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Author manuscript *Am J Transplant*. Author manuscript; available in PMC 2019 May 01.

Published in final edited form as:

Am J Transplant. 2018 May ; 18(5): 1110–1121. doi:10.1111/ajt.14586.

## Heme Oxygenase-1 Regulates Sirtuin-1 – Autophagy Pathway in Liver Transplantation: From Mouse-to-Human

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## Abstract

Liver ischemia-reperfusion injury (IRI) represents a major risk factor of early graft dysfunction and a key obstacle to expanding the donor pool in orthotopic liver transplantation (OLT). Although graft autophagy is essential for resistance against hepatic IRI, its significance in clinical OLT remains unknown. Despite recent data identifying heme oxygenase-1 (HO-1) as a putative autophagy inducer, its role in OLT and interactions with Sirtuin-1 (SIRT1), a key autophagy regulator, have not been studied. We aimed to examine HO-1 mediated autophagy induction in human OLT and in a murine OLT model with extended (20h) cold storage, as well as to analyze the requirement for SIRT1 in autophagy regulation by HO-1. Fifty-one hepatic biopsies from OLT patients were collected under an IRB protocol two hours after portal reperfusion, followed by Western blot analyses. High HO-1 levels correlated with well-preserved hepatocellular function and enhanced SIRT1/LC3B expression. In mice, HO-1 overexpression by genetically-modified HO-1 macrophage therapy, was accompanied by decreased OLT damage, increased SIRT1/LC3B expression, whereas adjunctive inhibition of SIRT1 signaling diminished HO-1 mediated hepatoprotection and autophagy induction. Our translational study confirms the clinical relevance of HO-1 cytoprotection and identifies SIRT1-mediated autophagy pathway as a new essential regulator of HO-1 function in IR-stressed OLT.

Disclosure

The authors of this manuscript have no conflicts of interest to disclose as described by the American Journal of Transplantation.

**Supporting Information** Supplementary Materials

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The authors declare that no conflict of interest exists.

## Introduction

Orthotopic liver transplantation (OLT) is the standard of care for patients with end-stage liver disease and those with hepatic malignancies, but liver graft shortage remains a major challenge. Liver ischemia-reperfusion injury (IRI), an innate immune-driven sterile inflammation response leading to hepatocellular death, is an inevitable consequence of multiple clinical conditions, including trauma, sepsis, hepatic tumor resection, and liver transplantation. Indeed, hepatic IRI has been recognized as a major risk factor for delayed early graft function, acute and chronic rejection as well as a key obstacle to expanding the donor organ pool. However, despite obvious clinical importance, mechanisms that account for liver IRI are only partially understood and no effective therapy is available to prevent or treat this condition in humans (1).

Autophagy is an evolutionarily conserved intracellular self-digesting pathway responsible for maintaining energy homeostasis and removing long-lived or damaged organelles and proteins (2). A growing body of evidence indicates a tissue protective function for autophagy in various pathologic states, such as aging, diabetes and neurodegenerative diseases (3) as well as in liver IRI (4-6). Although enhanced autophagy may represent a novel therapeutic target against IR-damage, its efficacy and mechanism in OLT settings remain to be assessed, while clinical relevance of autophagy pathway in liver transplant patients has not been studied before.

Heme oxygenase-1 (HO-1; *Hmox1*), a rate-limiting enzyme that catalyzes the conversion of heme into biliverdin, carbon monoxide, and free iron, exerts potent anti-oxidative, antiinflammatory and cytoprotective functions (7). We and others have reported on beneficial function of HO-1 in hepatic IRI murine models using chemical HO-1 inducers (8-11), adenovirus (Ad) gene transfer (8, 12) and genetically-modified animals (13). As macrophages are key mediators of innate immune-driven inflammation and the primary source of HO-1 in IR-stressed liver (10, 14), we have established a novel molecular transfer approach using *ex-vivo* genetically-modified HO-1-overexpressing macrophages. This regimen effectively transferred the target molecule into 40-50% of hepatic cells and successfully alleviated warm IRI in mice (15, 16); its putative efficacy in clinically-relevant OLT models has not been tested. Although anti-inflammatory phenotype is central to HO-1 function in liver IRI (17), recent studies have identified HO-1 as a novel hepatic autophagy inducer (18, 19); its role in liver transplantation remains unknown and putative regulatory mechanisms need to be studied.

