



Role of the PI3K/Akt signaling pathway in liver ischemia reperfusion injury: a narrative review

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Objective: To explore the role of phosphatidylinositol-3-kinase/protein kinase B and alpha serine/threonine protein kinase PI3K/PKB (also known as PI3K/Akt) signaling pathway in liver ischemia reperfusion injury.

Background: The PI3K/Akt signaling pathway is one of the major signal transduction pathways that regulates numerous cellular activities *in vivo*. The main functions of this pathway include induction of stem cell differentiation and metastasis, promotion of cell proliferation, inhibition of apoptosis, and regulation of tissue inflammation, tumor growth, and invasion. Liver ischemia reperfusion injury is an inevitable clinical problem that can occur during liver transplantation, liver resection, and various circulatory shock events, and it is one of the primary reasons for postoperative liver dysfunction, and poor disease outcome and patient prognosis. In recent years, it has been found that PI3K/Akt signaling pathway is closely related to liver ischemia reperfusion injury.

Methods: In this review, a large number of relevant literatures were collected to explain the biological basis of PI3K/Akt signaling pathway and its role in liver ischemia reperfusion injury. The review was based on a PubMed search using the terms “liver ischemia reperfusion injury”, “PI3K/Akt signaling pathway”, and “PI3K/Akt signaling pathway AND liver ischemia reperfusion injury”, so as to understand the complex interaction between them.

Conclusions: Activated PI3K/Akt signaling pathway can exert anti-inflammatory, antioxidant stress, anti-apoptosis and autophagy regulation effects through downstream related targeted pathways and proteins, thus alleviating liver ischemia-reperfusion injury. Therefore, the regulation of PI3K/Akt signaling pathway is expected to become an effective targeted pathway for clinical prevention and alleviation of liver ischemia reperfusion injury.

Keywords: Phosphatidylinositol-3-kinase/protein kinase B (PI3K/Akt); liver ischemia reperfusion injury (LIRI); oxidative stress; apoptosis; autophagy

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Introduction

Liver ischemia reperfusion injury (LIRI) commonly occurs during liver surgery, hemorrhagic shock, and other processes, and can lead to severe metabolic disorders, inflammatory immune responses, oxidative stress injury, and cell apoptosis. Studies have indicated that the pathogenesis of LIRI involves a number of different mechanisms, including oxidative stress, inflammatory responses, apoptosis, liver Kupffer cell activation and neutrophil aggregation, vascular cell adhesion molecule overexpression, autophagy, and calcium overload (1,2). The phosphatidylinositol-3-kinase/protein kinase B (PI3K/Akt) signaling pathway is one of the important pathways regulating various cellular functions and plays an important role in numerous physiological processes and pathological responses. The PI3K/Akt signaling pathway is activated during the early stage of LIRI, where it plays anti-apoptosis, anti-inflammatory, anti-oxidative stress, and autophagy regulation roles. In this review, the functional characteristics of this pathway and its role in the various pathogenesis processes of LIRI are discussed.

We present the following article in accordance with the Narrative Review reporting checklist (available at <https://apm.amegroups.com/article/view/10.21037/apm-21-3286/rc>).

PI3K/Akt signaling pathway

General information regarding PI3K

PI3K has dual activities of lipid kinase and protein kinase, which belongs to the lipid kinase family of phosphorylated phosphatidylinositol, a component of eukaryotic cell membranes (3). Based on sequence homology and lipid substrate characteristics, PI3K is divided into three categories (class I, II, and III), as shown in *Figure 1*, where the PI3K I class has been the most thoroughly studied (4). The PI3K I class exists as dimers that are composed of a regulatory subunit and a catalytic subunit. According to their different molecular structures, PI3K proteins can be divided into groups A and B (5,6). Class IA PI3K consists of a heterodimer between the P85 regulatory subunit and the P110 catalytic subunit (7). In mammals, the P85 subunits are further subdivided into α , β , and γ types that are encoded by the PI3K regulatory subunit (PIK3R) 1, PIK3R2 and PIK3R3, respectively. The specific functions of the P85 subunit include receptor binding, enzyme activation, and localization (8-11). The p110 subunit consists of α , β , and δ types that are encoded by the PI3K

