8). However, the observations of post-ischemic cell swelling, vacuolation, karyolysis, and cell content release, are characteristic features of necrosis (18). Therefore, whether cells die predominantly by apoptosis or necrosis needs to be determined in experimental or clinical settings.

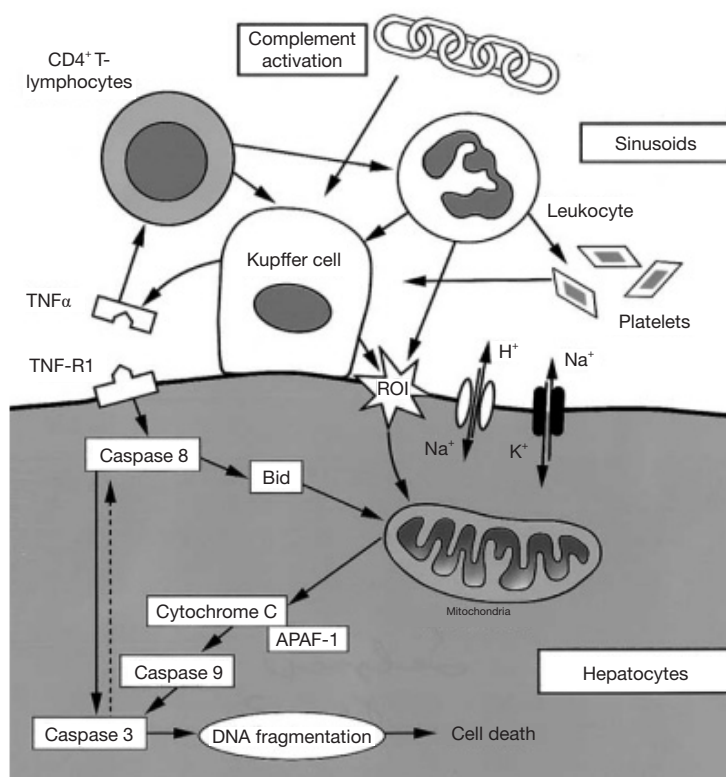


Figure 1 Mechanisms of warm ischemia/reperfusion injury of the liver.

Autophagy, although the first known role is its action during nutrient starvation, is rapidly studied and focused on its role in liver diseases including liver I/R (19-22). Autophagy is an intracellular degradative pathway that targets different cellular components to lysosome for degradation in order to maintain cellular homeostasis and energy supply for substrates. Increasing evidence has linked autophagy to a detrimental role in hepatic I/R injury. It clears out dysfunctional or abnormal mitochondria to make sure optimal cellular function and survival advantage. However, with insufficient or impaired autophagy, damaged mitochondria were accumulated, leading to uncontrolled formation of ROS and destructive mutation of mitochondrial DNA, and ultimately cell death. Hepatic I/R injury could be attenuated by inhibition of autophagy, which is mediated partly through down-regulation of NF- κ B and Akt serine/threonine phosphatase signaling pathways (23-26).

Wip1 (also known as PP2C δ), is a nuclear serine/threonine phosphatase which is encoded by protein phosphatase magnesium-dependent 1 delta, or PPM1D. It belongs to the Ser/Thr PP2C family of phosphatases (27).

Wip1 is a key regulatory protein in DNA repair pathways (28). In response to various stresses including UV, γ -radiation, and alkylating agents, wip1 is activated by MAPK and p53 (29-31). It is not only a direct transcriptional target of p53 but also negatively regulates p53, thus forming a negative regulatory feedback loop (32,33). As a protein phosphatase, activated Wip1 can directly dephosphorylate various proteins including p38 MAPK, checkpoint kinases CHK1/2, NF- κ B, ATM, uracil DNA glycosylase (UNG), MDM2, γ -H2AX, BAX and p53. By dephosphorylating these proteins, Wip1 functions as a key homeostatic regulator that reverses various signaling cascades and induces the damaged cells to re-enter the normal cell cycle following completion of DNA repair (32,34-42). Moreover, Wip1 is emerging as a potent regulator of tumorigenesis as it is frequently amplified/overexpressed in numerous primary human tumors, including breast, ovarian, and pancreatic cancers (43-46). Meanwhile, in addition to programmed cell death and autophagy, Wip1 also plays critical roles in multiplex cellular functions including cell proliferation and senescence, cell cycle arrest, immune-regulation and inflammatory response (47-49). It is found to be expressed in diverse immune-regulating cells like

