

Effects of ischemic preconditioning in a pig model of large-for-size liver transplantation

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OBJECTIVE: In most cases of pediatric liver transplantation, the clinical scenario of large-for-size transplants can lead to hepatic dysfunction and a decreased blood supply to the liver graft. The objective of the present experimental investigation was to evaluate the effects of ischemic preconditioning on this clinical entity.

METHODS: Eighteen pigs were divided into three groups and underwent liver transplantation: a control group, in which the weights of the donors were similar to those of the recipients, a large-for-size group, and a large-for-size + ischemic preconditioning group. Blood samples were collected from the recipients to evaluate the pH and the sodium, potassium, aspartate aminotransferase and alanine aminotransferase levels. In addition, hepatic tissue was sampled from the recipients for histological evaluation, immunohistochemical analyses to detect hepatocyte apoptosis and proliferation and molecular analyses to evaluate the gene expression of *Bax* (pro-apoptotic), *Bcl-XL* (anti-apoptotic), *c-Fos* and *c-Jun* (immediate-early genes), ischemia-reperfusion-related inflammatory cytokines (*IL-1*, *TNF-alpha* and *IL-6*, which is also a stimulator of hepatocyte regeneration), intracellular adhesion molecule, endothelial nitric oxide synthase (a mediator of the protective effect of ischemic preconditioning) and *TGF-beta* (a pro-fibrogenic cytokine).

RESULTS: All animals developed acidosis. At 1 hour and 3 hours after reperfusion, the animals in the large-for-size and large-for-size + ischemic preconditioning groups had decreased serum levels of Na and increased serum levels of K and aspartate aminotransferase compared with the control group. The molecular analysis revealed higher expression of the *Bax*, *TNF-alpha*, *I-CAM* and *TGF-beta* genes in the large-for-size group compared with the control and large-for-size + ischemic preconditioning groups. Ischemic preconditioning was responsible for an increase in *c-Fos*, *IL-1*, *IL-6* and *e-NOS* gene expression.

CONCLUSION: Ischemia-reperfusion injury in this model of large-for-size liver transplantation could be partially attenuated by ischemic preconditioning.

KEYWORDS: Liver Transplantation; Models; Animal; Reperfusion; Apoptosis; Ischemic Preconditioning.

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INTRODUCTION

With the improvement in the management of pediatric patients with terminal liver diseases and the technical refinement and advances in intensive care and immuno-

suppression, liver transplantation has been performed in increasingly young children (1); a large series of liver transplantations in recipients that were less than 10 or even 5 kg have been reported with satisfactory results (1,2). However, a problem encountered by surgeons in such cases is the graft-to-body weight ratio (GBWR). Classically, the ideal ratio varies from 1 to 3%. However, in small children, the ratio is higher than 5% (3). This situation is known as large-for-size (LFS) and is characterized by the discrepancy between the small size of the abdominal cavity and the large size of the graft, which can lead to hepatic dysfunction and a diminished blood supply to the liver graft (3). Higher incidences of graft dysfunction, vascular complications and

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acute rejection have been reported in the literature in cases involving LFS grafts (3). The mechanisms of injury involved in this scenario are not yet well established.

Among the main mechanisms of graft injury in transplantation is ischemia-reperfusion injury (IRI) (4), which has been related to complications such as poor graft function. Immediate-early genes (IEGs) encode transcription factors that regulate the expression of cellular response genes after injury and are associated with tissue repair and cell apoptosis (5,6). Many studies have shown that *c-Fos* and *c-Jun* are the IEGs that are activated during the early phase after IR and that they play an important role in the regulation of the tissue response to injury stimulation (7,8). Tissue inflammation triggered by IRI occurs immediately after reperfusion and is related to the production of cytokines (TNF-alpha, IL-1, IL-6) and increased expression of adhesion molecules (such as ICAM) (9). The inflammatory cascade that culminates in apoptotic cell death and the cellular mechanisms of regeneration are the elements involved in IRI (10,11,12,13,14). The apoptosis phenomenon results from alterations in the balance between the expression of pro-apoptotic (*Bax* and *Bak*) and anti-apoptotic (*Bcl-2* and *Bcl-XL*) genes. The activation of liver regeneration, which is essential for graft survival, is mainly mediated by IL-6 (15,16,17). After liver transplantation, IRI is responsible for the induction of pro-fibrogenic cytokines, represented mainly by TGF-beta (18).

Some techniques that attenuate IRI, such as ischemic preconditioning (IPC), have been reported and the nitric oxide (NO) produced by endothelial nitric oxide synthase (eNOS) is an important mediator of the effects of IPC. eNOS produces a protective action mediated by vasodilation and platelet aggregation inhibition, thereby improving the microcirculation in addition to inhibiting neutrophil activation and neutralizing reactive oxygen species (18,19,20). Furthermore, it has been demonstrated that IPC stimulates liver regeneration (21). This effect has been demonstrated in rats that underwent warm ischemia and hepatectomy and interleukin-6 (IL-6) has been shown to be a likely mediator (22). No experimental studies have evaluated early hepatic injury in the LFS transplant scenario or the effects of IPC in this situation. Therefore, this was the objective of the present experimental investigation, which utilized a LFS pig experimental model of liver transplantation.

■ MATERIALS AND METHODS

The institution's animal welfare regulatory committee approved all study protocols and all protocols were in compliance with the Guide for the Care and Use of Laboratory Animals published by the National Institutes of Health 86-23, which was revised in 1985. Our study was performed using 36 Landrace pigs, with body weights between 20 and 50 kg. All of the experiments were performed under general anesthesia and all of the animals were killed after the experiments. The operations were performed by the same team of surgeons (the first five authors).