catalytic subunits (PIK3C) A, PIK3CB, and PIK3CD, respectively (12). P110 α and p110 β primarily affect cell proliferation and insulin signaling in various tissues, while p110 δ is found only in leukocytes and is involved in immune function and inflammation (9,13). Class IB PI3K consists of the regulatory P101 subunit and the catalytic P110 γ subunit. P110 γ is primarily expressed in white blood cells, and its expression is reduced in the heart, pancreas, liver, and skeletal muscle. The P85 regulatory subunit of class IA PI3K consists of the SH3 domain, the homology of the Rho binding domain/breakpoint cluster, the C-terminal SH2 domain, and the junction region (6,9,14). The PI3K II class does not possess a regulatory subunit and instead only contains a catalytic subunit. This protein exists as three subtypes (α , β , and γ) that contain proline-rich domains, RAS binding domains, HR3 domains, HR2 domains, HR1 domains, PX domains, and C2 domains (6,15-17). Vps34 is the only member of the class III PI3K family, and this protein is a human homologous yeast gene product. Human PI3K is a serine/threonine kinase that is also known as Vps34 (9). Vps34 is a heterodimer formed by the regulatory subunit of P150 and the catalytic subunit of nutmeg acylation of P100, which phosphorylates PI to PI3P and mediates autophagy and protein synthesis under nutritional stress (9,18,19). Other studies have pointed out that although PI3P is widely present, its level does not change when cells are stimulated, so it is considered that PI3K class iii may be a housekeeping kinase that does not play a role in signal transduction (20).

General information regarding Akt

Akt is a serine/threonine kinase that functions as the central mediator of the PI3K pathway and plays a key role in numerous cellular processes, including glucose metabolism, apoptosis, cell proliferation, transcription, and cell migration (21). There are three subtypes of Akt that include PKB α (Akt1), PKB β (Akt2), and PKB γ (Akt3) (22,23), as shown in *Figure 2*. Akt1 is widely expressed in a number of tissues, while Akt2 is primarily expressed in insulin-sensitive tissues such as skeletal muscle, adipose tissue, and liver, and Akt3 is predominantly expressed in the testes and brain. Akt consists of a pH domain, an intermediate kinase domain, and a regulating carboxyl terminal domain in which the pH domain regulates Akt transposition (24). Akt is activated through two key phosphorylation processes. PIP3 as an intracellular second messenger, and the structure of the pH domain within Akt can facilitate binding to PIP3 to cause

Category	Component	Function
Class IA PI3K		
Regulatory subunit	<p>p85α/β — SH3 — P — BH — P — SH2 — iSH2 — SH2 —</p> <p>p55α/p50α — SH2 — iSH2 — SH2 —</p> <p>p55γ — SH2 — iSH2 — SH2 —</p>	Affecting cell proliferation and insulin signaling, immune function and inflammation
Catalytic subunit	p110 α / β / δ — ABD — RBD — C2 — PIK — Catalytic subunit —	
Class IB PI3K		
Regulatory subunit	p101 — p110 γ binding — G β γ binding —	Regulating leukocyte function
Catalytic subunit	p110 γ — RBD — C2 — PIK — Catalytic subunit —	
Class II PI3K		
Catalytic subunit	<p>PI3KC2α/β — CB — RBD — C2 — PIK — Catalytic subunit — PX — C2 —</p> <p>PI3KC2γ — RBD — C2 — PIK — Catalytic subunit — PX — C2 —</p>	Affecting cell migration, proliferation and survival
Class III PI3K		
Catalytic subunit	Vps34 — C2 — PIK — Catalytic subunit —	Mediating autophagy and protein synthesis under nutritional stress

Figure 1 Structural overview of each type of PI3K. SH3, Src homology 3; P, proline-rich region; BH, breakpoint cluster region homology; SH2, Src homology 2; iSH2, interSH2; ABD, adapter binding domain; RBD, Ras-binding domain; PI3K, phosphatidylinositol-3-kinase; CB, clathrin binding domain; PX, phox homology.