Sirtuin 1 (SIRT1), an NAD+-dependent type III histone/protein deacetylase involved in cellular senescence, inflammation and stress resistance (20), plays a key role in autophagy induction (21, 22). In addition to an anti-inflammatory role in pathogenesis of liver IRI (23), recent studies have demonstrated the significance of SIRT1-induced autophagy in hepatocyte resistance against IR-stress (24, 25). Molecular communication between SIRT1 and HO-1 in OLT has not been studied and the mechanism by which SIRT1 may regulate the HO-1 autophagy pathway remains to be elucidated.

To gain further insight into HO-1 mediated autophagy in liver transplantation, we used an *ex-vivo* genetically-modified HO-1 macrophage adoptive transfer approach in a clinically relevant murine OLT model and analyzed human OLT samples (n=51) in parallel. Our translational study confirms the clinical relevance of HO-1 hepatoprotection and identifies SIRT1-dependent autophagy as a novel and essential regulator of HO-1 function in OLT under IR-stress.

### Materials and Methods

### **Clinical liver transplant study**

Fifty-six consecutive adult primary OLT recipients were recruited under an IRB protocol (13-000143; May 2013 - August 2015). Routine standard of care and immunosuppressive therapy were administered as specified by UCLA liver transplant protocols. Study data were collected and managed using REDCap electronic data capture tools (26). Donor organs, procured using standardized techniques, were perfused with and stored in cold University of Wisconsin (UW) solution (ViaSpan; Bristol-Meyers Squibb Pharma). Cold ischemia time was defined as the time between the perfusion of the donor liver with UW solution and its removal from cold storage. Recipient venous blood was collected within the hour prior to the transplant and on post-operative days 1-14 (POD 1-14). Hepatocellular injury was evaluated by serum alanine aminotransferase (sALT) levels. Post-transplant biopsies (Bx) were obtained from the left liver lobe using a Tru-Cut biopsy needle approximately two hours after portal reperfusion (prior to the abdominal closure) and snap-frozen. Out of fifty-six cases, fifty-one were examined; four were excluded because the Bx samples were too small for Western blot analyses, and one was excluded due to unavailability of clinical data. Early allograft dysfunction (EAD) was defined by the presence of one or more of the following: total bilirubin 10 mg/dL (171 µmol/L) or INR 1.6 on POD 7, and ALT/AST >2,000 IU/L within the first 7 POD (27) (28). Post-transplant rejection was diagnosed by follow-up Bx done per clinical standard of care.

#### Animals

C57BL/6 mice at 6-8 weeks of age were used (Jackson Laboratory, Bar Harbor, ME). Animals were housed in the UCLA animal facility under specific pathogen-free conditions, received humane care according to the criteria outlined in the "Guide for the Care and Use of Laboratory Animals" (NIH publication 86-23 revised 1985). All studies were reviewed and approved by the UCLA Animal Research Committee.

#### Generation of genetically modified bone marrow-derived macrophages (BMM)

L929 cells (ATCC, Rockville, MD) were cultured in RPMI-1640 medium supplemented with 2 mmol/l l-glutamine, 100 U/ml penicillin, 100 µg/ml streptomycin, and 10% fetal calf serum. The conditioned media was collected from cells grown for 7 days. The BMM were generated according to standard procedures (15). In brief, bone marrow cells were removed from the femurs and tibias of C57BL/6 mice and cultured in Dulbecco's modified Eagle's medium supplemented with 10% fetal calf serum, 100 U/ml penicillin, 100 µg/ml streptomycin, and 2 mmol/l l-glutamine, with the addition of 15% L929-conditioned

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medium. Cells were cultured at  $5 \times 10^6$  cells/well for 7 days with 5% CO<sub>2</sub> and 95% air at 37 °C, and used for in vitro transfection.

Replication-defective recombinant adenovirus encoding HO-1 (AdHO-1) was generated as described (29). BMM ( $5 \times 10^{6}$ /well) were incubated for 1h with AdHO-1 or Ad $\beta$ -gal (at a MOI of 10). After medium change, cells were cultured for additional 48h prior to *in vivo* adoptive transfer. By 48h of transfection the X-Gal staining was >90%, compared with control BMM (15).