Eighteen large, white Landrace pigs (weighing 17 to 20 kg) underwent OLT with whole liver grafts and were divided randomly into three experimental groups, as follows:

1- Control group (n=6): the weights of the donors were similar to the recipients (17-20 kg);

2- Large-for-size group (LFS, n=6): the weights of the donors were nearly 2 times the weights of the recipients (40-50 kg);

3- Large-for-size + ischemic preconditioning (LFS + IPC, n=6): the weight distributions were similar to the LFS group and IPC was performed in the donors.

Anesthetic procedures and intraoperative monitoring

After fasting for a period of 12 hours, the donors were anesthetized and maintained with a continuous intravenous infusion of 0.005 mg/kg/h fentanyl and 3.5 mg/kg/h propofol.

After an open cut was made down the right carotid sheath, a double lumen catheter was placed into the internal jugular vein for invasive venous pressure monitoring and fluid administration. In addition, a single lumen catheter was inserted into the carotid artery for arterial pressure monitoring.

A total volume of approximately 300 ml of blood was collected from the donors before liver perfusion for infusion into the recipient pigs if necessary.

Donor procedure

In the donor operation, a long, longitudinal, midline thoracic and abdominal incision was performed to widely expose all organs. The portal vein, infrahepatic inferior vena cava and suprahepatic inferior vena cava were dissected. The hepatic artery was maintained in continuity with the celiac trunk and abdominal aorta up to the iliac bifurcation. In situ cold liver perfusion (with Euro Collins® and Ringer's lactate solutions) was then performed through both the portal vein and the aorta. The harvested graft was weighed, immersed in a plastic bag filled with Euro Collins® perfusion solution at 4°C and placed in a basin for the subsequent back-table procedure.

In the donors of the LFS + IPC group, after opening the abdominal cavity, the hepatic hilum was clamped with a vascular clamp. Blood flow was stopped for 15 minutes, followed by reperfusion for 15 minutes.

Recipient procedure

A total hepatectomy was performed via a longitudinal incision and the supraceliac aorta was dissected close to the diaphragmatic crura and clamped during the anhepatic phase. The graft was put in place and the suprahepatic inferior vena cava, infrahepatic inferior vena cava and portal vein were anastomosed. Before the completion of infrahepatic inferior vena cava anastomosis, the graft was flushed with 1 L of saline solution at 4°C through the portal vein.

All anastomoses were performed using continuous Prolene 5-0 or 6-0 sutures. After completion of the venous anastomoses, the liver graft was reperfused. The recipient hepatic artery was anastomosed to the donor celiac trunk patch. The biliary reconstruction consisted of a choledochus-choledochus stented anastomosis.

The abdominal wall was kept open and only a few stitches were placed in the skin to retain the bowels and to avoid hypothermia. After 3 hours, the animal was euthanized.

Blood and serum analyses

Blood from the recipient was sampled at baseline and at 1 and 3 hours after portal reperfusion for the evaluation of the



pH and the sodium and potassium levels. Aspartate aminotransferase (AST) and alanine aminotransferase (ALT) levels were measured in the blood samples collected 1 and 3 hours after reperfusion.

Tissue analyses

Hepatic tissue was sampled from the recipient at 1 and 3 hours after portal reperfusion. Each biopsy sample was divided into two sections. One was preserved in 10% formaldehyde for subsequent embedding in paraffin and the other was snap-frozen in liquid nitrogen at -170°C for molecular analyses.

Histological examination was performed via hematoxylin-eosin staining and light microscopy for the presence of centrilobular necrosis, sinusoidal neutrophils, steatosis, apoptotic bodies and sinusoidal dilatation. For each of these parameters, the following scoring system was used:

0 - absent; 1 - mild; 2 - moderate; 3 - intense; 4 - very intense.

■ IMMUNOHISTOCHEMICAL METHODS

Detection of hepatocyte apoptosis

For the detection of early apoptosis, M30 CytoDEATH, a mouse monoclonal antibody (Clone M30, Roche, Mannheim, Germany), was used to detect a specific epitope of cytokeratin 18 (cleaved cytokeratin 18), which is produced after cleavage by caspase.

The paraffin-embedded liver tissue specimens were sectioned into 5- μm -thick sections. These sections were dewaxed in xylene and hydrated in graded concentrations of alcohol. Endogenous peroxidase activity was blocked with 3% H_2O_2 in methanol for 15 minutes.

The M30 CytoDEATH (Roche) antibody was used at a dilution of 1:100 for 1 hour at room temperature. The sections were then incubated with anti-mouse-Ig Biotin (Chemicon AP302B) at 37°C for 30 minutes, followed by Streptavidin-POD (Roche) for 30 minutes. The location of the antibody was detected using diaminobenzidine, (DAB, Sigma[®]) before the sections were counterstained with hematoxylin, dehydrated with graded alcohol concentrations (70%, 90%, 100%) and HistoClear, and mounted with Entellan (Merck).

Detection of hepatocyte proliferation

To quantify cellular proliferation, immunohistochemical staining for Ki-67 was performed on the liver tissue samples using a rabbit anti-human Ki-67 monoclonal antibody (Clone SP6, Spring Bioscience, Pleasanton, California, USA).

Ki-67 is a nuclear protein that is detected in all active stages of the cell cycle except for G0 (quiescent) and is therefore a marker of cellular proliferation.

After fixation with 10% formaldehyde, the liver tissues were embedded in paraffin, cut into 5- μm -thick sections, deparaffinized and hydrated. The sections were then processed for antigen recovery, endogenous peroxidase blocking and immunohistochemical detection of Ki-67. For that purpose, the sections were incubated with the rabbit anti-human Ki-67 monoclonal antibody (Clone SP6, Spring Bioscience, Pleasanton, California, USA) at a 1:100 dilution. Staining was performed using DAB, followed by counterstaining with Harris hematoxylin, dehydration, and mounting with Entellan (Merck).