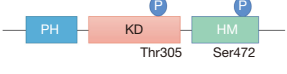
Category	Component	Distribution	Function
Akt1/PKB α		Widely distributed in the organism	Influencing placenta development, animal growth and adipogenesis
Akt2/PKB β		Insulin-sensitive tissues such as liver, skeletal muscle and adipose tissue	Maintaining glucose homeostasis
Akt3/PKB γ		Brain and testis	Affecting brain development after birth

Figure 2 The structure and function of each type of Akt. PH, pleckstrin homology domain; KD, kinase domain; HM, hydrophobic motif.

a conformation change in AT to expose the Ser473 and Thr308 sites to phosphoinositide dependent protein kinase 1 (PDK1) phosphorylation of the kinase Thr308 structure. This further facilitates the Akt activation process, where mammalian target of rapamycin (mTOR) compound 2 (mTORC2) can initiate Ser473 phosphorylation facilitates additional adjustment structure of the domain at the terminus of the carboxyl region, thereby fully activating Akt (25). Similar phosphorylation events were observed on the corresponding residues of Akt2 (Thr309 and Ser474) and Akt3 (Thr305 and Ser472). Phosphorylation of both

residues is necessary for maximum activation of Akt, and protein 2A phosphatase (PP2A) and leucine-rich repeat protein phosphatase function to dephosphorylate Thr308 and Ser473 of Akt, respectively, thus leading to Akt inactivation (26).

Types and stages of LIRI

LIRI is mainly divided into two types: (I) warm ischemia reperfusion injury, which occurs during liver resection, shock or trauma caused by various reasons, and can lead

to liver or even multiple organ failure; (II) cold ischemia reperfusion injury usually occurs during organ preservation and transplantation. Historically, it has been believed that liver cells are more sensitive to warm ischemia reperfusion, while liver sinusoidal endothelial cells (LSEC) are more sensitive to cold ischemia reperfusion. However, limitations on damage to specific cell types are rare under pathophysiological conditions (1,27).

LIRI includes two stages of ischemia and reperfusion: ischemia leads to decreased intracellular 5'-adenosine triphosphate (ATP) concentration, ATP-dependent Na⁺/K⁺ pump failure, cell edema, and increased cytoplasmic calcium concentration, all of which leads to cell damage. Liver reperfusion enhances cellular damage through oxidative stress and inflammatory response. The initial stage of ischemia reperfusion injury (within 2 hours after reperfusion) involves Kupffer cells releasing reactive oxygen species (ROS) and pro-inflammatory mediators such as tumor necrosis factor (TNF)- α , interleukin (IL)-6, IL-1, and arachidonic acid. ROS may lead to lipid peroxidation, mitochondrial permeability conversion, signal transduction pathway activation, caspase activation, hepatocyte and endothelial cell necrosis and apoptosis. Late stages (6–48 h after reperfusion) are characterized by neutrophil-mediated inflammatory responses. Cytokines, chemokines, and complement factors recruit neutrophils into the liver, where they damage liver cells by releasing ROS or proteases. Endothelin-1 (ET-1)-mediated vasoconstriction and increased platelet aggregation lead to microcirculation disorders and aggravate liver injury. Although there is a chronological sequence of events that occur during LIRI, these events may overlap or occur simultaneously in the body (28-30).

PI3K/Akt signaling pathway and LIRI

PI3K/Akt signaling pathway and anaerobic metabolism and acidosis

When the liver is in the ischemic stage, the tissue is in a state of ischemia and hypoxia, and cell normal redox process is blocked, the ATP in the cell was quickly consumption, anaerobic glycolysis increase, lead to the accumulation of lactic acid and ketone body acidic metabolites, and mitochondrial oxidative phosphorylation function decline, the organization cell pH decline, metabolic acidosis. After refluxes, the pH value returned to normal, and a large number of pH-dependent proteases, phospholipases

and other activities increased, but aggravated tissue and organ damage, resulting in LIRI, which is one of the most common mechanisms of LIRI (27).

Rafiee *et al.* found that acidic microenvironment can activate the PI3K/Akt signaling pathway and significantly enhance the phosphorylation of Akt (31). After stimulated by factors such as hypoxia and acidic environment, the regulatory subunit P85 of PI3K relieved the inhibition of catalytic pressure group P110, and the spatial conformation of PI3K dimer changed and was then activated, which promoted the formation of phosphatidylinositol 3,4,5-triphosphate (PIP3) on the membrane of PI3K. Signaling proteins are then recruited to regulate various intracellular activities such as cell survival, proliferation and differentiation (15).