#### Mouse liver cold ischemia and transplantation model

We used a well-established mouse model of *ex-vivo* hepatic cold storage and orthotopic liver transplantation (OLT) as described (30). Donor livers were stored in UW solution at 4°C for 20h prior to transplantation into syngeneic mice. BMM ( $5 \times 10^6$ ) transfected with AdHO-1/Adβ-gal ( $2.5 \times 10^9$  pfu) or untreated ( $5 \times 10^6$ ) were injected via portal vein prior to liver reperfusion at the completion of transplant surgery. In some experiments, SIRT1 inhibitor (EX527, Sigma Aldrich, St. Louis, MO) was administered (10mg/kg i.p.) at 30 min prior to OLT surgery (15, 16). Liver and serum samples were collected 6h after reperfusion, the peak of hepatocellular damage in this model. The sham group underwent the same procedures except for OLT.

#### Serum biochemistry

Serum alanine transaminase (sALT) levels, an indicator of hepatocellular injury, were measured by IDEXX Laboratories (Westbrook, ME).

#### Liver histology and IRI grading

Formalin-fixed paraffin-embedded OLT sections  $(5\mu m)$  were stained with hematoxylin and eosin (HE). The severity of hepatic IRI was graded using Suzuki criteria (31).

**Quantitative RT-PCR analysis**—RNA was extracted from liver tissue samples using RNAse Mini Kit (Qiagen, Germantown, MD). A total of 5.0µg of RNA was reverse-transcribed into cDNA. Quantitative PCR was performed using DNA Engine with Chromo 4 Detector (MJ Research, Waltham, MA) (32). The primers sequences are listed (Table S1). The expression of the target gene was normalized to the housekeeping HPRT.

**Western blot assay**—Proteins were extracted from liver tissues and their concentration was measured (BCA Protein Assay Kit, Thermo Scientific). Equal amount of protein was electrophoresed, blotted, and incubated with primary Ab, secondary HRP-conjugated Ab, and developed. Primary Ab detecting HO-1 (Enzo Life Sciences, Farmingdale, NY), SIRT1, LC3B,  $\beta$ -actin (Cell Signaling Technology, Danvers, MA) were used. To compare target protein expression in multiple human OLT samples, densitometry quantification was conducted using a reference sample and normalization with  $\beta$ -actin as reported (23).

#### Immunohistochemistry

OLT-infiltrating neutrophils were detected using monoclonal rat anti-Ly6G Ab (BD Biosciences, San Jose, CA). Immunostaining signals were visualized with a labeled polymer

in the EnVision + system horseradish peroxidase kit (Dako). Positively-stained cells were counted blindly (10 HPF/section). LC3B expression in OLT was detected using rabbit anti-LC3B Ab and signals were visualized with secondary Ab: Alexa Fluor 488 anti-rabbit IgG.

#### TdT-mediated dUTP nick end labeling (TUNEL) assay

Cell death in formalin-fixed paraffin-embedded liver sections (5µm) was detected by Apop Tag Plus Peroxidase in Situ Apoptosis Kit (Millipore, Temecula, CA). Results were scored semi-quantitatively by blindly counting the number of positive cells in 10 HPF/section.

**Statistical Analysis**—In mouse experiments, group comparisons were performed using a Student *t*-test. For human data, continuous values were analyzed by Mann-Whitney U test and categorical variables by Fisher's exact test. Spearman's correlation coefficient (r) was used to evaluate the strength of linear relationship between variables. The cumulative survival rate was analyzed by Kaplan-Meier method, and differences between groups were compared using a log-rank test. JMP for Windows 8.0 (SAS Institute, Cary, NC) was used for statistical analyses. A *p*-value of <0.05 was considered statistically significant.

### Results

#### AdHO-1 transfected BMM alleviate IRI in OLT

We first aimed to validate the efficacy of HO-1 induction following the adoptive transfer of AdHO-1 BMM in a clinically relevant mouse IRI-OLT model. Cold-stored liver grafts were infused with BMM via portal vein immediately prior to reperfusion. At 6h after reperfusion, AdHO-1 BMM-treated OLT showed increased HO-1 expression as compared with naive or Adβ-gal transfected BMM-treated controls (Fig. 1A). AdHO-1 BMM-conditioned OLT displayed attenuated sinusoidal congestion, edema/vacuolization and hepatocellular necrosis as compared with control OLT infused with untreated or Adβ-gal transfected cells (Fig. 1B). These correlated with depressed Suzuki grading of histological liver damage (OLT: + AdHO-1 BMM =  $1.7\pm0.6$  vs. OLT: + BMM =  $3.7\pm1.2$  or IR + Adβ-gal =  $3.3\pm0.6$ ; n=4-6/group, p<0.05, Fig. 1C) and preserved liver function, evidenced by sALT levels (OLT: +AdHO-1 BMM =  $3.900\pm2.642$  vs. OLT: +BMM =  $7.997\pm2.068$  or IR + Adβ-gal =  $7.726\pm2.129$  IU/L; mean±SD, n=4-6/group, p<0.05, Fig. 1C) in AdHO-1 BMM-treated OLT. Thus, infusion of AdHO-1 BMM into the graft enhanced HO-1 expression and minimized the adverse effects of hepatic IRI in OLT.