The stained tissue sections were examined at magnifications of 150 \times , 600 \times and 1500 \times (under oil immersion) by a

researcher who was blinded to the experimental groups. The stained hepatocyte nuclei were counted at a magnification of 600 \times using an ocular lens containing a 1- cm^2 grid (Nikon). Among all of the nuclei included in the grid area (500 – 800 nuclei), all nuclei stained for Ki-67 and cleaved cytokeratin 18 were counted and expressed as percentages of the total number of nuclei.

Molecular analysis

The expression of the pro-apoptotic *Bax*, anti-apoptotic *Bcl-XL*, IEGs *c-Fos* and *c-Jun*, *IL-1*, *IL-6*, *TNF-alpha*, *TGF-beta*, *ICAM* and *eNOS* genes were evaluated using quantitative reverse-transcriptase polymerase chain reaction (qRT-PCR). In all experiments, the *β -actin* gene was used as a house-keeping control gene.

Total RNA was isolated from all liver samples using the TRIzol reagent (Invitrogen, Carlsbad, CA, USA). Approximately 100.0 mg of tissue was fragmented (Mikro-Dismembrator U; Sartorius AG, Goettingen, Germany) after the addition of liquid nitrogen and homogenization in 1 mL of TRIzol reagent. The RNA was then isolated following standard procedures. Total RNA was quantified by spectrophotometry via a Biophotometer (Eppendorf AG, Hamburg, Germany) at an absorbance of 260 nm and the purity was assessed by measuring the 260/280 nm ratio. This ratio ranged from 1.8 to 2.0 for all samples.

The integrity of the isolated RNA sample was assessed by denaturing agarose gel electrophoresis through the visualization of the 18S and 28S ribosomal RNA bands after ethidium bromide staining. Complementary DNA (cDNA) was prepared from 2.0 μg of total RNA by reverse transcription using 200.0 U of SuperScript III RNase H-RT (Invitrogen) and oligo (dT)s as primers. The resulting cDNA solution was stored at -20°C .

Quantitative real-time PCR was performed in a 15.0- μL reaction mix using 7.5 μL of Platinum SYBER Green qPCR SuperMix-UDG (Invitrogen Carlsbad, CA-USA), 0.3 μL of gene-specific forward and reverse primers (10 μM), 1.0 μL of cDNA, and 5.9 μL of nuclease-free water. The primers used are shown in Table 1.

The cycling conditions were as follows: initial template denaturation at 95°C for 1 minute, followed by 40 cycles of denaturation at 95°C for 20 seconds, annealing at 60°C for 30 seconds, and extension at 72°C for 30 seconds. The reactions were prepared separately for each of the genes and were amplified in a Rotor-Gene RG-3000 (Corbett Research, Australia). The reactions were run in triplicate. Blank controls were included in parallel for each master mix. Amplification was followed by a melting curve analysis to check the PCR product specificity. The data were analyzed by relative quantification and the $2^{-\Delta\Delta\text{Ct}}$ value was calculated (23).

Statistical analysis

Data were stored in Microsoft Excel and transferred to R (www.r-project.org) for statistical analysis and figure creation. The anthropometric and blood analysis data were compared using analysis of variance with a post-hoc Tukey test because the data followed a parametric distribution. The results of the immunohistochemical analyses are shown as the median (minimum and maximum values). The results of the molecular analyses are presented as the 25th and 75th percentiles, the median and the minimum and maximum values in box-plot graphics. The differences between the



Table 1 - Primer sequences.

Gene	Primers
β actin	Sense: 5'-tcc aga ggc gct ctt cca-3' Antisense: 5'-cgc act tca tga tgc agt tga-3'
Bax	Sense: 5'-ctg gga ctt ctt gct gga ttt gc-3' Antisense: 5'-cag ggt gaa gca ggt gaa gat c-3'
Bcl-XL	Sense: 5'-gga gct ggt ggt tga ctt tc-3' Antisense: 5'-gtt tcc tct tct gat tca gtc-3'
eNOS	Sense: 5'-aca tct gca acc aca tca agt acg-3' Antisense: 5'-cac cag ttg gct gtt cca gat c-3'
IL-6	Sense: 5'-cga gcc cac cag gaa cga aag-3' Antisense: 5'-aag cag ccc cag gga gaa ggc-3'
IL-1	Sense: 5'-caa gga cag tgt ggt gat gg-3' Antisense: 5'-cat cat tca gga tgc act gg-3'
TGF-beta	Sense: 5'-cct gga tac caa cta ctg ctt c-3' Antisense: 5'-gtg tct agg ctg cag atg ta g-3'
TNF-alpha	Sense: 5'-cca cca acg ttt tcc tca ct-3' Antisense: 5'-ggc act gag tgc atc atc ct-3'
c-Fos	Sense: 5'-acg gtg act gct atc tgc ac-3' Antisense: 5'-tct tct tct ggg gac aac tg-3'
c-Jun	Sense: 5'-aac ttg aaa gcg cag aac tc-3' Antisense: 5'-aac agt ctg gcc tca aaa cg-3'
ICAM	Sense: 5'-ttc ctt gga tgg act gtt cc-3' Antisense: 5'-cgt ctg cca gca tta tct ca-3'

groups were compared using a nonparametric Kruskal-Wallis test followed by Dunn's multiple comparisons test. Differences with a p value less than 0.05 were considered significant.

RESULTS

The mortality rate was 16.5% (1/6) in all groups. The death in the control group was related to bleeding from a laceration on the graft surface. In the LFS and LFS + IPC groups, the deaths were associated with hemodynamic instability immediately after aortic unclamping.

The anthropometric data and transplant characteristics of the groups are shown in Table 2. The recipient weights, total ischemia times and warm ischemia times were similar between the groups. The donor weights and GBWRs were significantly higher in the LFS and LFS + IPC groups (p=0.005).