Metabolic acidosis occurs during LIRI, and the acidic microenvironment affects the activity and function of lymphocytes. Studies have found that regulatory T cells (Tregs) can significantly relieve the inflammatory response caused by LIRI, and have a certain protective effect on injured organs (32-34). The acidic microenvironment of LIRI down-regulated the function of Tregs and promoted the apoptosis of Tregs. It is known that V-ATPase located in internal acidic vesicles and plasma membrane plays a role in regulating pH gradient across membrane (35,36). Tregs expressed more V-ATPase in acidic microenvironment. Some scholars have found that omeprazole can inhibit PI3K/Akt/mTOR signaling pathway by reducing v-ATPase expression, improve the induction efficiency of Tregs in acidic microenvironment, and then improve LIRI acidic microenvironment and reduce LIRI (37,38).

PI3K/Akt signaling pathway and oxidative stress

Due to ischemia and hypoxia that occur during LIRI, oxygen free radicals are stimulated through xanthine oxidase system, phagocyte system and mitochondrial respiratory chain during reperfusion, resulting in increased production of ROS and insufficient scavenging capacity, thus resulting in liver cell damage. Increased ROS can damage liver cells by generating a variety of lipid peroxides that damage the structure and function of LSEC, cause DNA mutations in the nucleus of liver parenchyma tissues, inhibit oxidative phosphorylation of mitochondria, and cause microcirculation disorders (2,39-41).

PI3K is a redox sensitive kinase that can be activated by changes in intracellular ROS levels (42). During the process of LIRI, the increase in intracellular ROS and

oxidative stress levels cause compensatory activation of the PI3K/Akt signaling pathway. The activated PI3K/Akt signaling pathway promotes the generation of heme oxygenase-1 (HO-1) by upregulating the expression of Nrf2, thus alleviating oxidative stress injury during LIRI (43-47). In addition, activation of PI3K/Akt pathway can activate endothelial nitric oxide synthase (eNOS) in vascular endothelial cells, increase the formation of nitric oxide (NO), and further improve LIRI (48-50).

PI3K/Akt signaling pathway and inflammatory response

In the early stages of ischemia reperfusion, Kupffer cells are activated, which, in addition to producing a large number of ROS, increases the release of pro-inflammatory chemokines and cytokines that promote and amplify subsequent neutrophil-mediated inflammatory responses (51,52). Stimulated by pro-inflammatory factors and chemokines at the early stage, neutrophils adhere to and aggregate in the liver space. These neutrophils release a large number of proteases (e.g., collagenase, elastase, cathepsin G) that act on cell membranes and matrix components, thereby promoting cellular degradation (53). Neutrophils also drive inflammation of liver damage by releasing inflammatory mediators, including proteolytic enzymes, lipocalin 2, arachidonic acid metabolites, and ROS (54); In response to Mac-1 adhesion, neutrophils release myeloperoxidase (MPO) and proteases through NADPH oxidase and thrashing, leading to superoxide formation. It can directly cause liver endothelial injury or indirectly induce tissue injury by triggering other inflammatory mediators (55).

During LIRI, Kupffer cell activation significantly increases the expression of high-mobility group B1 (HMGB1), and this activates Toll-like receptor 4 (TLR4) to trigger the recruitment of MyD88 and subsequently promote the activation of downstream signaling pathways, ultimately leading to significant increases in NF- κ B, TNF- α , and IL-6 (56). PI3K can inhibit the activation of NF- κ B and its downstream pro-inflammatory cytokines by inactivating TLR4 signaling through preventing the recruitment of Toll-IL-1 resistance domain junction protein (TIRAP) to the cell membrane, thus alleviating LIRI (57-59). Blocking of PI3K/Akt leads to increased transcription of NF- κ B and the release of TNF- α , IL-1 β , and IL-6 to ultimately exacerbate LIRI (60). Shen *et al.* determined that activation of the PI3K/Akt signaling pathway during LIRI increased the expression of IL-4 and IL-10, decreased the expression of IL-1 β and TNF- α ,

and reduced the liver inflammatory response (50). Our recent study found that methyl eugenol can reduce LIRI by activating the PI3K/Akt signaling pathway, inhibiting the expression of inflammatory factors and reducing the apoptosis rate (61).

