



Alteration of *Connexins* Gene Expression by Silibinin during Hepatic Warm Ischemia-Reperfusion Injury in a Rat Model

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Abstract

Background: Ischemia-reperfusion injury (IRI) is an injurious phenomenon that is the primary determinant of liver dysfunction after surgery and transplantation. The present evidence demonstrated that *connexin 43* (*Cx43*), *Cx32*, and *Cx26* are the essential gap junction proteins involved in the liver IRI.

Objectives: This study aimed to characterize the beneficial effects of silibinin on *Cx43*, *Cx32*, and *Cx26* gene expression during warm hepatic ischemia-reperfusion (IR).

Methods: A total of 32 male Wistar rats weighing 250-300 g were randomly divided into four equal groups of eight animals in each group as follows: 1) control (laparotomy+normal saline), 2) laparotomy+silibinin (30 mg/kg) (SILI), 3) liver IR procedure+normal saline (IR), and 4) liver IR procedure+silibinin (30 mg/kg) (IR+SILI). After 1 h of ischemia followed by 3 h of reperfusion, blood samples and tissue sections were gathered to assess the serum liver markers and evaluate the liver histological changes as well as gene expression, respectively.

Results: The obtained data proved no considerable differences between control and SILI groups in all experiments. Furthermore, the gene expression of *Cx26*, *Cx32*, and *Cx43* was significantly induced in the IR group, compared to the control group. Silibinin markedly reduced *Cx26* and *Cx32* mRNA expression, whereas increased *Cx43* mRNA expression. Moreover, serum alanine aminotransferase and aspartate aminotransferase levels were markedly elevated in the IR group ($P < 0.001$), compared to the control group. However, in the IR+SILI group, silibinin could significantly decline these elevations, compared to the IR group. In addition, silibinin diminished hepatic tissue damages during IR.

Conclusion: Silibinin could attenuate liver injury through better cell-to-cell communication via lowering *Cx32* and *Cx26*, as well as increasing *Cx43* gene expression, respectively.

Keywords: Connexin, Gap junction, Ischemia-reperfusion, Silibinin

1. Background

Liver ischemia-reperfusion injury (IRI) is a pathophysiological process that occurs during partial or total obstruction of hepatic blood flow and oxygen depletion in the liver tissue and is sometimes necessary during surgical procedures. The renovation of blood flow to the ischemic liver tissue and re-oxygenation triggers a molecular cascade that provokes further hepatocellular dysfunctions or even death (1). During reperfusion, many molecular mechanisms are involved in the pathogenesis of liver IRI and hepatocellular damage, including anaerobic metabolism, oxidative stress, mitochondrial dysfunction, intracellular Ca^{2+} overload, inflammatory response, neutrophil, and macrophage infiltration (2, 3). However, the actual molecular mechanisms of hepatic IRI have remained obscure.

Gap junctions (GJs) are specialized intercellular channels that directly connect the cytoplasmic compartments of the two adjacent cells and allow rapid intercellular exchange of ions and small molecules <1000 Da, highlighting their role in the

homeostasis of cells and tissues (4). A GJ channel is made up of two connexons or hemichannels from the two adjacent cells in a head-to-head organization. Each connexon contains six transmembrane protein monomers, which are called *connexin* (*Cx*). According to their molecular weight, *Cxs* are encoded by a conserved gene family with 21 different members in mammals, which are expressed in a cell-dependent pathway (5,6). The *Cx26*, *Cx32*, and *Cx43*, the building units of liver GJ, are expressed in the mature hepatocytes and Kupffer cells (5). A previous study has reported that *Cx32* plays an essential role in hepatic IRI by modulating hepatocyte apoptosis and damage (7). In acetaminophen-induced liver toxicity, *Cx43* can lead to effective intercellular communication as part of the hepatic protective response (8). Moreover, during hepatic IRI and inflammation, an increase occurs in the hepatic *Cx43* expression leading to the converse changes in *Cx32* and *Cx26* expression (8-11).

2. Objectives

Silibinin, the principal active constituent found in

silymarin, is extracted from the milk thistle plant (*Silybum marianum*). It has several therapeutic uses, including hepato-protective, anti-cancer, anti-oxidant, and anti-inflammation (12-14). In general, studies have shown that silibinin could suppress the tumor necrosis factor- α (TNF- α) mRNA and protein expression. On the other hand, TNF- α represses the expression of *Cx43* in cardiac cells; therefore, silibinin can indirectly impede the repressive effect of TNF- α on the *Cx43* gene (15). Accordingly, this study aimed to investigate the silibinin effect on *Cx43*, *Cx32*, and *Cx26* gene expressions and liver tissue damages during IR in a rat model.