Table 1 Suzuki's score for I/R injury evaluation

Suzuki's score	Severity (sinusoidal congestion)	Severity (vacuolization of hepatocyte cytoplasm)	Parenchymal necrosis (percentage)
0	None	None	None
1	Minimal	Minimal	Single cell
2	Mild	Mild	~30%
3	Moderate	Moderate	~60%
4	Severe	Severe	>60%

I/R, ischemia/reperfusion.

hematopoietic progenitor cells, neutrophils, macrophages and B/T lymphocytes, and hence *Wip1* knock-out mice undoubtedly displays immunodeficiency and susceptibility to viral infection (50). To date, the role of *Wip1* in liver I/R injury has not been studied yet. In this study, we are aimed to discover the functional role and underlying mechanism of *Wip1* in acute liver injury by utilizing partial (2/3) liver warm I/R injury model.

Methods

Wip1 knock-out mice

In order to generate C57BL *wip1*^{-/-} mice, 129sv genetic background *wip1* knock-out mice were intercrossed with C57BL/6 genetic background wild type mice, then the 129sv/C57BL/6 *wip1*^{+/-} springs were intercrossed with C57BL/6 wild type mice for 7 serial generations to get C57BL/6 *wip1*^{+/-}. Finally, these C57BL6 *wip1*^{+/-} mice were mated to breed C57BL6 *wip1*^{-/-} and *wip1*^{+/+} littermates.

Liver injury assessment

Wip1^{-/-} and their wild-type littermates at 10–14 weeks with the same gender were assigned as experimental group and control group respectively. A 70% hepatic warm I/R model was established and used in this study. Time of ischemia was set as 60 and 90 min. After 6 or 24 h of reperfusion, peripheral blood was collected and mice liver was harvested shortly after euthanasia. The severity of liver injury was assessed through serum ALT levels and Suzuki's scores using H&E staining of the ischemic liver lobes. The TUNEL assay was utilized for evaluation of apoptosis.

Analysis of liver ALT enzyme

The level of serum ALT enzyme was detected with a

chemical analyzer (Olympus AU1000, Tokyo, Japan).

H&E staining

Immediately after collection the liver tissues were fixed in 10% formalin for more than 24 hours, and embedded in paraffin. The tissue blocks were cut into 3- μ m sections. After that, the sections were then stained with hematoxylin-eosin (HE) and the pathological variations were observed under a light microscopy. Suzuki classification was performed to evaluate the severity of I/R injury, consisting three parameters includes sinusoidal congestion, vacuolization of hepatocyte cytoplasm, and parenchymal necrosis. Suzuki's scores were graded numerically and implicated at *Table 1*.

TUNEL assay

Apoptosis of histological sections were detected using TUNEL kit (Beijing Ding Guo Biological Co., Ltd., Beijing, China) according to the manufacturer's instructions. Briefly, the sections were firstly deparaffinated and hydrated, and digested with proteinase K for 20 min. After extensive rinse with 0.01 M PBS, the sections were then incubated with terminal deoxynucleotidyl transferase and digoxigenin-dUTP at 37 °C for 1 h, followed by incubation with a biotin-conjugated anti-digoxin antibody and streptavidin-biotin complex for 30 min each. After extensive washes with PBS, the slides were immersed in DAB solution. All slides were counterstained with Mayer's hematoxylin. Quantitative analysis is calculated as a ratio of TUNEL-positive cells versus total cells which is counted under a microscope field.