PI3K/Akt signaling pathway and apoptosis

Apoptosis is closely related to LIRI, and apoptosis of hepatocytes and hepatic sinusoidal endothelial cells is an important cause of LIRI. The process of apoptosis is extremely complex, and the generation of oxygen free radicals, cell energy metabolism disorder, intracellular calcium overload, the activation of cytokines and caspase and B lymphocyte tumor-2 (Bcl-2) family gene expression is closely related to the mechanism disorder will induce a variety of diseases. Disorders of the mechanisms underlying apoptosis, including the death receptor pathway, the mitochondrial pathway, the perforin/granzyme pathway, and the endoplasmic reticulum pathway, may induce various diseases. The mitochondrial pathway, also known as endogenous apoptosis pathway, is the most important pathway for mediating apoptosis. When cells receive a variety of endogenous apoptosis signal stimulations, the Bcl-2 protein regulates mitochondrial permeability by irreversibly opening the mitochondrial permeability transition pore (mPTP), ultimately resulting in clearance of cytochrome C into the cytoplasm. Apoptosis protein activated factor 1 (Apaf-1) then interacts with this released cytochrome C to facilitate increased levels of the caspase 9 precursor that then becomes active through an auto-cleavage process. Activated caspase 9 activates a downstream caspase cascade that is mediated by caspase 3 to induce apoptosis (62,63).

The PI3K/Akt signaling pathway is a classical antiapoptotic pathway. When the PI3K/Akt signaling pathway is activated, Akt inhibits caspase 3-mediated cell death by phosphorylating Bcl-2/Bcl-XL-associated death promoter (Bad). Concurrently, the depolymerization of Bcl-2 with phosphorylated Bad further promotes cell survival. Studies have demonstrated that activation of the PI3K/Akt/mTOR signaling pathway can significantly increase Bcl-2 protein levels, reduce the expression of Bax and caspase 3 proteins, protect cells from apoptosis, and significantly improve LIRI (3,27,64-68).

PI3K/Akt signaling pathway and autophagy

Autophagy is a lysosomal dependent catabolic process that

plays an important role in regulating cell homeostasis by degrading damaged organelles and misfolded proteins. Autophagy is not only important for the dynamic balance of proteins, but it also acts a quality control system under internal and external stress conditions and is closely related to the occurrence and development of a variety of physiological processes (such as nutritional deficiency, ischemia, hypoxia, and others) and diseases (such as neurodegenerative diseases, immune diseases, and others). During LIRI, autophagy can be activated by adverse factors such as reduced ATP synthesis, elevated production of ROS, and calcium overload. Mitochondrial autophagy can rapidly remove the aging and damaged mitochondria and excess oxygen free radicals through the lysosomal mechanism to thus maintain the normal function of mitochondria and alleviate LIRI. However, when the content of oxygen free radicals and calcium ions exceeds the clearance rate of autophagy, autophagy cannot clear all damaged mitochondria. This eventually leads to irreversible cell death (69). Currently, it remains undetermined whether autophagy plays a protective or injurious role in LIRI. It should be noted that class I PI3K and class III PI3K play different roles in autophagy regulation. Class I PI3K inhibits autophagy through PI3K/Akt/mTORC1 pathway, while class III PI3K enhances autophagy by inducing Beclin1 (70).

It has been demonstrated that Akt regulates autophagy levels by directly phosphorylating and inhibiting adenosine monophosphate-activated protein kinase (AMPK)-mediated phosphorylation of tuberous sclerosis complex 2 (TSC2), thus inducing complete inhibition of TSC2 and activation of mTOR (71). Yang *et al.* found that downregulation of the PI3K/Akt/mTOR signaling pathway resulted in increased expression of Beclin-1, Atg7, and LC3 and decreased expression of p62, thus indicating that autophagy can be properly enhanced to protect the liver from ischemia-reperfusion injury (72). A recent study showed that octreotide and melatonin can enhance autophagy by regulating AMPK/PI3K/Akt/mTOR/ULK1 signaling, and subsequently reduce LIRI (73).