Blood analysis: All animals developed acidosis, but there was no difference in the pH values among the groups during the procedure. At 1 hour and 3 hours after reperfusion, the animals in the LFS and LFS + IPC groups had decreased serum levels of Na and increased serum levels of K compared with the control group (Figure 1). The AST levels were markedly higher in the LFS and LFS + IPC groups at both 1 and 3 hours after portal reperfusion

Table 2 - Anthropometric data and transplant characteristics of the groups (mean ± sd).

	Control	LFS	LFS + IPC
Donor weight (kg)	20.3 ± 0.58	44.8 ± 6.21*	48.24 ± 2.89**
Recipient weight (kg)	20.86 ± 1.08	22.2 ± 4.76	20.6 ± 2.79
GBWR	2.9 ± 0.22	6.3 ± 0.73*	6.4 ± 0.98***
Total ischemia time (min.)	138.8 ± 17.8	147.6 ± 8.3	152.0 ± 14.5
Warm ischemia time (min.)	44.8 ± 3.7	46.8 ± 2.7	47.0 ± 3.4

*p=0.005 relative to control.
**p=0.001 relative to control.
***p=0.004 relative to control.

compared with the control group. No alterations were observed in the levels of ALT (Figure 2).

Tissue analysis: Histological alterations were observed in all groups, including marked sinusoidal dilatation and the presence of sinusoidal neutrophils and steatosis (Figure 3). However, there was no difference between the experimental groups regarding these findings.

Immunohistochemical analysis: The numbers of hepatocytes that stained positive for Ki-67, expressed as the percentage of the total number of counted cells, are shown in Table 3. The comparisons between the groups demonstrated that there were no significant differences. Regarding the apoptotic index represented by cleaved cytokeratin-18-positive cells, the statistical analysis showed that there were no differences between the groups.

Molecular analysis: The results of the molecular analyses are shown in Figure 4. Higher Bax gene expression was observed at 3 hours after reperfusion in the LFS group compared with the control and LFS + IPC groups (p=0.004 and p=0.002, respectively). Bcl-XL gene expression did not differ among the groups. c-Fos gene expression was higher in the LFS + IPC group compared with the control and LFS groups (p=0.02 and p=0.004, respectively). This elevated gene expression persisted after 3 hours of reperfusion (p=0.02 compared with the control group). c-Jun gene expression did not differ among the groups and the times after reperfusion.

Regarding cytokines, adhesion molecules and e-NOS, elevated TNF-alpha gene expression was observed in the LFS group compared with the LFS + IPC group (p=0.015 and p=0.043 1 hour and 3 hours after reperfusion, respectively). I-CAM expression was elevated at 3 hours after reperfusion in the LFS group compared with the control and LFS + IPC groups (p=0.026 and p=0.009, respectively). After 1 hour, no differences were observed. IL-1 and e-NOS gene expression was significantly elevated in the LFS + IPC group compared with the control, at both time points after reperfusion (p=0.004 and p=0.002; p=0.003 and p=0.028, respectively). Additionally, e-NOS gene expression was elevated in the LFS + IPC group compared with the LFS group, 3 hours after reperfusion (p=0.017). IL-6 gene expression in the LFS+IPC group was increased 3 hours after reperfusion compared to the control group (p=0.03). Finally, TGF-beta gene expression was elevated in the LFS group compared with the control and LFS + IPC groups at both time points after reperfusion (p=0.009 and p=0.012 after 1 hour; p=0.026 and p=0.021 after 3 hours, respectively).

DISCUSSION

Given the complexity of liver transplantation in children, the difference between the recipient weight and the size of liver grafts that are most commonly harvested from adults may lead to an LFS clinical scenario, which can involve multiple complications.

We previously reported an original LFS experimental model of liver transplantation in large animals with aortic clamping (23). One technical difficulty in porcine liver transplantation is the occasional lethal hemodynamic instability caused by simultaneous portal and caval clamping; to compensate for this, venovenous bypass (VVB) is often used during the anhepatic phase (24). However, VVB substantially increases the morbidity and complexity of the