#### AdHO-1 gene transfer promotes anti-inflammatory signature in OLT

As anti-inflammatory function is one of the key HO-1 biological features (17), we examined whether AdHO-1 gene transfer alters the anti-/pro-inflammatory gene expression profile in IR-stressed OLT. Indeed, AdHO-1 BMM infused directly into the graft increased mRNA coding for anti-inflammatory IL10 and TGF $\beta$ , while decreasing pro-inflammatory TNF $\alpha$ , MCP1 and CXCL10 as compared with naïve/Ad $\beta$ -gal BMM treated controls (Fig. 2A/B). This confirms HO-1 overexpression exerts an anti-inflammatory function and inhibits inflammatory cytokine-induced OLT injury. We also tested whether AdHO-1 gene transfer may affect cytokine-induced neutrophil OLT influx. As expected, HO-1 overexpression reduced neutrophil (Ly6G) numbers in IR-stressed OLT infused with AdHO-1 BMM (Fig.

3A/B), indicating HO-1 gene transfer enhanced the anti-inflammatory phenotype and diminished cytokine-induced inflammation.

# AdHO-1 gene transfer attenuates hepatocellular death while enhancing SIRT1/LC3B expression in OLT

We next aimed to determine whether AdHO-1 gene transfer affects cell death and autophagic activity in IR-stressed OLT. As shown in Fig. 3A/B, a reduced frequency of TUNEL+ apoptotic/necrotic cell death was observed in AdHO-1 BMM-treated OLT as compared with naive BMM or Adβ-gal BMM controls. Moreover, we found that AdHO-1 BMM therapy not only upregulated hepatic SIRT1 (a key autophagy regulator) but also enhanced LC3B (an autophagosome marker) expression, as compared with controls (Fig. 4A-C), underscoring the importance of the HO-1 – SIRT1 axis in cell autophagy regulation. Taking together, these results are consistent with the idea that HO-1 overexpression exerts cytoprotective and autophagic functions in IR-stressed OLT.

#### HO-1 expression correlates with IR-liver damage and SIRT1/LC3B signaling in human OLT

Having shown AdHO-1 BMM-facilitated hepatoprotection accompanied by increased SIRT1/LC3B levels in a murine OLT model, we next aimed to validate the relevance of these findings in human liver transplantation. Liver post-reperfusion biopsies (Bx) from fifty-one human OLTs were screened by Western blots for HO-1, SIRT1 and LC3B expression (Fig. 5A). Hepatic HO-1 expression profile exhibited weak negative correlation with sALT levels at POD1 (r=-0.3306, p=0.0178, Fig. 5B), indicating higher HO-1 was associated with milder IR-liver injury. In addition, graft HO-1 expression showed positive correlation with SIRT1 (r=0.4229, p=0.0020, Fig. 5C) and weak positive correlation with LC3B (r=0.3594, p=0.0096, Fig. 5D).

To evaluate the impact of hepatic HO-1 on clinical outcomes, fifty-one OLT samples were split evenly into low (n=26) and high (n=25) HO-1 expression groups (Fig. 6A). The patients' demographic data and clinical parameters are shown (Table S2). There was no correlation between HO-1 expression on and donor background, including age, gender, weight, BMI or pre-procurement sALT level. We also found no correlation between HO-1 expression and recipient factors, including age, gender, weight, BMI, race, disease etiology, presence of HCC, ABO-compatibility, MELD score, pre-transplant sALT or dialysis. Furthermore, there was no correlation between HO-1 expression and cold ischemic time or intra-operative blood loss.

