

Cold Storage Followed by Transplantation Induces Immunoproteasome in Rat Kidney Allografts: Inhibition of Immunoproteasome Does Not Improve Function

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Abstract:	Background It is a major clinical challenge to ensure the long-term function of transplanted kidneys. Specifically, the injury associated with cold storage of kidneys compromises the long-term function of the grafts after transplantation. Therefore, the molecular mechanisms underlying cold-storage-related kidney injury are attractive therapeutic targets to prevent injury and improve long-term graft function. Previously, we found that constitutive proteasome function was compromised in rat kidneys after cold storage followed by transplantation. Here, we evaluated the role of the immunoproteasome (iproteasome), a proteasome variant, during cold storage (CS) followed by transplantation. Methods

	Established in vivo rat kidney transplant model with or without CS containing vehicle or iproteasome inhibitor (ONX 0914) was used in this study. The iproteasome function was performed using rat kidney homogenates and fluorescent-based peptide substrate specific to $\beta 5i$ subunit. Western blotting and quantitative RT-PCR were used to assess the subunit expression/level of the iproteasome ($\beta 5i$) subunit. Results We demonstrated a decrease in the abundance of the $\beta 5i$ subunit of the iproteasome in kidneys during CS, but $\beta 5i$ levels increased in kidneys after CS and transplant. Despite the increase in $\beta 5i$ levels and its peptidase activity within kidneys, inhibiting $\beta 5i$ during CS did not improve graft function after transplantation. Summary These results suggest that the pharmacological inhibition of immunoproteasome function during CS does not improve graft function or outcome. In light of these findings, future studies targeting immunoproteasomes during both CS and transplantation may define the role of immunoproteasomes on short- and long-term kidney transplant outcomes.	
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Cold Storage Followed by Transplantation Induces Immunoproteasome in Rat Kidney Allografts: Inhibition of Immunoproteasome Does Not Improve Function

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Keywords: kidney transplantation; cold storage; immunoproteasome, interferon-gamma

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Key Points

- Cold storage increases the severity of graft dysfunction in a time dependent manner, and prolonged cold storage decreases animal survival
- Cold storage plus transplant increases iproteasome levels/assembly in renal allografts; IFN-g is a potential inducer of the iproteasome
- Inhibiting iproteasome ex vivo during renal cold storage did not confer graft protection after transplantation

Abstract

Background: It is a major clinical challenge to ensure the long-term function of transplanted kidneys. Specifically, the injury associated with cold storage of kidneys compromises the long-term function of the grafts after transplantation. Therefore, the molecular mechanisms underlying cold-storage-related kidney injury are attractive therapeutic targets to prevent injury and improve long-term graft function. Previously, we found that constitutive proteasome function was compromised in rat kidneys after cold storage followed by transplantation. Here, we evaluated the role of the immunoproteasome (*iproteasome*), a proteasome variant, during cold storage (CS) followed by transplantation.

Methods: Established in vivo rat kidney transplant model with or without CS containing vehicle or iproteasome inhibitor (ONX 0914) was used in this study. The *iproteasome* function was performed using rat kidney homogenates and fluorescent-based peptide substrate specific to $\beta 5i$ subunit. Western blotting and quantitative RT-PCR were used to assess the subunit expression/level of the *iproteasome* ($\beta 5i$) subunit.

Results: We demonstrated a decrease in the abundance of the $\beta 5i$ subunit of the *iproteasome* in kidneys during CS, but $\beta 5i$ levels increased in kidneys after CS and transplant. Despite the increase in $\beta 5i$ levels and its peptidase activity within kidneys, inhibiting $\beta 5i$ during CS did not improve graft function after transplantation.

Summary: These results suggest that the pharmacological inhibition of immunoproteasome function during CS does not improve graft function or outcome. In light of these findings, future studies targeting immunoproteasomes during both CS and transplantation may define the role of immunoproteasomes on short- and long-term kidney transplant outcomes.

Introduction

It continues to be a challenge to optimize long-term graft function, especially when the transplanted kidneys come from deceased donors (up to 70%)¹⁻⁷. These donor kidneys are flushed with and stored in cold storage (CS) solutions (static or machine perfusion)⁸⁻¹⁰, until the recipient is identified. While it is well-known that CS is detrimental to graft function after transplantation¹¹⁻¹⁵, the mechanisms of CS-mediated renal injury are largely unknown.