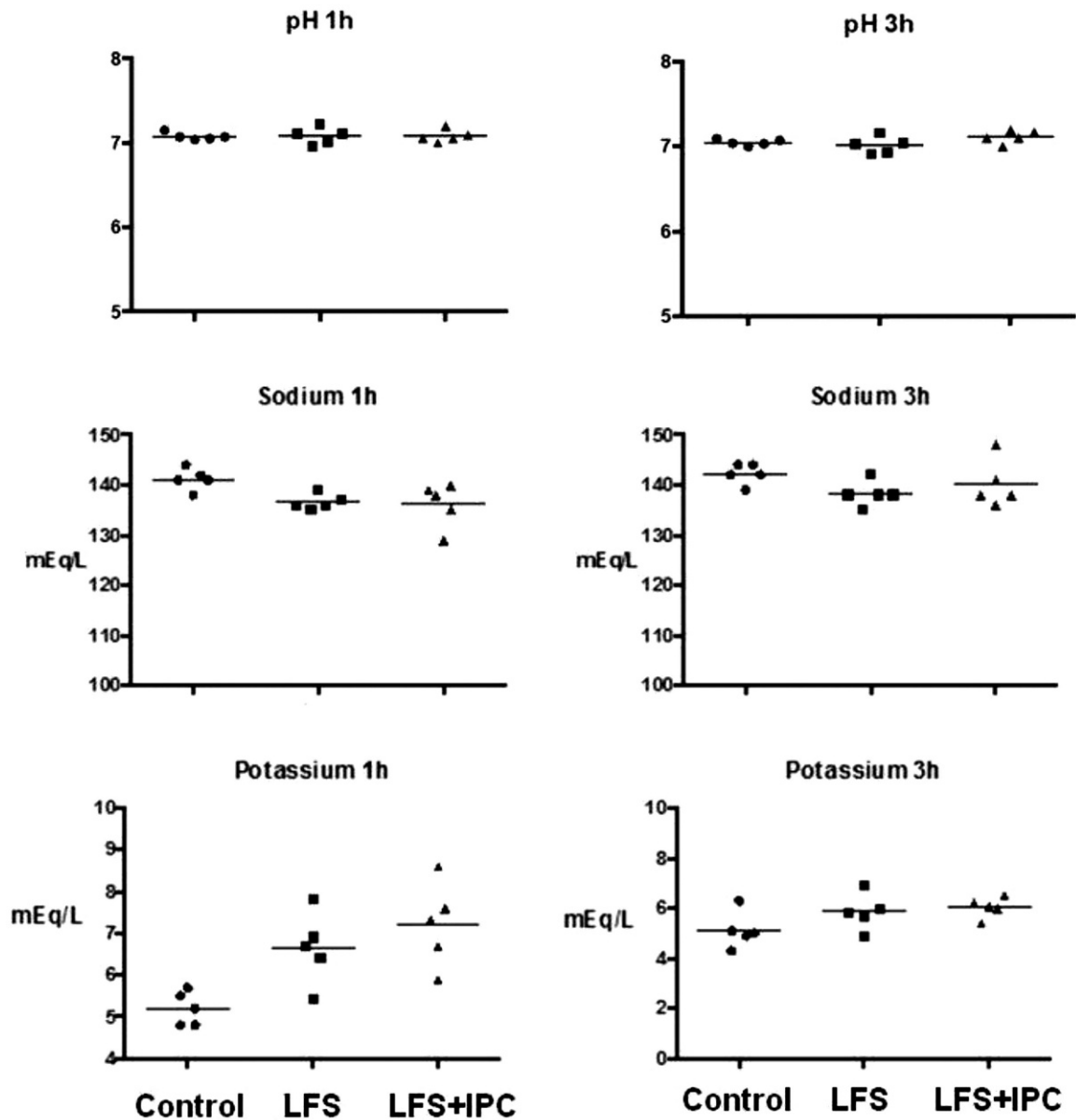


Figure 1 - The results of serum analyses. There were no differences in the pH values among the groups. Decreased serum levels of Na and increased serum levels of K were observed at 1 hour and 3 hours after reperfusion in the LFS and LFS + IPC groups compared with the control group ($p=0.021$ and $p=0.047$ for Na values and $p=0.028$ and $p=0.004$ for K values, respectively).

procedure (25,26). Aortic clamping during the anhepatic phase is well tolerated, results in excellent hemodynamic stability and considerably reduces the operative time because VVB and mesocaval anastomosis construction become unnecessary (27). Although there are potential harmful consequences of aortic clamping, these complications are dependent on the level of aortic cross clamping (clamping of the thoracic aorta produces more complications than clamping of the infrarenal aorta), the duration and patient comorbidities (28). In our experiments using young, healthy pigs and an anhepatic phase lasting no more than 45 minutes, we believe that the complication rate should be low.

Up to 60% of the indications for liver transplantation in children are due to biliary atresia (1), a condition that progresses with hypoplasia or even thrombosis of the portal

vein (29) and these complications lead to an early indication for liver transplantation. The low flow in the portal vein results in difficulty with graft perfusion and this difficulty can be even more pronounced when the graft is relatively large. Using a flowmetry study, we compared portal flow in control and LFS pig liver transplantations. We observed a significant decrease in portal flow from the donor to the recipient in the LFS group, and this decrease was correlated with the higher expression of inflammatory genes related to IRI and progressive elevation of AST levels (30). To simulate the clinical situation in humans, we initially attempted to use pigs weighing approximately 10 kg, but they were still suckling and did not tolerate the procedure; deaths occurred during the anhepatic phase. Therefore, we began to use weaning pigs (weighing approximately 20 kg), which produced better survival indices.