Western blot

The expression of *wip1*, Akt, mTOR, p70S6K, S6 and

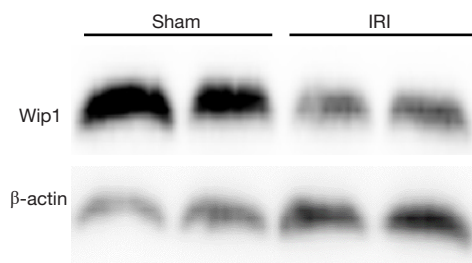


Figure 2 Decreased wip1 protein level after liver ischemia/reperfusion in wild-type mice. Western blot analysis was used to detect the wip1 protein level in wild type mice (n=30) after 60 min of ischemia and 6 h of reperfusion. β -actin was used as an internal control.

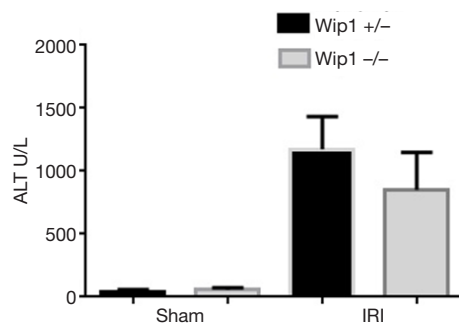


Figure 3 Wip1 knock-out leads to a decreased serum ALT levels.

their phosphorylated protein levels were determined through western blot assay. Briefly, total proteins were extracted from the liver tissues by using appropriate cold lysis buffer based on the manufacturer's instructions. The protein concentrations were determined by BCA method (Beijing Ding Guo Biological Co., Ltd.), following by sample loaded onto 12% SDS-PAGE, separated electrophoretically, and transferred onto a PVDF membrane. After blocking non-specific binding sites with 5% dried skim milk in TBST at room temperature for 1 h, the membrane was then incubated overnight at 4 °C with primary antibodies, followed by incubation with horseradish peroxidase-conjugated secondary antibodies (1:3,000) at room temperature for 1 h. After extensive washes, protein expressions were detected by an enhanced chemiluminescence (ECL) method, imaged using automated wash machine TL-760S (Tyrone medical equipment factory, Taixing, Jiangsu, China), and analyzed by BandScan software. The data were adjusted to correspond internal reference expression [integrated

optical density (IOD) value of target protein versus IOD of correspond internal reference] in order to eliminate the variations of protein expression. All antibodies used in the study were purchased from Cell Signaling Technology (Danvers, MA, USA).

Real-time RT-PCR assay

Total RNA was extracted from the liver tissues by using RNA extraction kit (Omega, Philadelphia, PA, USA) following the manufacturer's protocol followed by real-time RT-PCR assay to detect wip1 mRNA levels. Briefly, reverse transcription (RT-PCR) was performed by using First Strand cDNA Synthesis Kit (Thermo Fisher Scientific, Waltham, MA, USA) according to the manufacturer's instructions, and the level of wip1 mRNA transcription was quantified by real-time PCR with Power SYBR Green PCR Master Mix (Applied Biosystems, Waltham, MA, USA) and ABI 7500 real-time PCR System (Applied Biosystems). Meanwhile, a no-template reaction was used as a background control in parallel and the *GAPDH* gene was selected as the internal control in our study.

Statistical analysis

All of the data were analyzed by using GraphPad Prism statistical software and displayed as means \pm SD. Differences among groups were determined by utilizing one-way ANOVA or unpaired Student's *t*-test. P value <0.05 and <0.01 were considered to be statistical significant.