3. Methods

A total of 32 mature male Wistar rats weighing 250-300 g were obtained from the Research Center and Experimental Animal House of Mazandaran University of Medical Sciences, Sari, Iran. All of the experimental methods of Animal Breeding and Research of Mazandaran University of Medical Sciences were approved by the Ethical Committee of Mazandaran University of Medical Sciences, Sari, Iran. The animals were maintained and adapted in the laboratory under normal and standard conditions (12 hours of light/dark cycles and moisture of 55% \pm 5 at 23 \pm 2 $^{\circ}$ C). After a one-week adaptation, the rats were randomly assigned into four groups each containing eight rats. These groups included: 1) Control (Vehicle) (animals were under laparotomy and received 0.9% sodium chloride [normal saline] intraperitoneally [IP] 30 min before laparotomy and immediately after the liver inserted into the abdominal cavity of the body), 2) Silibinin (SILI) (animals were under laparotomy and received 30 mg/kg silibinin 30 min before laparotomy and immediately after the liver inserted into the abdominal cavity of the body), 3) Ischemia-reperfusion (IR) (animals received normal saline 30 min before ischemia and immediately after reperfusion), and 4) IR+SILI (animals received silibinin [30 mg/kg] before ischemia and immediately after reperfusion).

3.1. Surgical procedure

All animals were under fasting for 18 h before surgery; however, drinking water was unlimitedly accessible. They were then anesthetized by a mixed solution with an IP injection of ketamine (50 mg/kg) and xylazine (8 mg/kg). After creating a longitudinal incision on the ventral midline and bringing out the liver in IR and IR+SILI groups, the left branch of the triad port (including hepatic artery, portal vein, and biliary duct) was obstructed by a single metallic bulldog clamp for 60 min to create complete ischemia in the median and left lobes. Subsequently, the clamp was removed, and the liver was inserted into the abdominal cavity of the body to allow reperfusion for

3 h. The Vehicle and SILI groups were prepared similarly; however, vascular clamping was not applied. After the experimental periods, the animals were sacrificed, and the blood samples, as well as the liver tissue sections, were collected for further studies. Liver tissues were stored in RNAlater solution at -70 $^{\circ}$ C. Furthermore, the serum separated from the blood samples was stored at -20 $^{\circ}$ C.

3.2. Silibinin administration

Silibinin (~98% purity) was purchased in a lyophilized powder form with a dihydrogen succinate disodium salt formula (Legalon) manufactured by Rottapharm/Madaus (Cologne, Germany). It was injected twice (IP) in a total dosage of 60 mg/kg 1 h before laparotomy and immediately after the beginning of reperfusion each time 30 mg/kg (16,17).

3.3. Biochemical analysis

Serum levels of alanine aminotransferase (ALT) and aspartate aminotransferase (AST) enzymes were measured by a biochemical auto analyzer (BT-3000-plus, Biotechnica, Italy) using the Pars Azmoon kit (Karaj, Iran).

3.4. Total RNA extraction and real-time PCR

According to the manufacturer's guidelines, an RNeasy plus mini kit (Qiagen, Germany) was used to extract the total RNA of all liver tissue samples. A UV spectrophotometer (Thermo Scientific, USA) was also utilized to evaluate RNA concentration at 260 nm and its purity at an absorbance ratio of 260/280. In addition, RNA quality was confirmed by two sharp-band detections for 18S and 28S ribosomal RNA by resolving electrophoresis in agarose gel stained with SYBR Green. According to the kit protocol, a concentration of 1 μ g of RNA per reaction was applied for cDNA synthesis (EURx, Poland). The reverse transcription Real-time PCR was performed according to the following steps: 50 ng of cDNA (2 μ l), 10 pM of specific primers (1 μ l of forward and reverse), 12.5 μ l SYBR Green PCR Master Mix reagent (EURx, Poland), and DD water up to 25 μ l total volume. The PCR cycles were as follows: UNG pre-treatment at 50 $^{\circ}$ C for 2 min, initial denaturation at 95 $^{\circ}$ C for 12 min, and 40 cycles (95 $^{\circ}$ C for 15 sec [denaturation]), 58 $^{\circ}$ C for 30 sec (annealing), and 72 $^{\circ}$ C for 30 sec (final extension). It is worth mentioning that the *GAPDH* gene was used for the normalization of the results, and the sequences of the primers are listed in Table 1.