Liu *et al.* revealed that shikonin treatment could improve LIRI by upregulating the expression of PI3K and p-Akt and inhibiting autophagy. PI3K can upregulate the expression of the Bcl-2 protein, and the Bcl-2/Beclin-1 complex plays a key role in the regulation of apoptosis and autophagy (74). A recent study demonstrated that upregulation of Bcl-2 could inhibit the disassociation of Beclin-1 and Bcl-2, thus leading to decreased Beclin-1 expression and inhibition

of autophagy (75). Various doses of shikonin significantly decreased the expression of LC3 and Beclin-1 and increased the expression of P62, thus suggesting that shikonin treatment can inhibit autophagy by activating the PI3K/Akt pathway and can thereby prevent hepatic ischemia reperfusion injury (64).

PI3K/Akt signaling pathway and heat shock proteins (HSPs)

HSPs are also known as emergency proteins and exist as a group of structurally highly conserved peptides. HSPs possess a variety of biological activities, and their most important role is to act as molecular chaperones where they aid in facilitating the correct folding of newly synthesized proteins, influence the refolding of proteins destroyed by stress, and maintain the stability of various protein structures. Studies have revealed that there are multiple HSP expressions during LIRI (76,77). HSP70 is the most highly conserved and most important HSP. The expression level of HSP70 in normal cells is typically low. In response to various stimuli such as ischemia, hypoxia, and inflammatory responses, HSP70 can migrate from the cytoplasm to the nucleus and become expressed at high levels. HSP70 possesses immunomodulatory properties, and in combination with IL-2, it can enhance the ability of Tregs to increase the secretion levels of the immunosuppressive cytokines IL-10 and TGF- β . Human HSP70 enhances the regulatory activity of Tregs primarily through modulating the interaction between TLR2 and T cells, and PI3K/Akt is involved in HSP70-dependent activation of Tregs (78). Zheng *et al.* revealed that HSP70 can protect the liver from ischemia-reperfusion injury by stimulating the PI3K/Akt signaling axis to regulate the differentiation of Tregs (79).

PI3K/Akt signaling pathway and non-coding RNA

Non-coding RNA refers to a type of RNA that does not participate in encoding proteins during the process of gene expression transcription and instead plays an important role in the process of transcription and post-transcription modification. During LIRI, the expression of certain non-coding RNAs within cells becomes abnormal, and these non-coding RNAs are primarily involved in cell energy metabolism, apoptosis, autophagy, oxidative stress responses, and inflammation (80).

MicroRNAs (miRNAs) can recognize and target mRNA to regulate the expression of target genes at the

post-transcriptional or translational level. Studies have demonstrated that miR-494 can upregulate the PI3K/Akt pathway by targeting PTEN to reduce cell apoptosis and improve LIRI in rats (81).

Long non-coding RNAs (lncRNAs) can interact with all types of intracellular macromolecules, including proteins, DNA, and RNA. lncRNA can be used as signal, bait, guide, skeleton, and enhancement molecule to modulate gene expression at the transcriptional and post-transcriptional levels. Cui *et al.* demonstrated that lncRNA H19 could enhance autophagy-induced hypoxia/reoxygenation injury in liver cancer cells through downregulation of the PI3K/Akt/mTOR pathway (82).

The main biological function of small interfering RNA (siRNA) is to participate in RNA interference (RNAi). RNAi induces gene silencing by targeting mRNA for degradation or by preventing mRNA from being translated into proteins. During the process of LIRI, siRNA-induced β -catenin deficiency and STAT3 silencing are used to inhibit PTEN and activate the PI3K/Akt pathway, and this can downregulate the immune function of dendritic cells (DCs), inhibit TLR4-driven inflammatory responses, and reduce cell apoptosis (83,84).