Results

Wip1 knock-out attenuated liver I/R injury in mice

In order to investigate the role of wip1 in liver I/R injury, firstly we were interested in examining the expression level of wip1 during I/R injury in wild-type mice. As shown in *Figure 2*, after 60 min of ischemia and 6 h of reperfusion, wip1 expression was remarkably reduced in wild-type I/R mice compared with sham-operated mice, indicating that wip1 may be a key regulatory protein during liver injury. In addition, after 90 min of ischemia and 6 h of reperfusion, the wip1 mRNA transcription was also significantly down-regulated in wild-type mice (data not shown). Wip1 knock-out was confirmed by genotyping (data not shown). In order to determine the functional role of wip1 expression in I/R injury, we next investigated the impact of wip1-/-

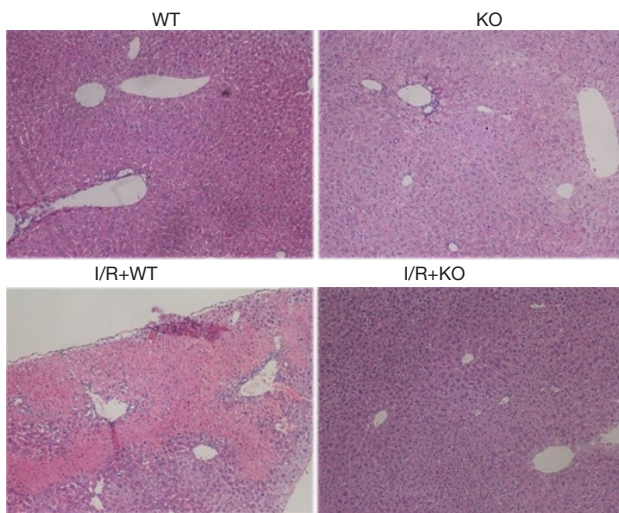


Figure 4 Fewer pathological changes after I/R injury in *wip1^{-/-}* mice. H&E staining was used to evaluate the severity of pathological changes in wild-type mice (left) and *wip1^{-/-}* mice (right) after 90 min of ischemia and 24 h of reperfusion (200 \times).

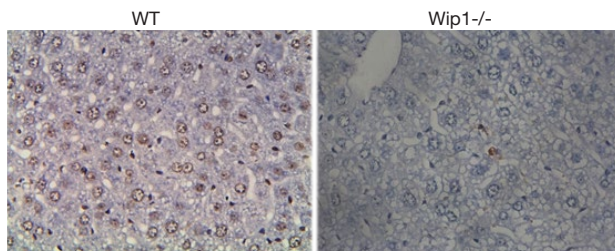


Figure 5 Decreased apoptosis in *wip1^{-/-}* mice compared with WT mice by TUNEL assay. Apoptotic cell morphology was inspected at 400 \times under light microscope.

on liver injury by examining the possible changes of serum ALT levels. As shown in *Figure 3*, after 60 min of ischemia and 6 h of reperfusion, the serum levels of ALT were significantly decreased in *wip1^{-/-}* mice (851.3 \pm 270.9 U/L) compared with wild-type mice (1172.5 \pm 237.1 U/L) ($P < 0.01$), indicating that *wip1* down-regulation in return may protect liver tissues against I/R injury. Moreover, there is no significant difference in serum ALT levels between *wip1^{-/-}* mice (56.6 \pm 12.9 U/L) and wild-type mice (44.3 \pm 11.3 U/L) in sham operated mice ($P = 0.369$), excluding the possibility that operation itself may leads to a decreased ALT level. Hence, we concluded that liver injury was attenuated in *wip1* knock-out mice.

Wip1 knock-out alleviates pathological changes after I/R injury

Next, to confirm the protective role of *wip1^{-/-}* in liver injury, Suzuki's score was conducted and graded numerically to evaluate the severity of liver injury, and morphological changes in sinusoidal congestion, vacuolization of hepatocyte cytoplasm, and parenchymal necrosis were evaluated and detected by H&E staining (see *Table 1*).

As shown in *Figure 4*, after 90 min of ischemia and 24 h of reperfusion, the severity of liver morphological changes following I/R injury was significantly compromised by *wip1^{-/-}*, with the Suzuki's scores of 5.25 \pm 0.43 in *wip1^{-/-}* mice compared with 7.75 \pm 0.43 in wild-type mice ($P < 0.01$), confirming that *wip1^{-/-}* may protect liver tissues against I/R injury.