3.5. Histopathological study

Liver tissue sections were fixed in 10% phosphate-buffered saline formalin, dehydrated with increasing ethanol grades, washed with xylene, and finally embedded into paraffin blocks for sectioning. The thin sections, about 3-5 μ m, were cut from paraffin-embedded samples with a microtome.

Table 1. Primer sequences

Genes	Forward Primer	Reverse Primer	Product Length
<i>Cx26</i>	5'- CACTTCTGACCAACCCAGGAG -3'	5'- GCTCTGTAGTGTGCCCAAT -3'	192
<i>Cx32</i>	5'- CAGACAGCCTGCATACATTC-3'	5'- TAGAATGCCGATTCACGCCA-3'	133
<i>Cx43</i>	5'- CATTGGGGAAAGGCGTGAG-3'	5'- GAAGCTTCCCCAAGGCACTC-3'	204
<i>GAPDH</i>	5'- AGTGCCAGCCTCGTCTCATA-3'	5'- GATGGTGATGGGTTCCCGT-3'	248

After deparaffinization with xylene, samples were stained with Hematoxylin and Eosin. Ultimately, all prepared slides were observed under an optical microscope (18).

3.6. Statistical Analysis

The data were analyzed in SPSS software (version 18), and REST-RG software was used to analyze the real-time PCR data. All results have been reported as the mean±standard error of the mean (mean±SEM). The mean difference between groups was also analyzed by one-way ANOVA, followed by Tukey's multiple comparison tests. A p-value less than 0.05 was considered statistically significant.

4. Results

4.1. Biochemical results

The comparison of liver tissue damage indices in the control and SILI groups did not show any significant changes ($P>0.05$). Furthermore, the IR caused a sharp increase in serum AST and ALT values ($P<0.001$). After the injection of silibinin, the levels of these indicators were significantly reduced, compared to the IR group ($P<0.001$) (Table 2).

4.2. Results of liver histopathology

4.2.1. Vehicle group

As shown in Figure 1, the structure of portal space, especially the hepatic artery branch, bile ducts, and the epithelium lining of them are intact and without any damage. In zones I and II of the classic lobules, hepatocytes display the circular and clear nuclei with one or two nucleoli and an intact cytoplasm space full of pink mitochondria. The intact sinusoidal space, Kupffer cells, and immune cells (neutrophils and small lymphocytes) are well identifiable. In zone I of the classic lobules, bile ducts with the simple cuboidal epithelium in the duct wall, and also the cross and longitudinal sections of the

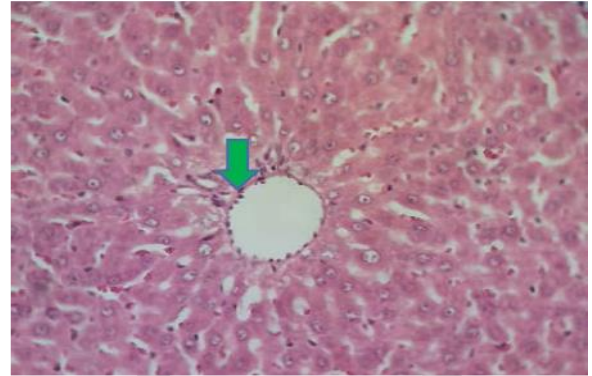
Table 2. Comparison of the markers of liver tissue damage in the four groups

Group	AST	ALT
Vehicle	232.25±21.6	71.89±15.9
IR	2256±442.6***	1384.7±293.8**
SILI+IR	889.13±152.0###	817.38±324.8###
SILI	257.25±29.2	82.51±7.6

The data were expressed by mean±SEM of eight rats per group, and the significance level was considered the values of $P<0.001$.

*** and ### show significant differences, compared to the control group and the IR group, respectively ($P=0.000$).

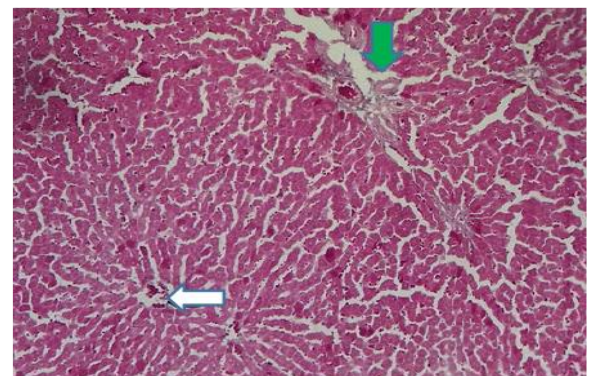
AST: aspartate aminotransferase; ALT: alanine aminotransferase; SILI: Silibinin; IR: ischemia/reperfusion.