PI3K/Akt signaling pathway and exosomes

Exosomes are a class of tiny vesicles possessing a lipid bilayer structure and ranging in size from 30 to 150 nm in diameter. Exosomes can transmit information between cells, and the proteins, DNA, and RNA that they carry can be absorbed by the recipient cells to facilitate specific functions. Studies have revealed that hepatocyte-, DC-, and mesenchymal bone marrow stem cell-derived exosomes can mitigate LIRI through different mechanisms (79,85,86). DC-derived exosomes inhibit the immune inflammatory cascade and reduce inflammation-induced hepatocyte necrosis, and Zheng *et al.* revealed that DC-derived exosomes can activate the PI3K/Akt/mTOR signaling pathway in the initial CD4⁺ T lymphocytes by delivering HSP70 to these lymphocytes. This induced the differentiation of the initial CD4⁺ T lymphocytes into Tregs and indirectly inhibited the differentiation of the initial CD4⁺ T lymphocytes into Th17 cells. Amplified Tregs can induce the formation and maintenance of immune tolerance, and at the same time, the reduction of Th17 production can further inhibit the production of immune inflammatory cascade, thus inhibiting the necrosis and apoptosis of liver cells induced by inflammation to

ultimately improve LIRI (79).

Conclusion and prospects

LIRI is a serious clinical problem that is considered to be the potential mechanism underlying the dysfunction and injury of organs other than the liver itself. Recent studies demonstrated that the PI3K/Akt signaling pathway is closely associated with LIRI, and PI3K/Akt signaling can improve LIRI by influencing the anti-oxidative stress responses, anti-inflammatory processes, regulation of immune cells, regulation of autophagy, and reduction of cell apoptosis (as shown in *Figure 3*).

However, in the process of LIRI, how the PI3K/Akt signaling pathway coordinates the functions of the above pathological phenomena and whether there are interactions between these phenomena still need to be further discussed. In addition, most current studies are only limited to the involvement of PI3K/Akt signaling pathway in LIRI, but how PI3K/Akt signaling pathway is activated during LIRI has not been clarified. In addition, most of the current studies are focused on animal models, and different experimental conditions and modeling will inevitably produce different experimental results. Whether various intervention methods have the same safety and effect in clinical application remains to be further studied.

It is noteworthy that the abnormal activation of PI3K/Akt signaling pathway is involved in the regulation of tumor cell survival, proliferation, invasion and migration in the process of various tumors. By inhibiting the expression of each molecule in this pathway and blocking this signaling pathway, tumor growth and metastasis can be inhibited and anti-tumor effect can be achieved (87,88). Therefore, PI3K/Akt will show different effects in different diseases and cells, so clinical intervention of PI3K/Akt signaling pathway needs to be dealt with in combination with the actual situation.

Therefore, in the future, studies on PI3K/Akt signaling pathway in LIRI can focus on the following points: (I) what is the specific molecular mechanism by which PI3K/Akt signaling pathway is activated or inhibited during LIRI? (II) Whether targeted intervention of PI3K/Akt signaling pathway can reduce liver ischemia-reperfusion injury in large animal experiments, such as pigs and monkeys, or even in clinical process? (III) In the prevention and treatment of LIRI, whether there will be adverse reactions after activation of PI3K/Akt signaling pathway, and how to evaluate its safety? (IV) For liver cancer patients, is it