Wip1^{-/-} led to a decreased programmed cell death following liver I/R injury

Wip1 is an important regulator in a number of critical cellular functions including proliferation, cell cycle arrest, senescence and programmed cell death, autophagy, immunoregulation as well as inflammatory response (47-49), whereas the underlying mechanisms through which *wip1^{-/-}* protected against liver injury was still unknown. We hypothesized that *wip1^{-/-}* may protect liver tissues from I/R injury via suppressing apoptosis. In order to test this hypothesis, TUNEL assay was performed to detect variations of *in situ* apoptotic cell percentage by comparing liver samples from *wip1^{-/-}* mice with wild-type mice. As shown in *Figure 5*, after 90 min of ischemia and 24 h of reperfusion, the rate of apoptosis was unsurprisingly reduced in *wip1^{-/-}* mice (30.4% \pm 3.7%) compared to wild-type mice (62.3% \pm 5.6%) ($P < 0.01$), supporting our hypothesis that *wip1^{-/-}* plays a protective role in liver injury by inhibition of programmed cell death.

Wip1^{-/-} plays a protective role in liver injury via activating PI3K/Akt signaling pathway

The mechanisms of liver I/R injury are complicated by various factors. It can be mediated by release of ROS, lipid peroxidation, cell apoptosis and autophagy, and many other cellular molecules and signaling pathways. Previous studies have demonstrated that *wip1^{-/-}* prevented liver I/R injury by inhibiting apoptosis, yet the mechanism underlying

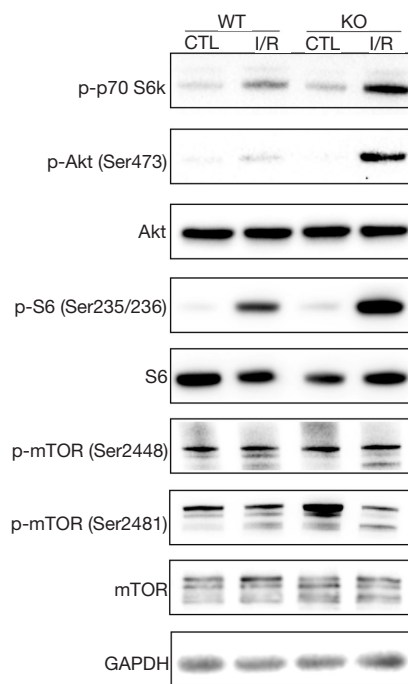


Figure 6 Increased PI3K/Akt activation in *wip1*^{-/-} mice after I/R injury. The expression levels of Akt, p-Akt, mTOR, p-mTOR, S6, p-S6 and p-p70 S6K were detected by western blot, and GAPDH expression was used as an internal control (n=3).

remained unclear. It is well known that the PI3K/Akt pathway plays an important role in cellular survival and apoptosis. Therefore, we planned to investigate whether *wip1*^{-/-} could activate PI3K/Akt pathway to protect liver tissues against I/R injury. The protein levels of Akt, p-Akt, mTOR and p-mTOR are shown in *Figure 6* by comparing wild-type mice or *wip1*^{-/-} mice with sham-operation or I/R injury. After 90 min of ischemia and 6 h of reperfusion, the protein levels of mTOR, p-mTOR (Ser2448), Akt and S6 ribosomal protein showed no significant changes while the protein levels of p-Akt (Ser473), p-p70 S6K (Thr389) and p-S6 (Ser235/236) were clearly increased in the *wip1*^{-/-} mice following I/R injury compared with wild-type mice. Taken together, these results provided strong evidence that *wip1*^{-/-} attenuated apoptosis at least partly by activating the PI3K/Akt pathway.

In summary, our study has revealed that *wip1* expression was suppressed during hepatic I/R injury and *wip1*^{-/-} in return attenuated I/R injury, suggesting a critical role of *wip1* expression in liver self-regulation during I/R injury. Moreover, our study has demonstrated for the first time that *wip1*^{-/-} exerted its protective effect on I/R injury through

inhibiting apoptosis by at least partially activating PI3K/Akt pathway.