**Figure 1.** Histology of normal liver tissue sections with the normal saline injection into rat peritoneal (magnification 400 x)

hepatic artery branch are visible at the margin of the classical lobule and inside the portal space. The hepatic artery endothelium with prominent nuclei in the lumen specifies the health of the artery.

4.2.2. Silibinin group

In Figure 2, the classic hepatic lobule appears healthy in different zones. The components of portal space, including branches of the blood vessel and their walls, as well as the bile ducts appear to be intact. Highly stainability of Remak trabeculae with healthy nuclei, clear nucleoli, and minimal apoptotic vacuoles show the health of the liver tissue in the portal space and the surrounding classical zone I. Normal sinusoidal spaces are visible with intact endothelial lining and a large number of Kupffer cells. In general, the presence of silibinin did not cause any damage to the tissue structure of the rats' liver. In zones I, II, and III, the hepatocyte cords and sinusoidal spaces among them are completely healthy and undamaged.

**Figure 2.** Evaluation of the liver sections treated with silibinin alone (magnification 100x)

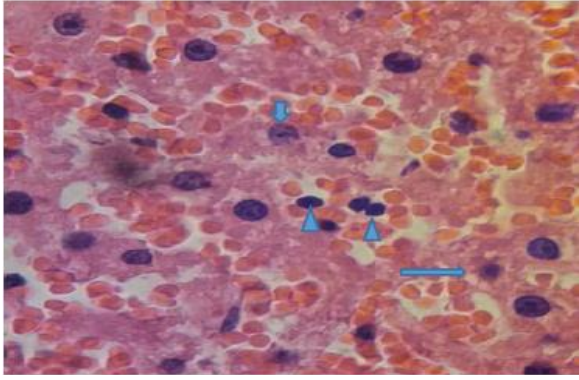


Figure 3. Examination of the liver sections with 1-hour ischemia (magnification 400x)

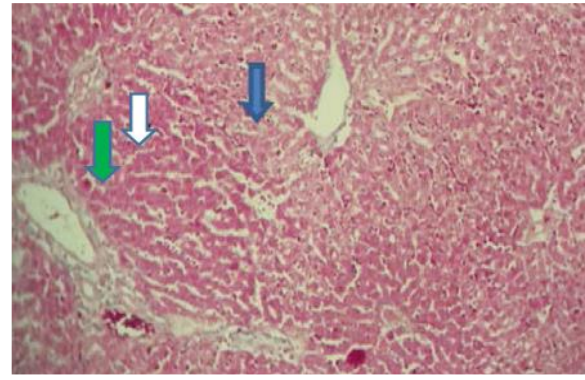


Figure 4. Evaluation of the liver sections with 1-hour ischemic treated with the protective effect of Silibinin (magnification 100x)

4.2.3. Ischemia group

The sinusoidal spaces in zone III are congested (hyperemia and atherosclerosis). In the sinusoidal wall, endothelial cells are severely damaged and desquamated; however, the central vein endothelium appears healthy. Some hepatocytes have become signet ring cells due to the extensive destruction of cytoplasmic organelles and the attachment of these apoptotic vacuoles. Hepatocyte Remak trabecules are visible with clear nuclei, as well as pale and vacuolated cytoplasm. Extensive infiltration of RBCs indicates sinusoidal congestion with extensive endothelial shedding overlying them (Figure 3).

4.2.4. I/R + SILI group

Figure 4D illustrates the silibinin protective effects against IR. Zones I and II of the hepatic lobules have a healthy structure. The healthy hepatocytes with many mitochondria have given a pink to red cytoplasm to the hepatic Remak trabecules. Circular nuclei with clear nucleoli, sinusoids with an intact endothelium, and a low density of red blood cells are observed in the sinusoidal space. The structure of the

hepatic Remak trabecules and their interstitial sinusoids appear to be intact. However, the fading of zone III (blue arrow) surrounding the central vein shows some damage due to decreased mitochondrial density and an increased number of apoptotic vacuoles. The protective effect of silibinin in reducing the extent and severity of injury in these tissue sections is observed, compared to the IR group. Hepatic classic lobules in 100x magnification in zones I and II were marked with green and white arrows, respectively.