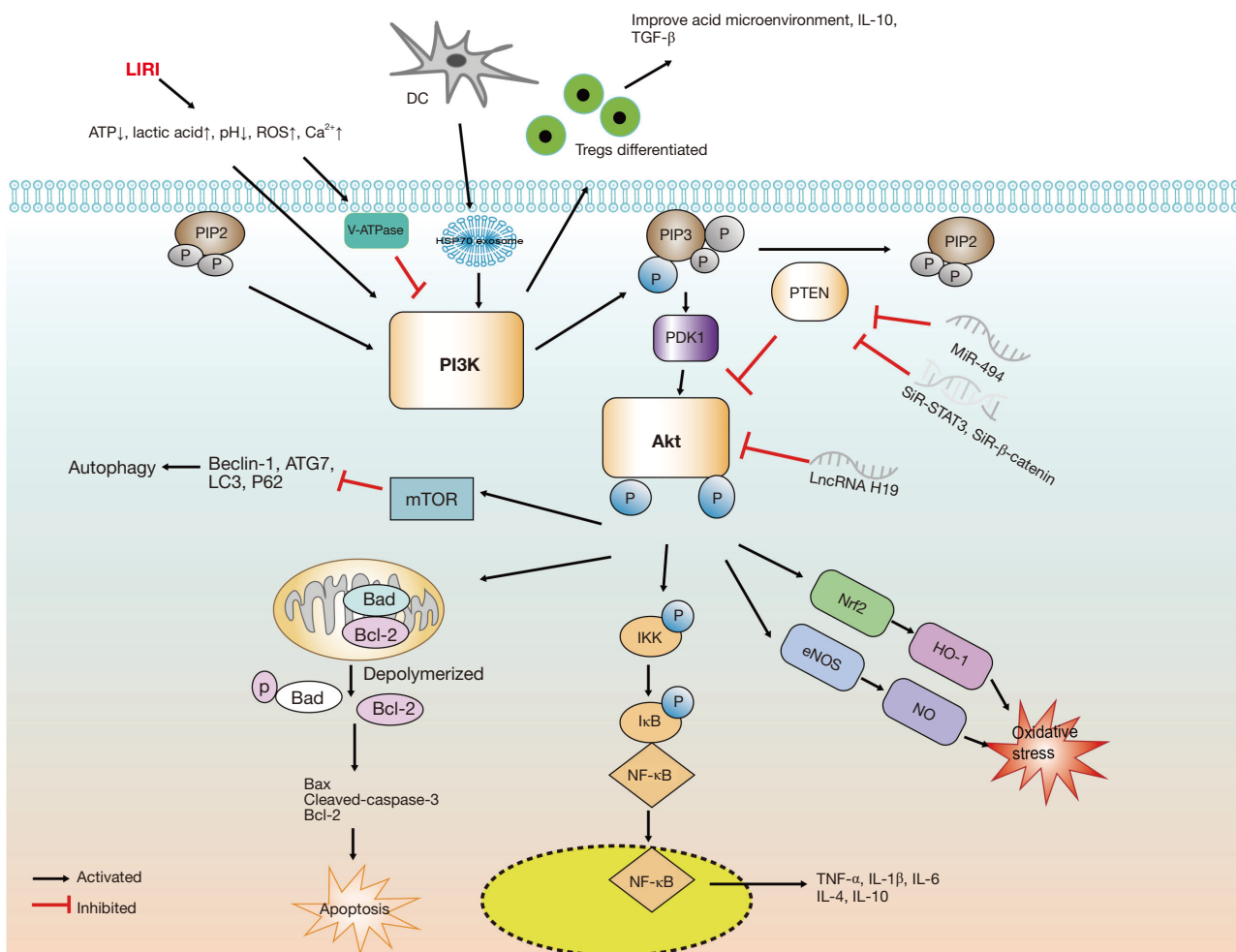


Figure 3 The relationships between liver ischemia reperfusion injury and PI3K/Akt signaling pathway. LIRI, liver ischemia reperfusion injury; ATP, adenosine triphosphate; ROS, reactive oxygen species; DC, dendritic cell; IL, interleukin; TGF, transforming growth factor; PIP2, phosphatidylinositol-4,5-diphosphate; PIP3, phosphatidylinositol-3,4,5-triphosphate; PTEN, phosphatase and tensin homolog; PDK, phosphoinositide dependent protein kinase; PI3K, phosphatidylinositol-3-kinase; Akt, alpha serine/threonine protein kinase; mTOR, mammalian target of rapamycin; ATG, autophagy-related protein; LC3, autophagy microtubule-associated protein light chain β ; Bad, Bcl-2/Bcl-XL-associated death promoter; Bcl-2, B lymphocyte tumor-2; IKK, I κ B kinase; I κ B, inhibitor proteins of NF- κ B; NF- κ B, nuclear factor- κ B; Nrf2, nuclear factor E2 related factor 2; HO-1, heme oxygenase-1; eNOS, endothelial nitric oxide synthase; NO, nitric oxide; TNE, tumor necrosis factor.

feasible to intervene PI3K/Akt signaling pathway to prevent and treat LIRI?

Although there are still some unclarified questions about the effect of PI3K/Akt signaling pathway on LIRI, the PI3K/Akt signaling pathway has the potential to become a target for the prevention and treatment of LIRI, providing a research direction for the prevention and treatment of ischemia reperfusion.

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Footnote

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