Discussion

Hepatic I/R injury is a complex pathological process associated with liver transplantation, shock, and trauma. The process of hepatic I/R injury is a serious threat to human health. I/R injury activate Kupffer cells which generate ROS and lead to release of lysosomal enzymes and various pro-inflammatory cytokines including TNF- α (9,10). Further binding of these cytokines to their respective receptor during early stages of reperfusion or release of oxygen free radicals is shown to initiate the complex apoptotic machinery and eventually leads to cell death of hepatocytes (10).

Wip1 (also known as PP2C δ), is an Mg²⁺-dependent nuclear serine/threonine phosphatase encoded by PPM1D gene (27). Wip1 is a key regulatory protein in DNA repair pathways (28). It is activated by MAPK and p53 in response to a sort of stresses, including UV, γ -radiation, and alkylating agents (29-31). Activated Wip1 induces damaged or injured cells re-enter the normal cell cycle process by directly dephosphorylating various key proteins participating in DNA damage repair including CHK1/2, NF- κ B, p38 MAPK, UNG, ATM MDM2, γ -H2AX, BAX and p53 (32,34-42), indicating its important role in regulating DNA repair pathways. In our study, we have found that *wip1* expression level was significantly reduced after I/R injury, and more importantly, *wip1* knock-out mice presented an ameliorated liver injury, indicating that I/R injury may down-regulate *wip1* expression which in return protect liver tissues from injury. Therefore, *wip1* down-regulation may serve as a self-protective regulator following liver injury.

Several pathways have been reported to associate with I/R injury, including ROS/MAPK pathway (51), ROS/JNK/Bcl-2 pathway and HMGB1/TLR4/NF- κ B pathway. The PI3K/Akt pathway, a well-known cell survival pathway, plays a critical role in the regulation of cell proliferation and cell apoptosis (52). Moreover, the PI3K/Akt signaling pathway exerts a strong protective effect on I/R injury through the inhibition of apoptosis (53-56). We therefore investigated whether *wip1*^{-/-} protected liver tissues against I/R injury through the activation of the PI3K/Akt pathway. To investigate the anti-apoptotic mechanism of *wip1*^{-/-}, we first performed the TUNEL assay to confirm its anti-apoptotic effect after I/R injury, followed by measuring the

protein levels of Akt, mTOR, S6 and their phosphorylated proteins. Our results showed that *wip1-/-* did not change the expression of these proteins at their baseline levels. However, I/R injury up-regulated p-Akt, p-p70 S6K and p-S6 protein levels, and *wip1-/-* further markedly up-regulated these protein level whereas the expression of total Akt and S6 remains unchanged. These results indicated that *wip1-/-* prevent apoptotic cell death probably through activating PI3K/Akt pathway.

Hepatic I/R injury involve various complex and multifactorial mechanisms and these mechanisms need to be further explored. In our present study, we investigated the protective effect of *wip1-/-* on hepatic I/R injury by activating PI3K/Akt pathway. However, other studies have found that Akt as well as NF- κ B activation could also attenuate I/R injury by inhibition of autophagy (23-26), and whether or not *wip1-/-* protects liver tissues from I/R injury by inhibiting autophagy or via other mechanisms like NF- κ B activation remains to be determined. In addition, as a serine/threonine protein phosphatase, *wip1* can directly dephosphorylate various downstream proteins like CHK1/2, ATM and BAX to participate in a number of biological processes, particularly DNA damage response and cell cycle regulation, and thus whether *wip1-/-* exerts its protective effect on I/R injury through other mediators and biological functions waits to be addressed. We are currently working toward answering these questions.

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Footnote

Conflicts of Interest: All authors have completed the ICMJE uniform disclosure form (available at <http://dx.doi.org/10.21037/tcr.2017.12.16>). The authors have no conflicts of interest to declare.

Ethical Statement: The authors are accountable for all aspects of the work in ensuring that questions related to the accuracy or integrity of any part of the work are appropriately investigated and resolved. The study was approved by the Peking Union Medical College Hospital Review Board (No. PUMCH-2014-A230854).

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