Ischemia-reperfusion injury (IRI) is an unavoidable consequence of CS and transplantation, and the IRI is known to induce numerous pathophysiological pathways leading to cell stress and organ damage¹⁶. Following stress, the ubiquitin proteasome system (UPS) and mitochondria play a critical role in maintaining protein homeostasis and energy metabolism. The constitutive proteasome, which is part of the UPS, has 3 to 7 protease active sites (β -catalytic subunits - β 1, β 2, and β 5) and it selectively degrades modified or damaged proteins to small peptides¹⁷⁻¹⁹. Similarly, the mitochondrial electron transport chain produces energy (ATP), and reactive oxygen species (ROS) are a byproduct. Previously, we showed that CS-induced mitochondrial oxidative stress and impaired mitochondrial function contributed to kidney injury during preservation in both rat and pig models²⁰⁻²². Our follow-up study with a clinically relevant model of CS followed by transplantation (CS+Tx) demonstrated decreased peptidase activity of the β 5 subunit of the constitutive proteasome, increased mitochondrial dysfunction, and severe renal damage/dysfunction after CS+Tx compared to transplantation without CS²³⁻²⁵. These results clearly indicate that extended CS negatively affects kidney function in our experimental model of CS+Tx.

The immunoproteasome (*i*proteasome) is a proteasome variant normally found in immune cell compartments^{26, 27}. In response to inflammation, constitutive proteasome subunits (β 1, β 2, and β 5) are exchanged for *i*proteasome subunits (β 1i/LMP2, β 2i/LMP10, and β 5i/LMP7) in most non-immune cells^{28, 29}. ROS activate the *i*proteasome, which, unlike the constitutive

proteasome, is resistant to oxidative stress and can function without ATP^{26, 27, 30}. Recent studies indicate that inhibiting the *i*proteasome with ONX 0914—a specific, reversible inhibitor of the $\beta 5i$ subunit³¹—protects against cardiac and neuronal ischemia–reperfusion injury^{32, 33}. It is not known whether CS mediates activation of *i*proteasome in kidney grafts, and if so, what role $\beta 5i$ plays in CS-mediated renal damage and dysfunction. Here, we hypothesized that CS+Tx increases $\beta 5i$ peptidase activity in the *i*proteasome in renal grafts; we predicted that the addition of ONX 0914 during CS would blunt $\beta 5i$ function, leading to improved graft function after transplantation. Here, we used rat models of kidney transplantation and evaluated *i*proteasome activity in the kidney grafts. Similarly, we evaluated inhibition of $\beta 5i$ activity during renal CS and evaluate the effects on allograft function after transplantation.

Materials and Methods

Animals

Lewis or Fischer rats were obtained from Charles River Laboratory. All animal experiments were performed according to the animal use protocol approved by the Institutional Animal Care and Use Committee, and in compliance with institutional/NIH guidelines.

Rat Surgery

Rat kidney transplant surgery was performed as described previously^{23-25, 32-36}. Postoperatively, rats were given saline solution (5-10 ml, subcutaneously [SC]) to replace lost fluids and buprenorphine (2 mg/kg, SC) to alleviate pain.

Cold storage followed by transplant (CS+Tx) surgery: For the syngeneic model, Lewis rats were used as donors and recipients. For the allogeneic model, Fischer rats were used as donors, and Lewis rats were used as recipients. The right and left kidneys from the donor rat were flushed with and stored in University of Wisconsin (UW) solution, referred to here as CS solution, at 4°C

for 4 h or 18 h. For *iproteasome* inhibition studies, the donor kidney was flushed with and stored in CS solution containing ONX 0914 (β 5i inhibitor, 100 nM) or DMSO (vehicle control). In the recipient rat, the native left kidney was removed, and the donor left kidney (from 4- or 18-h CS groups) was orthotopically implanted with end-to-end anastomosis (CS+Tx or CS+ONX 0914+Tx group, n=4). For 1-day surgery groups and 7-day survival surgery groups (both syngeneic), the native right kidney was removed at the time of the transplant so that renal function/rat survival depended entirely on the transplanted left kidney. For the 9-day surgery group (allogeneic), the native right kidney was removed on day 7 post-surgery, and the rats survived for an additional 2 days to assess the function of the transplanted (left) kidney.

Autotransplant (ATx) surgery: ATx surgery was included in these studies so that the effect of CS could be isolated from the effect of transplant surgery alone. ATx (n=4) was performed as described for the CS+Tx group above, except that the left kidney was removed, flushed with saline, and immediately transplanted back into the same rat without CS exposure; a right nephrectomy followed immediately.