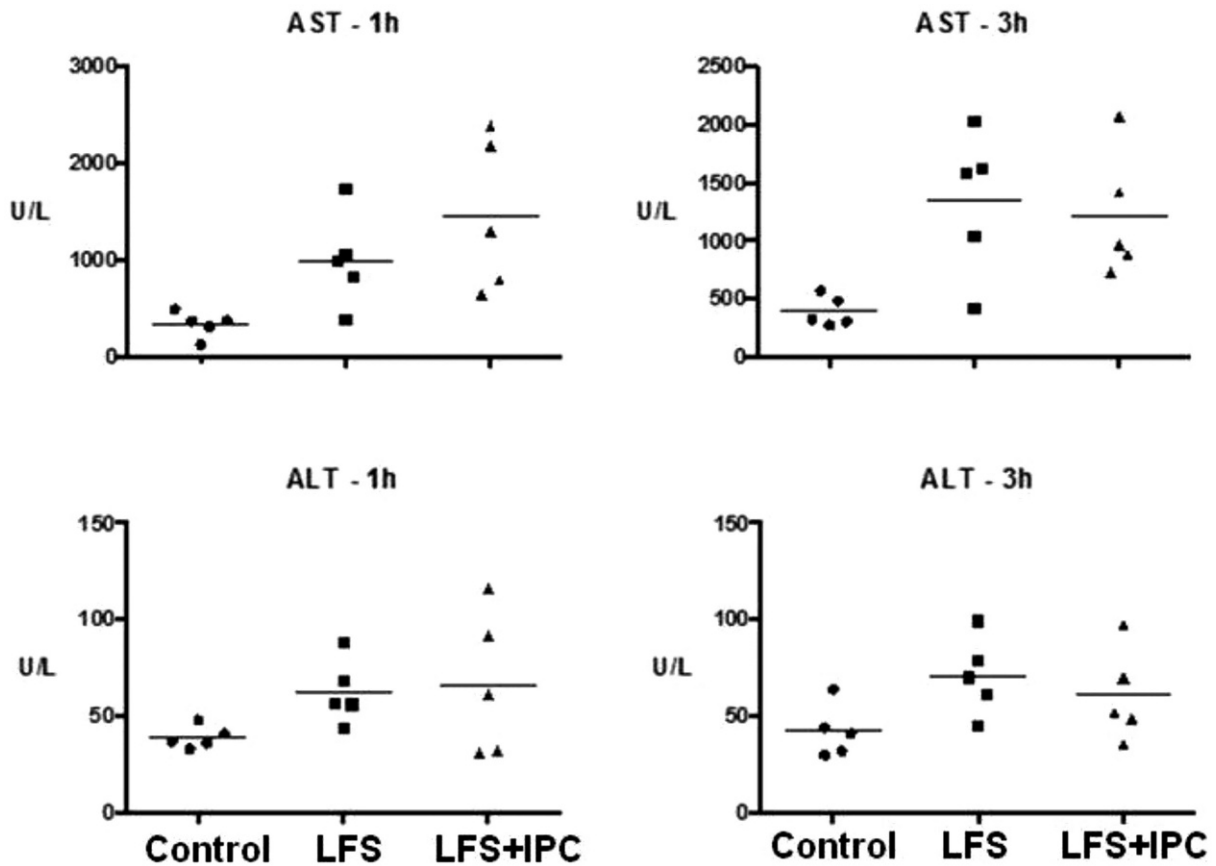


Figure 2 - The AST levels were markedly higher in the LFS and LFS + IPC groups at 1 and 3 hours after portal reperfusion compared with the control group ($p=0.013$ and $p=0.002$, respectively). No alterations were observed in the ALT levels.

In the present study, we aimed to assess the early phase of IRI in LFS transplantation cases. The abdominal cavity was kept open after the experiment in an attempt to avoid the deleterious effects of increased abdominal pressure and inferior vena cava obstruction.

The GBWR was significantly higher in the LFS and LFS + IPC groups (mean 6.3% and 6.4%, respectively). This ratio is consistent with liver transplantation in children. Although the numbers of transplantations and recipients per group were not high, they were similar to many pig experimental

studies in the literature. The complexity of the procedure and the difficulties associated with the maintenance of large animals in urban laboratories justify the low number of animals per group in the current study. However, this number was sufficient to detect many significant differences among the groups. Another point of discussion is the cold ischemia time, which was relatively short compared with liver transplantations with cadaveric donors but prolonged compared with liver transplantations with living donors, which represent the majority of pediatric transplantations in many centers. Although the period of three hours after reperfusion is a relatively short follow-up, we focused our study on the early repercussions of IRI and the expression of genes that are triggered immediately in this process. Thus, the present model is suitable for assessing injury in the LFS

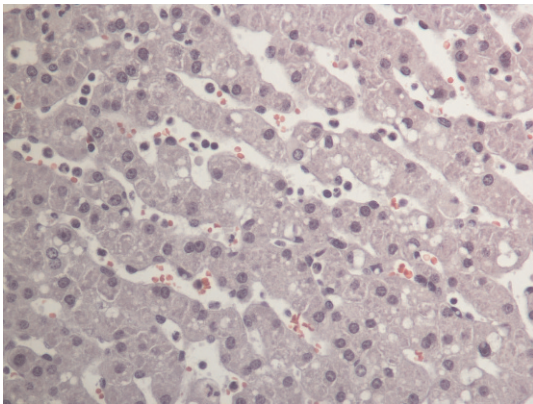


Figure 3 - The histological aspects of a transplanted liver at 3 hours after reperfusion. Note the presence of sinusoidal dilatation, steatosis and sinusoidal neutrophils.

Table 3 - Results of immunohistochemical analyses (median, minimum and maximum values). No differences were observed among the groups.

	Control		LFS*		LFS + IPC*	
	1 h	3 h	1 h	3 h	1 h	3 h
Ki-67	10 (5-10)	15 (8-25)	7.5 (3-15)	20 (6-25)	15 (6-25)	10 (3-15)
Cytokeratin	1 (0-3)	2 (0-4)	2 (1-5)	1 (0-3)	1 (0-4)	2 (0-4)

* $p=0.7$ and $p=0.8$ relative to the control group at 1 h and 3 h, respectively, for Ki-67.

* $p=0.6$ and $p=0.8$ relative to the control group at 1 h and 3 h, respectively, for cytokeratin.