4.3. Real-time PCR results

Real-time PCR data analysis demonstrated no statistical differences between the control and SILI groups in terms of the mRNA levels of *Cx43*, *Cx32*, and *Cx26*. Furthermore, the mRNA expression of *Cx32*, *Cx26*, and *Cx43* was significantly enhanced in the IR group, compared to the control group. In the IR+SILI group, silibinin reduced the *Cx26* and *Cx32* mRNA expression, compared to the IR group, whereas it had an inductive effect on the *Cx43* gene expression ($P < 0.001$) (Figure 5).

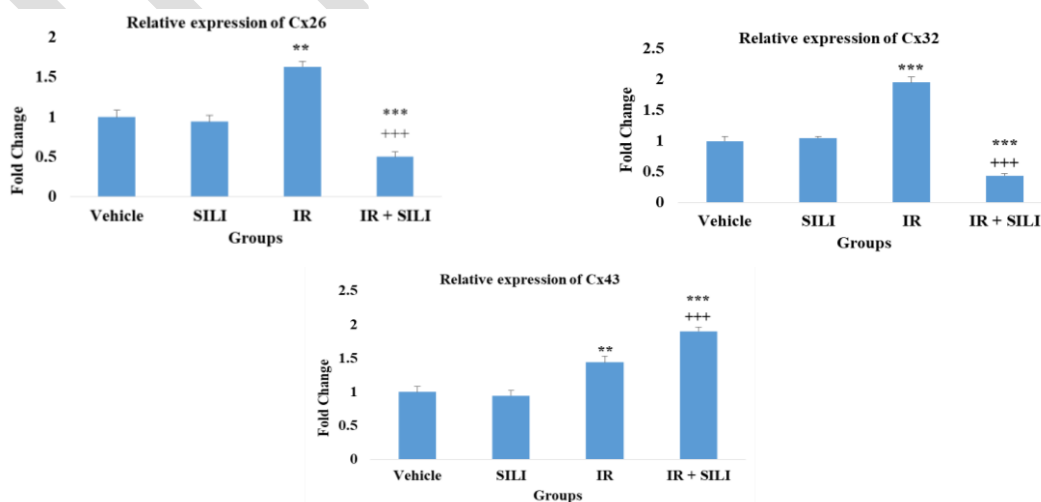


Figure 5. Relative *Cx26*, *Cx32*, and *Cx43* gene expressions among the four studied groups. ** $P < 0.01$ and *** $P < 0.001$, compared to the control group. +++ $P < 0.001$, compared to the IR group, SILI: Silibinin; IR: Ischemia/reperfusion; Cx: connexin

5. Discussion

Research has shown that the kind of Cxs and the change in GJ channels play an important role in the IR damage in the liver tissue. It has been revealed that the transmission rate of IR damage signals is increased through GJs, which causes a chain of damages that lead to changes in cellular metabolism (7). The role of GJ inhibitors in reducing IR damage has been demonstrated in *in vitro* and *in vivo* models (4, 19, 20). Recently, herbal medicines have been considered to reduce the effects of liver ischemia. The positive effects of silibinin on hepatocyte regeneration and IR damage have been proven (21); however, its effects on genes involved in gap junctions have not been studied so far. In the present study, the effect of silibinin was examined on the serum indicators of liver tissue injury, as well as *Cx43*, *Cx32*, and *Cx26* gene expression during warm hepatic IR. The IRI frequently occurs during transplantation and hepatic surgery; moreover, it promotes molecular reactions that enhance damage in the liver parenchyma (22-24). Several studies reported that GJ proteins are involved in liver IRI (7, 25). The GJs are the intercellular channels that mediate the passive diffusion of ions and small molecules, such as calcium, sodium, potassium, ATP, cyclic adenosine monophosphate, glutathione, and inositol triphosphate between the two adjacent cells. The diffusion of these substances via the GJs regulates a large number of physiological processes and tissue homeostasis. The *Cx26*, *Cx32*, and *Cx43* are the predominant GJ proteins detected in various liver cells, including hepatocytes, sinusoidal endothelial cells, and Kupffer cells (5).