Consistent with Fig. 5C/D data, the high HO-1 group showed increased SIRT1 (low HO-1:  $0.83\pm0.10$  vs high HO-1:  $1.25\pm0.10$ , mean $\pm$ SEM, p<0.05, Fig. 6B left) and LC3B (low HO-1:  $0.87\pm0.08$  vs high HO-1:  $1.13\pm0.10$ , mean $\pm$ SEM, p<0.05, Fig. 6B right) expression. Representative Western blots are shown (Fig. 6C). In addition, the high HO-1 group exhibited significantly lower sALT levels at POD1 (low HO-1:  $809\pm367$  vs high HO-1:  $255\pm34$  IU/L, mean $\pm$ SEM, p<0.05, Fig. 6D). To examine the relationship between HO-1 expression and OLT outcomes, we analyzed cumulative post-transplant survival, with the median follow-up of 740 days (range, 4-1432 days). None of the patients underwent secondary liver transplantation. Despite consistent trends, the improved survival in the high

HO-1 expression group failed to reach statistical significance when compared with the low HO-1 expression group (p=0.1963, Fig. 6E). The high HO-1 expression group was characterized by lower frequency of early graft dysfunction (3.8% vs 8.0%) and post-OLT rejection episodes (3.8% vs 16.0%). However, these differences failed to reach statistical significance (p=0.5150 and p=0.1871, respectively). Representative TUNEL and LC3B staining in low vs high HO-1 Bx groups are shown (Fig. 6F).

## SIRT1 inhibition abrogates hepatoprotection and depresses autophagy despite HO-1 induction in mouse OLT

Based on the aforementioned preclinical and clinical data, preservation of hepatocellular function (sALT) and enhanced autophagy (LC3B) in HO-1 overexpressing OLT was accompanied by increased SIRT1 signaling, a putative regulator of autophagy and stress resistance against hepatic IR-damage (25). Hence, we hypothesized SIRT1 might be essential in cytoprotection after HO-1 induction in IR-stressed OLT. As shown in Fig. 7A, adjunctive SIRT1 inhibition prior to transplantation (EX527, 10mg/kg i.p.) recreated cardinal features of hepatic IRI in otherwise well-functioning AdHO-1 BMM-treated OLT, evidenced by increased sALT levels and Suzuki histological grading scores (Fig. 7B). Moreover, SIRT1 inhibition depressed LC3B expression in AdHO-1 BMM-treated liver grafts. These data indicate the SIRT1 – autophagy pathway is required for HO-1 mediated hepatoprotection against IRI in OLT.

## Discussion

This study is the first to document HO-1 mediated autophagy enhancement in human liver transplant settings and in a clinically relevant mouse OLT model. Although antiinflammatory signaling is critical for HO-1 function against IR-stress, our findings document autophagy regulation as a novel mechanistic component of its biological effects. Our clinical study showed positive correlation between HO-1 and SIRT1 levels in human OLT (Fig. 5C/D and Fig. 6B/C), while SIRT1 inhibition in parallel mouse studies diminished autophagy phenotype seen otherwise after HO-1 induction in IR-resistant OLT (Fig. 4A/B and Fig. 7C/D). These data are consistent with the notion that HO-1 regulates hepatic autophagy in a SIRT1 dependent manner. Although previous studies suggested the role of p38 MAPK in HO-1 induced liver autophagy (18, 33), it remains unknown as to how HO-1 may regulate hepatic p38 signaling. In the context of our present data, a recent finding of liver-specific SIRT1 increasing p38 translocation/phosphorylation via YAP/MKK3 dependent manner (34) implies SIRT1-p38 signaling may underlie the autophagy regulation by HO-1. These complex molecular mechanisms require further in-depth analyses.

It is well documented that hepatocyte autophagy is essential in liver stress resistance (4, 35, 36). Indeed, the loss of Atg4B increased sensitivity of aged mouse livers to warm IRI (5), whereas rapamycin increased autophagy and protected against IR-stress (6). In contrast, recent studies in myeloid-specific Atg5 knockout mice have shed a new light on macrophage autophagy regulation (37-39). While macrophage Atg5 deficiency was shown to limit acute toxic liver injury by downregulating IL1 $\beta$  (38), the impaired macrophage Atg5 promoted proinflammatory macrophage polarization in obese mice (37). Although macrophages are

pivotal in innate-immune driven IR-damage, the significance of macrophage autophagy in liver IRI, using myeloid autophagy-deficient animals, awaits future research.