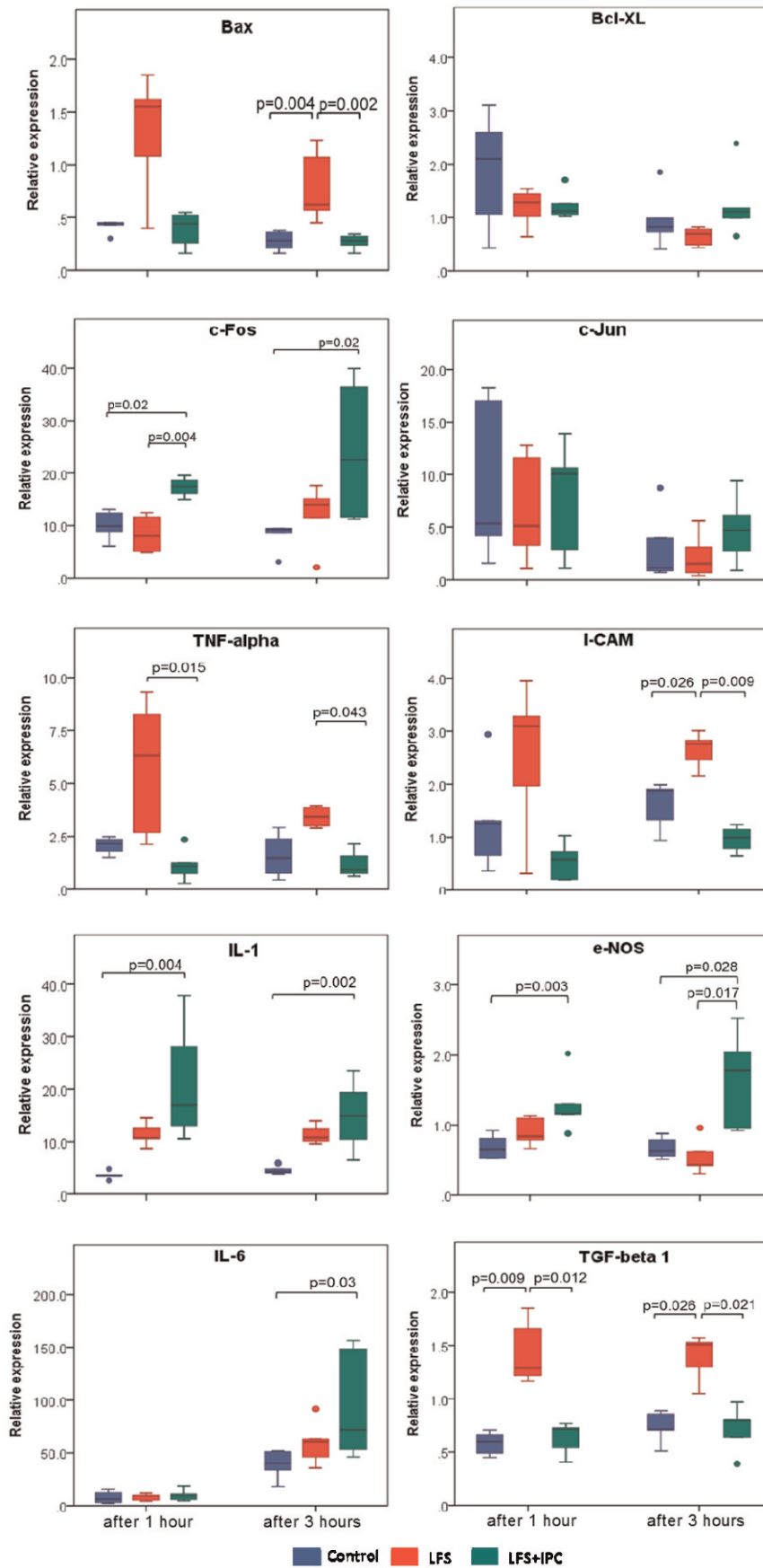


Figure 4 - The results of molecular analyses. Note the positive effects of IPC on gene expression.



scenario and we aimed to evaluate acute graft lesions in this context.

The standardization of the IPC interval (15 minutes of portal clamping followed by 15 minutes of reperfusion) was based on a literature review. Schulz et al. demonstrated in a study conducted in pigs that IPC with 10-minute intervals had no protective effect on IRI (31), whereas the utilization of longer IPC times (15 minutes) produced significant effects (32). Similar to the clinical scenario, all animals developed acidosis that was not related to the size of the graft because there were no differences in the pH values between the groups. The acidosis was likely related to conditions of hypoperfusion caused by the clamping of the inferior vena cava and the portal vein during the anhepatic phase in addition to the clamping of the aorta.

Hyperkalemia was observed in all groups, especially in the first hour after reperfusion. This condition is also similar to that observed in clinical transplantation. The liver graft, which is stored in a cold preservation solution, is the main source of potassium in this situation. During reperfusion, a large quantity of potassium is released from the graft into the bloodstream (33,34). Higher values were observed in the LFS and LFS + IPC groups and as described above, this can be attributed to the higher mass of the transplanted liver graft. After 3 hours, the serum potassium values in the LFS and LFS + IPC groups decreased and were comparable to the control group values.

Intraoperative natremia disturbances are less often discussed in the literature. The sodium values were lower in the LFS and LFS + IPS groups one hour after reperfusion. During the period of ischemia, cellular swelling occurs because of the influx of sodium and water into the intracellular space (35). As the liver mass was higher in the LFS and LFS + IPC groups, this result may have been caused by a greater accumulation of intracellular sodium in these groups.

The pathophysiology of hepatic ischemia/reperfusion injury (IRI) includes a number of complex and diverse mechanisms involving the interactions between hepatocytes, Kupffer cells, neutrophils, macrophages, sinusoidal endothelial cells and platelets. IRI is one of the main causes of graft dysfunction after transplantation and contributes to decreased graft and patient survival (14-16). Therefore, the identification of the factors that contribute to increased tissue damage, as well as measures to attenuate this damage, is critical to improving survival. In the current investigation, we aimed to study whether IPC has positive effects on the graft in the experimental LFS liver transplantation model. IPC consists of an adaptive mechanism in which brief episodes of ischemia protect an organ from injury caused by prolonged ischemia. IPC contributes to improving hepatic microcirculation and regeneration, thereby attenuating IRI. Biochemical, histological and molecular biology analyses were used in this study to evaluate the effects of IRI and IPC in the LFS liver transplantation model.

The serum AST levels were higher in the LFS and LFS + IPC groups during both time periods studied. This change highlights the potential deleterious effect of implanting a disproportionately large graft into a recipient. The AST levels in the LFS group increased from the first to the third hour after reperfusion; however, in the LFS + IPC group, the AST levels tended to decrease from the first to the third hour. Similarly, immunohistochemical analyses of cellular

proliferation revealed that IPC was responsible for a tendency towards an increased peak in Ki-67 expression. These facts may suggest that IPC provokes an earlier but less harmful response to IRI.