Evidence demonstrated that Cxs are involved in liver IRI. Nakashima et al. demonstrated that the *Cx26* and *Cx32* expression levels were down-regulated during liver tissue ischemia; however, they were increased during the early reperfusion phase of the rat model (9). In the same line, Patel et al. reported that increased expression of the *Cx32* gene was associated with liver injury, whereas its inhibition following hepatic reperfusion protected the liver against IRI (7, 25). In line with the previous studies, our results showed that *Cx26*, *Cx32*, and *Cx43* mRNA levels were up-regulated during the hepatic IR. It was also revealed that silibinin administration during IR could reduce endothelial damages, inflammation, and glycogen depletion; in addition, it preserved the mitochondrial membrane, which protected the liver against the IRI (21, 26, 27). It was also determined in this study that silibinin could reduce the expression of *Cx26* and *Cx32*; however, it could induce the expression of the *Cx43* gene. Maes et al. pointed out that an increase in the *Cx43* expression happened upon acetaminophen intoxication converse to the changes in *Cx32* and *Cx26* expression. They elucidated that increased expression of *Cx43* might be due to hepatocyte de novo production in a classical

response to stress and insult.

The above-mentioned study also revealed that *Cx43*-deficient animals were more susceptible to oxidative stress, inflammation, and death, compared to the wild-type animals. Therefore, *Cx43* may have a protective role in acetaminophen toxicity (8). Since the mechanism of liver damages in high-dose acetaminophen intoxication and ischemia-reperfusion is fundamentally similar, it can be concluded that the increased *Cx43*-expression during IR may also play a protective role in this regard.

Hepatic IRI involves several cells in the liver organ, such as hepatocytes, sinusoidal endothelial cells, neutrophils, and Kupffer cells, which trigger multiple molecular pathways, including toll-like receptor (TLR) signaling activation and reactive oxygen species production (28, 29). The TLRs cause neuroinflammation by the triggering of NF- κ B and pro-IL-1 β . Polyphenols, such as silibinin, lignans, flavonoids, phenolic acids, stilbenes, and phenolic alcohols can impede the increase of inflammatory cytokines by TLRs/NF- κ B/STAT (signal transducer and activator of transcription) signaling path. Furthermore, some of the polyphenols can reduce neural apoptosis through the modulating of the TLRs/MyD88/NF- κ B (30).

Ji-Hyeon Song et al. concluded that silibinin decreased the oxidative stress-related proteins and increased the glutathione and catalase expression. Furthermore, it could impede the TLR4-TAK1 signaling cascade and subsequent mediators of inflammation in RAW264.7 macrophage cells triggered by lipopolysaccharide (31). The anti-inflammatory effects of silibinin are mostly due to inhibiting NF- κ B nuclear translocation and activation that results in the inhibition of inflammatory cytokines, such as TNF- α . TNF- α reduced the *Cx43* expression; accordingly, silibinin could indirectly prevent the repressive effect of TNF- α on the *Cx43* gene and increased the expression of the gene (15).

Some histopathological and serum parameters were also investigated in this study to evaluate the impact of silibinin on liver tissue injury. Studies showed that serum levels of ALT and AST were raised during reperfusion with a maximal peak at 12 h after reperfusion (32). The effect of silibinin on ALT and AST levels are consistent with the results of the previous reports. Silibinin can decrease the damage and destruction of hepatocytes, as well as the level of these indices. Furthermore, it could prevent inflammatory cell infiltration, hepatocyte damage, and edema during IR (21). In this study, our histological results confirmed that silibinin declined neutrophil infiltration, endothelial damage, cellular degeneration, and vocalization in IR rats.

6. Conclusion

In this study, silibinin's liver-protective effects

were proved by reducing the expression of specific GJ proteins Cx26 and Cx32, and the induction of Cx43 gene expression in the liver tissue under IR conditions. Furthermore, the histological study of the liver tissue suggested that silibinin dramatically reduced the serum levels of liver enzymes and liver damage. Our data documented the efficacy of silibinin in liver protection against IRI.

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Footnotes

Authors' Contribution: Abbas Khonakdar-Tarsi designed the article. Amin Ghobadi and Akram Mousavi performed laboratory tests. Other authors participated in searching and writing the manuscript. All authors read the manuscript.

Conflict of Interests: The authors have no conflict of interest to declare.

Ethical Approval: The protocol of the experiments was approved by the Ethics Committee of Mazandaran University of Medical Sciences, Sari, Iran (Code: IR.MAZUMS.REC.1398.964).

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