Sham surgery: Rats underwent the same procedure for right nephrectomy but without renal transplantation (n=4). The right kidney was saved immediately as an untreated control kidney. The left kidney remained in the rat, which lived for the same number of days as CS+Tx or ATx animals.

Tissue Sample Collection

For all groups, kidneys and blood were collected under anesthesia 1 or 9 days post-surgery, and animals were euthanized. Kidneys were immediately processed with formalin for histology or flash-frozen and saved at -80°C . Isolated blood was collected in a tube with (plasma) or without heparin (serum), as appropriate. The blood was then centrifuged (5,000 g) at 4°C for 5 min, and the supernatant was saved at -80°C . For survival surgery, blood from the

same rats was collected 1, 3, and 7 days after surgery in a tube containing heparin and used for renal function assays.

In Vitro Cell Model

Normal rat kidney proximal tubular cell cultures (NRK-52E, ATCC) were maintained in warm growth medium (DMEM plus 5% FCS and 1% penicillin/streptomycin). NRK cells at 70% confluence were treated with growth medium containing IFN- γ (10 ng/ml, R&D Systems) or TNF- α (10 ng/ml, R&D Systems) at 37°C for 1 day^{35, 36}. Control NRK cells were treated with vehicle only. After treatment, NRK cells were washed 2 times with cold PBS (4°C) and used to prepare RIPA lysates^{23, 35, 36}.

Renal Function and Serum Chemistry

Blood chemistry was determined in heparinized blood (arterial) using an iSTATTM hand-held clinical chemistry analyzer and CHEM8⁺ cartridges as described by the manufacturer (Vetscan®, Abaxis, USA)^{23-25, 32-36}.

Histology and Immunohistochemistry

Paraffin sections from sham, CS+Tx, and CS+ONX +Tx kidneys (9 days post-surgery) were assessed for tubular injury using periodic acid-Schiff reaction as described previously²³. Masson's Trichome staining was performed to assess renal fibrosis. Histological sections were evaluated in a blinded fashion. All parameters culminating in a tubular injury score were in the scale of 0-5 as described previously²³. KIM-1 (Novus Bio, 1:800) and NGAL (Novus Bio, 1:500) immunohistochemistry on renal sections were performed as described previously³⁷.

Immunoproteasome Function

Peptidase activity of the $\beta 5i$ subunit of the immunoproteasome was measured in renal tissue homogenate by hydrolysis of the fluorogenic peptide substrate ²³, specific for $\beta 5i$ /PSMB8 Acetyl-Ala-Asn-Trp-AMC (100 μ M) (South Bay Bio) in the presence or absence of ONX 0914 (10 μ M) (Cayman Chemical). Fluorescence was measured (excitation, 380 nm; emission, 460 nm) with a BioTek Synergy H1 plate reader (Agilent) ²³.

SDS-PAGE/Native PAGE and Immunoblotting

For SDS-PAGE, renal extracts from whole-kidney homogenates were prepared with RIPA lysis buffer (Pierce) ^{38, 39}. For native PAGE, Renal extracts from whole-kidney homogenates were prepared with 0.9% digitonin lysis buffer as described in ⁴⁰. Renal extracts (20-30 μ g) were resolved with a Bis-Tris (4%–12%) gel under denaturing (SDS-PAGE) or non-denaturing condition (native PAGE⁴¹) and then transferred to a PVDF membrane. After transfer, the membrane was incubated with 1X Red Alert Western Blot Stain (Millipore) for 10 min at room temperature. The membrane was quickly washed with ddH₂O to visualize protein bands, and images were taken with a FluorchemTM 8900. Western blot analysis was performed with antibodies for $\beta 5i$ (1:1000; LSBio). β -actin (1:1000; Sigma) or GAPDH (1:1000; Signalway) were used as loading control. Probed membranes were washed, incubated with horseradish peroxidase-conjugated secondary goat anti-mouse or goat anti-rabbit antibodies (1:30,000; Seracare KPL), and assayed for enhanced chemiluminescence (Thermo Fisher Scientific). Densitometry was performed with AlphaEase FC software (Alpha Innotech). For all SDS-PAGE western blot analyses, a densitometry ratio of target protein to the corresponding loading control was considered for statistical evaluation. For all native gel western blot analyses, a densitometry ratio of $\beta 5i$ to the corresponding whole-lane density of Red AlertTM stain was considered for statistical evaluation ³⁵.