The histological analysis revealed no differences between the groups. This result is most likely explained by the fact that the study only analyzed the acute phase of the experiment. Nevertheless, it is important to note that some typical IRI-related changes, such as the presence of sinusoidal dilatation, sinusoidal neutrophils and hepatic steatosis, were observed.

Apoptosis was evaluated based on *Bax* (pro-apoptotic) and *Bcl-XL* (anti-apoptotic) gene expression. *Bax* gene expression was significantly higher in the LFS group compared with the control and LFS + IPC groups at 3 hours after reperfusion. This finding supports the hypothesis that apoptosis due to IRI is exacerbated in LFS scenarios. In addition, IPC plays a protective role by decreasing *Bax* gene expression.

Regarding the anti-apoptotic effects of IPC, *Bcl-XL* gene expression did not differ between the studied groups, but there was a tendency for higher *Bcl-XL* expression in the LFS + IPC group at 3 hours after reperfusion. The difference between the LFS and LFS + IPC groups almost reached statistical significance ($p=0.05$), demonstrating that IPC likely plays an anti-apoptotic role in IRI.

The co-existence of apoptosis and hepatocyte regeneration in IRI shows that liver tissue remodeling is regulated by both processes. IEGs are important factors in the cellular proliferation induced by IRI as a part of tissue repair (9,10). Cells lose the ability to respond to proliferation signals by the blockade of Fos protein; thus, c-Fos is considered an important factor in the cell regeneration signal transduction pathway (36). Some studies suggest that the apoptosis/proliferation processes are doubly regulated by different *c-Fos* and *c-Jun* mRNA expression levels. *c-Fos/c-Jun* heterodimers are thought to promote cell regeneration and *c-Jun* homodimers may lead to cell apoptosis (10,11). In fact, we observed that *c-Jun* levels were similar in all groups; however, IPC was responsible for increasing *c-Fos* levels. This result may be one explanation for the protective effect of IPC against hepatocyte apoptosis.

Tumor necrosis factor- α (TNF- α), a cytokine produced by numerous cell types in response to inflammatory stimuli, is a well-known mediator during ischemia/reperfusion (IR) that plays a central role in hepatocyte injury. TNF- α has a multifactorial effect in hepatic IRI because of the many interactions between TNF- α and reactive oxygen species, nitric oxide, adhesion molecules, and various cytokines and chemokines (37). Zhang et al., studying the action of mast cell stabilization in acute lung injury after autologous orthotopic liver transplantation in rats, observed an upregulation of the serial pro-inflammatory cytokines TNF- α , IL-1-beta and IL-6, peaking at 4 hour after reperfusion (38). In our study, transplantation of a disproportionately large graft was responsible for increased TNF- α expression compared with the control group; this may be related to increased ICAM expression, suggesting more intense IRI in the LFS situation. Czigány et al., studying the effects of remote ischemic preconditioning in liver IRI in rats, observed that preconditioning was responsible for a protective effect on liver morphology and was associated with a decrease in TNF- α serum levels (39). Analogously, ischemic preconditioning was responsible for decreased TNF- α levels as well



as decreased ICAM expression in the LFS situation, thereby attenuating IRI.

Interestingly, in our study, IPC led to an increase in IL-1 expression. Shin et al., analyzing IPC-induced neuroprotection, observed that IPC was responsible for a parallel increase in IL-1 β and IL-1 receptor antagonist expression in the ischemic cortex, resulting in a significant reduction in the brain cortex infarct area (40). Unfortunately, we did not measure IL-1 receptor antagonist expression, but it is reasonable to assume that in the LFS+IPC group, higher IL-1 receptor antagonist expression would also be observed.

Along these lines, *eNOS* gene expression was significantly higher in the LFS + IPC group than in the other groups at 3 hours after reperfusion. IPC resulted in increased *eNOS* expression. This enzyme is responsible for NO production, which in turn contributes to improving microcirculation, thereby reducing tissue injury (41). Liver regeneration is closely related to the expression of IL-6, which is released in IRI by Kupffer cells. Previous studies have demonstrated the role of IL-6 in liver regeneration and the increased release of IL-6 promoted by IPC (20,42). At 1 hour after reperfusion, there was no difference between the groups. However, after 3 hours, IL-6 gene expression was higher in the LFS + IPC group. This finding suggests that IL-6, along with other factors, may play an important role in the protective mechanisms of IPC in attenuating IRI.

A relationship between IRI and the stimulation of fibrogenesis has been suggested and studied in the literature. TGF- β is one of the main cytokines related to liver fibrogenesis and has been related to the transdifferentiation of stellate cells into myofibroblasts and to the deposition of portal collagen. Yu et al., using a model of selective portal vein ligation, aimed to determine whether IR in clamped lobes facilitated liver regeneration of non-clamped lobes (43). In fact, the authors observed that IR pretreatment of the clamped lobes facilitated the liver regeneration of non-clamped lobes after selective portal vein ligation, which may have resulted from down-regulated TGF- β 1 expression in the non-clamped lobes. Our results suggest that the more intense IRI in the large graft stimulated TGF- β production and that IPC is responsible for the attenuation of these liver fibrogenesis stimuli.

In summary, IRI is more pronounced in this model of LFS liver transplantation and is partially attenuated by IPC, mainly through the improvement of the gene expression of early molecular markers. The long-term implications and the graft and animal survival were not examined in this study because this research was conducted to assess the acute phase of IRI. Future studies to evaluate the long-term outcomes, liver perfusion and techniques to modify liver perfusion are important to establish a better understanding of the biological processes involved in this scenario.

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■ AUTHOR CONTRIBUTIONS

Leal AJ, Tannuri AC, Belon AR and Guimarães RR performed the experiments. Coelho MC, Gonçalves JO, Serafini S and Melo ES

performed the laboratory analyses. Tannuri U performed the final revision of the manuscript.

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