Despite numerous preclinical reports identifying HO-1 as a major cytoprotective molecule (10, 15, 16), Geuken reported high "pre-transplant" HO-1 expression in human donor livers paradoxically correlated with augmented post-transplant injury (40). As "post-transplant" HO-1 features specific to clinical liver transplantation have not been well defined, we have recently reported that high HO-1 expression in human OLT (n=21) at the time of reperfusion are associated with suppressed IR-liver damage (41). In agreement with the latter, our present data from human OLT recipients (n=51) consistently demonstrated negative correlation between HO-1 levels and post-transplant sALT levels (Fig. 5B and Fig. 6D), implying therapeutic benefit of HO-1 induction in clinical liver transplantation. Unlike hepatocytes under basal conditions, liver macrophages (infiltrating and tissue-resident) are the major HO-1 producers, both in human OLT and IR-stressed murine livers (10, 41).

Although aging and obesity are poor-prognostic factors influencing HO-1 expression, we found no correlation between post-transplant HO-1 levels and donor/recipient demographic parameters or surgical factors (Supp. Table 2). As HO-1 is not only a cytoprotective molecule but also a stress-inducible heat shock protein-32 (hsp32), its expression increases with the severity of IR-stress. Paradoxically, high HO-1 levels were associated with a more attenuated liver damage in the clinical arm of our study (Fig. 6D). We speculate that the divergent HO-1 expression in OLT might be caused by variable individual inductivity of HO-1 in response to IR-stress, resulting in variable susceptibility against IRI. Indeed, short guanine-thymine nucleotide repeats polymorphism in the promotor region on chromosome 22q13.1 of human HO-1 was shown to enhance HO-1 expression in response to oxidative stress (42), whereas a more recent study associated HO-1 polymorphism with post-OLT hepatocellular damage (43). As others failed to correlate HO-1 polymorphism with its expression profile (40), more studies on putative factors influencing human HO-1 responses are warranted.

We have also recently reported on the anti-inflammatory and hepatoprotective role of SIRT1 activation in a warm liver IRI mouse model alongside negative correlation between SIRT1 and T-bet/IL-1 $\beta$  levels in twenty-one clinical OLT cases (23). The positive SIRT1 regulation by HO-1 seen in a mouse IRI model was then confirmed in twenty-one OLT patient cohort (41). The current study documents the functional significance of HO-1 - SIRT1 axis in a clinically-relevant mouse hepatic cold ischemia/OLT model (Fig. 4A/B) and in fifty-one human OLT cases (Fig. 5C and 6B). Although in our previous clinical study (n=21), post-OLT "high" HO-1 expression was accompanied by a statistically significant improvement in OLT survival (41), we failed to detect a similar statistically significant survival differences in the current (n=51) liver transplant patient cohort (Fig, 6E, p=0.1963). As discussed earlier (41), the link between graft and recipient long-term outcomes and IRI severity remains controversial. The presence of early lesions on the time-zero biopsy in older grafts (n=16)did not translate into early graft dysfunction or graft loss (44), while histological IRI grading in post-reperfusion liver biopsies and peak transaminases (n=55) did not show any correlation with clinical outcomes in the first post-transplant month (45). Likewise, despite preferable trends, our present study (n=51) failed to detect significant differences in EAD,

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post-OLT rejection or overall survival. In marked contrast, severe IRI on the time-zero biopsy and peak of post-transplant ALT in 476 OLT cases were significantly associated with a 1-year graft survival in an earlier study (46). The correlation between graft/recipient outcomes and IRI severity may thus require considerably larger number of study subjects.

A chemical HO-1 inducer (COPP; cobalt protoporphyrin) was recently shown to increase SIRT1 expression in hepatocytes, while adjunctive competitive HO-1 inhibitor (SnMP; tin mesoporphyrin) abrogated SIRT1 upregulation, indicating HO-1 dependent SIRT1 induction by exogenous COPP (47, 48). Likewise, in our current study, transfer of genetically modified HO-1-overexpressing BMM consistently enhanced liver graft SIRT1 expression levels (Fig. 4A/B). The mechanisms accounting for HO-1 mediated SIRT1 upregulation remain largely unknown. As oxidative stress suppresses hepatic SIRT1 (23, 49), anti-oxidant properties may be the key for HO-1 to regulate SIRT1 expression.

Adenovirus is an efficient vector for *in vivo* gene therapy due to its ability to mediate transgene expression in various cell types. However, serious off-target effects, including activation of innate immune responses, thrombocytopenia, coagulopathy, and liver damage restrict systemic use of Ad vector in clinical practice (50). In our present and previous preclinical studies (15, 16), we used replication-defective recombinant Ad to generate HO-1-overexpressing BMM. Indeed, the *ex-vivo* gene modified BMM local infusion upregulated hepatic HO-1/LC3B expression and alleviated IRI in a mouse OLT model. This regimen makes Ad-related adverse effects negligible (51), while direct BMM infusion into the liver improves macrophage transfer efficacy as a cellular delivery system for HO-1 trafficking into the disease site (52). The present study using *ex-vivo* gene modified HO-1-overexpressing BMM in a clinically relevant mouse model is likely to validate and support future experiments in translational and clinical settings.