Quantitative Real-Time PCR

Total RNA was isolated from kidney tissue with an RNeasy Kit (Qiagen), and quantitative real-time PCR was carried out with Superscript III (Invitrogen) reverse-transcribed mRNA, the SYBR Green PCR Kit (Thermo Fisher Scientific), and the PCR primers (**Table 1**)³⁵. The PCR reaction involved 45 cycles under the following conditions: 95°C for 10 s and 60°C for 20 s, and 75°C for 10 s. Amplification of the target gene was normalized to amplification of TATA box binding protein (*Tbp*) and to the levels of an appropriate control using the delta Ct ($2^{-\Delta\Delta Ct}$) method^{35, 39}.

Statistical Analysis

Data are presented as the mean $\pm$ standard error of the mean (SEM) (GraphPad Prism). Data (n=4 per group) were analyzed with a 1-way ANOVA with Tukey's post hoc test or Mann-Whitney U test, 2-way ANOVA (to compare treatment and time effects) and an unpaired Student's t-test was used when comparing differences between the means of 2 groups at a 95% confidence level. Differences with $p < 0.05$ were considered statistically significant.

Results

CS+Tx upregulates the iproteasome

We previously reported that proteasome (constitutive) function was decreased in transplanted rat kidneys 1 day after surgery when the kidneys had been subjected to 18-h CS (CS+Tx group)²³. To determine whether the function of *iproteasome* is dysregulated, we evaluated $\beta 5i$ subunit levels and function under the same conditions (**Fig. 1**). CS+Tx increased both the level of *iproteasome* $\beta 5i$ subunit (**Fig. 1A**) and its function (**Fig. 1B**) at 1 day post-surgery compared to the ATx and sham controls.

*IFN- γ upregulates the i*proteasome in rat kidney proximal tubular cells

We demonstrated that 18-h CS+Tx (syngeneic) increased IFN- γ in rat kidneys^{35, 36}. Here, we observed increased TNF- α expression in rat kidney grafts (syngeneic) after 18-h CS+Tx (**Fig. 2**). To determine whether these cytokines (IFN- γ and TNF- α) have the potential to induce the i

proteasome in rat kidneys, NRK cells were treated with recombinant IFN- γ or TNF- α . Immunoblot assays of renal lysates demonstrated that IFN- γ (**Fig. 3A**), but not TNF- α (**Fig. 3B**), increased $\beta 5i$ subunit levels in NRK cells.

CS+Tx increases renal dysfunction and mortality

Previously, we showed that renal CS (4-h or 18-h) exacerbated organ dysfunction after transplantation (1 day post-surgery)^{24, 25}. To further assay survival, 4 groups (sham, ATx, syngeneic 4-h CS+Tx, and syngeneic 18-h CS+Tx; in which native kidneys were removed and rats lived with transplanted donor or sham kidney only) were assigned to a 7-day survival and renal function study (**Fig. 4**). As expected, serum creatinine (SCr) and blood urea nitrogen (BUN) values (1 day post-transplant) were significantly more elevated in the 18-h CS+Tx animals than in the 4-h CS+Tx animals, indicating a severe loss of graft function reciprocal to the duration of CS (**Fig. 4A and B**). While SCr and BUN values in the ATx and 4-h CS+Tx groups returned to sham levels at 3 days post-transplant, these values continued to increase in the 18-h CS+Tx group (**Fig. 4A and B**). The sham, ATx, and 4-h CS+Tx animals all survived for up to 7 days post-surgery (**Fig. 4C**). Most importantly from a clinical perspective, animals in the 18-h CS+Tx group (syngeneic) did not survive longer than 3 days after transplant (**Fig. 4C**).

iproteasome is induced persistently in kidney allografts 9 days post-surgery

Next, we attempted to evaluate if CS contributes to a persistent increase of *iproteasome* in renal allografts, and we evaluated $\beta 5i$ in kidneys in our allogeneic transplant model (Fischer donor and Lewis recipient) with various CS times (4 h or 18 h). $\beta 5i$ levels increased significantly in rat kidneys in both 4-h CS+Tx and 18-h CS+Tx groups 9 days post-surgery compared to the sham group (**Fig. 5**).