In our current study, the injection of BMM (Ad $\beta$ -gal) to OLT increased hepatic HO-1 expression as compared with untreated BMM (no virus) treated graft (Fig. 1A). This result is consistent with our previous findings in a mouse model of liver warm IRI (16). As a foreign pathogen, virus infection itself can cause cellular stress to activate antioxidant system, accompanied by increased antioxidant gene expression, such as manganese superoxide dismutase, HO-1, indoleamine-2, 3-dioxygenase, and glutathione peroxidase (53). We speculate that increased HO-1 expression following Ad $\beta$ -gal BMM injection was induced by virus-related BMM stress, which was accompanied by increased SIRT1 and slightly higher LC3B levels (Fig. 4A). However, Ad $\beta$ -gal BMM failed to ameliorate liver damage in IRstressed OLT (Fig. 1B/C) as "pure" viral-mediated HO-1 induction may not be proficient enough to protect the liver graft. In contrast, unlike the Ad $\beta$ -gal BMM controls, infusion of Ad HO-1 BMM markedly enhanced local HO-1 (Fig. 1A) and SIRT1/LC3B (Fig. 4A) levels leading to attenuated hepatocellular injury in IR-stressed OLT (Fig. 1B/C).

In conclusion, our study with genetically modified HO-1-overexpressing BMM transfer in a mouse OLT model and parallel clinical examination of fifty-one human liver transplants not only confirmed the clinical relevance of HO-1 hepatoprotection but also identified a novel regulatory role of SIRT1 mediated autophagy as a part of cytoprotective function in transplant recipients. The present data, in agreement with our recent reports (15, 16),

validate a novel investigative tool in which host macrophages can be transfected *ex-vivo* with cytoprotective HO-1 and then infused, if needed, into prospective recipients to mitigate IR-mediated inflammation during liver transplantation, resection or trauma. As decreasing donor organ quality represents one of the most challenging problems, our findings provide the rationale for a new and clinically attractive gene targeted strategy to "rejuvenate" extended criteria donor livers and improve their function via HO-1–SIRT1–autophagy pathway in transplant recipients.

### Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

### Acknowledgments

This work was supported by NIH grants: R01 DK102110; R01 DK107533; and R01 DK062357 (JWKW); NIH PO1 AI120944 and Keck Foundation Award 986722 (JWKW and EFR); and Ruth L. Kirschstein National Research Service Award T32CA009120 (RAS). We thank Ko Takanashi at UCLA-TPCL for immunohistochemical assistance.

## Abbreviations

Ad	adenovirus
ALT	alanine aminotransferase
BMM	bone marrow derived macrophages
Bx	biopsy
HE	hematoxylin and eosin
НО-1	heme oxygenase-1
IRI	ischemia-reperfusion injury
OLT	orthotopic liver transplantation
POD	post-operative day
sALT	serum alanine aminotransferase
SIRT1	sirtuin 1
TUNEL	TdT-mediated dUTP nick end labeling

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#### Figure 1. Adoptive transfer of AdHO-1 BMM ameliorates IR-damage in OLT

Mouse (C57BL/6) livers subjected to 20h of cold storage were transplanted to syngeneic mice. BMM (5×10<sup>6</sup>) transfected with AdHO-1 or Adβ-gal ( $2.5 \times 10^9$  pfu) or untreated BMM were infused into liver grafts via portal vein immediately prior to reperfusion. Liver grafts and serum samples were analyzed at 6h post-OLT. (**A**) Representative (n=3) Western blot-assisted detection of HO-1; β-actin expression served as an internal control. (**B**) Representative H&E staining (n=3; original magnification, ×100). (**C**) Suzuki histological grading of liver IRI and sALT levels. (n=4-6/group); \*p<0.05 vs. OLT: +BMM (AdHO-1). Data shown as mean±SD.

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## Figure 2. Adoptive transfer of AdHO-1 BMM inhibits pro-inflammatory signature and enhances anti-inflammatory program in OLT

Mouse (C57BL/6) livers subjected to 20h of cold storage and transplanted to syngeneic mice were infused immediately before reperfusion via portal vein with BMM (naïve, AdHO-1 or Ad  $\beta$ -gal). Samples were analyzed at 6h post-OLT. qRT-PCR-assisted detection of mRNA coding for IL10 and TGF $\beta$  (A), TNF $\alpha$ , MCP1, CXCL10 (B). Data were normalized to HPRT gene expression (n=4/group) \* p<0.05 vs. OLT: +BMM (AdHO-1). Data shown as mean±SD.

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## Figure 3. Adoptive transfer of AdHO-1 BMM suppresses neutrophil trafficking and mitigates hepatocellular death in OLT

Mouse (C57BL/6) livers subjected to 20h of cold storage and transplanted to syngeneic mice were infused via portal vein with BMM (naïve, AdHO-1 or Ad  $\beta$ -gal). Samples were analyzed at 6h post-OLT. (**A**) Representative immunohistochemical Ly6G (upper panels) and TUNEL (lower panels) staining. (**B**) Quantification of Ly6G (left) and TUNEL (right) positive cells/HPF (n=4-6/group). \* p<0.05 vs. OLT: +BMM (AdHO-1). Data shown as mean±SD.

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Figure 4. Adoptive transfer of AdHO-1 BMM enhances SIRT1/LC3B expression in OLT

Mouse (C57BL/6) livers subjected to 20h of cold storage and transplanted to syngeneic mice were infused via portal vein with BMM (naïve, AdHO-1 or Ad  $\beta$ -gal). Samples were analyzed at 6h post-OLT. (**A**) Representative (n=3) Western blot-asisted detection of SIRT1 and LC3B;  $\beta$ -actin expression served as an internal control. (**B**) qRT-PCR-assisted detection of mRNA coding for HO-1 and SIRT1. Data were normalized to HPRT gene expression (n=4/group) \* p<0.05 vs. OLT: +BMM (AdHO-1). Data shown as mean±SD. (**C**) Representative (n=3) immunohistochemical detection of LC3B in OLT at 6h postreperfusion.

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Figure 5. Post-transplant HO-1 expression negatively correlates with hepatocellular damage and positively with SIRT1/LC3B expression in human OLT

(A) Post-transplant liver biopsies (Bx; 2h post-reperfusion) were collected from fifty-one OLT patients. Bx samples were analyzed by Western blots with  $\beta$ -actin normalization. (B) Relationship between post-transplant HO-1 expression and sALT levels at postoperative day 1 (POD1). (C/D) Relationship between post-transplant HO-1 and SIRT1 (C) and LC3B (D) expression. r: Spearman's correlation coefficient.

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## Figure 6. Increased post-transplant HO-1 levels correlate with attenuated hepatocellular damage and enhanced SIRT1 – LC3B pathway in human OLT

(A) Bx samples collected from the fifty-one OLT patients were classified into low HO-1 (n=26) and high HO-1 (n=25) groups. (B) Western blot-assisted expression of SIRT1 and LC3B. Data shown in dot plots and bars indicate mean±SEM. # p<0.05 (Mann-Whitney U test). (C) Representative Western-blots assisted detection of HO-1, SIRT1 and LC3B (Case A/B: low HO-1 group, Case C/D: high HO-1 group) (D) sALT levels at postoperative day 1-14 (POD1-14) shown as mean±SEM. Dotted line indicates low HO-1, while the solid line high HO-1. # p<0.05 (Mann-Whitney U test). (E) The cumulative probability of OLT survival (Kaplan-Meier method). Dotted line indicates low HO-1, while the solid line high HO-1 groups (log-rank test). (F) Representative TUNEL and LC3B staining pattern in OLT.

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## Figure 7. SIRT1 inhibition depresses LC3B expression and abrogates hepatoprotection in AdHO-1 BMM-conditioned IR-OLT

Mouse (C57BL/6) livers subjected to 20h of cold storage were transplanted to syngeneic mice and conditioned with AdHO-1 BMM with or without adjunctive SIRT1 inhibition (EX527, 10mg/kg i.p). Samples were analyzed at 6h post-OLT. (**A**) Representative H&E staining (original magnification, ×100). (**B**) sALT levels (left) and Suzuki histological grading of liver IRI (right). \*p<0.05 vs. OLT: + AdHO-1 BMM + DMSO (n=3-4/group). Data shown as mean±SD. (**C**) Representative (n=3) Western blot-asisted detection of LC3B;  $\beta$ -actin expression served as an internal control. (**D**) Representative (n=3) immunhistochemical detection of LC3B.