Renal iproteasome composition is dysregulated during CS and after transplantation

The *iproteasome* is a multimeric enzyme complex (i26S/i30S) and is composed of a catalytic core (i20S) with one or two regulatory complexes⁴². $\beta 5i$ is a key subunit of the catalytic core of the *iproteasome*⁴³⁻⁴⁵. To examine the abundance of *iproteasome* composition (i26S/30S and i20S) under various CS and transplant conditions, we performed western blotting of kidney extracts (allogeneic model) under non-denaturing conditions. Western blot with anti- $\beta 5i$ antibody revealed a significant decrease in the abundance of the i26S/30S *iproteasome* subunits after 4-h or 18-h CS, but the i20S catalytic core decreased only after 18-h CS (**Fig. 6A**). Conversely, both 4-h CS+Tx and 18-h CS+Tx increased i26S/30S levels in kidney allografts at 9 days post-surgery, but i20S levels remained unchanged (**Fig. 6B**).

Inhibiting $\beta 5i$ subunit during CS does not protect renal injury after CS+Tx

Finally, we investigated whether the CS-mediated induction of the *iproteasome* negatively affects renal function. Specifically, we tested whether pharmacologically inhibiting $\beta 5i$ with ONX 0914 during CS could mitigate renal injury/dysfunction. In our survival studies, the mortality rate was high for rats in the syngeneic 18-h CS+Tx group at 7 days post-surgery (**Fig. 4**). Therefore, we opted to use the allogeneic rat transplant model with 4-h CS to test the therapeutic effect of ONX 0914. Treating kidneys ex vivo with ONX 0914 did not improve renal function or tubular

injury/fibrosis after CS+Tx (**Fig. 7**). In parallel, we tested several other blood chemistry parameters with the iSTAT CHEM 8+ cartridge. These parameters were not affected at 4-h CS+Tx, and ONX 0914 did not affect blood chemistry, except that it decreased plasma chloride levels (**Fig. 8**). Overall, these results suggest that pharmacological inhibition of $\beta 5i$ during renal CS does not offer protection against renal injury after transplantation.

Discussion

The donor organs are vulnerable to damage caused by cold ischemia. We reported previously that CS reduced mitochondrial respiration and increased ROS, a byproduct of respiration in donor kidneys²⁰⁻²², suggesting that the CS induced cold-ischemic stress. Such stress impairs several pathways, including the constitutive proteasome²⁶. The constitutive proteasome is susceptible to stress, but its variant, the immunoproteasome (*i*proteasome), is resistant to stress, including ROS^{26, 27, 30}. Here, we showed that CS mediated a massive induction of $\beta 5i$, a catalytic subunit of the *i*proteasome, in kidney grafts (both syngeneic and allogenic), suggesting abnormal or aberrant activation of the *i*proteasome. Although the *i*proteasome composition (i20S and i26/30S) was upregulated in transplanted kidneys that were exposed to CS, it was intriguing that some *i*proteasome complexes (i20S and i26/30S) were downregulated during CS. Given that kidney CS increased mitochondrial dysfunction and ROS²⁰⁻²², we postulate that ROS upregulated the *i*proteasome upon transplantation. Future studies are needed to establish the link between mitochondrial dysfunction (ROS) and $\beta 5i$ induction following CS and CS+Tx.

The *i*proteasome is resistant to stress and plays a key role in processing the antigenic peptides and stimulating adaptive immune system⁴⁶. We previously observed CS-mediated induction of several inflammatory pathways, including IFN- γ and complement, and infiltration of CD68-positive macrophages in kidney grafts after CS+Tx³⁴⁻³⁶. Here, we report a substantial

upregulation of the *i*proteasome in both syngeneic and allogeneic models in response to CS (4-18 h), and this induction was sustained in transplanted kidneys for up to 9 days post-surgery. We further showed that exogenous IFN- γ , but not TNF- α , induced the *i*proteasome in proximal tubular cells, suggesting that CS+Tx induced the *i*proteasome (β 5i subunit) via IFN- γ . IFN- γ is a potent inducer of *i*proteasome ⁴⁷. However, future studies should dissect the role of IFN- γ signaling in upregulating *i*proteasome function during CS+Tx. An important function of the *i*proteasome is to process antigenic peptides within antigen-presenting cells (e.g., macrophages and dendritic cells) and to present these peptides to adaptive immune cells (e.g., T cells) through the major histocompatibility complex ⁴⁸. Although CS increased the number of macrophages within transplanted kidneys ³⁴, a limitation of our study was that it lacked an assessment of *i*proteasome levels within the infiltrated macrophage compartment. Similarly, future studies could investigate a link between *i*proteasome and the adaptive immune system (e.g., T cells) because activation of the adaptive immune system is a key mechanism of transplant rejection ⁴⁹.

Currently, kidneys from living donors produce better outcomes than kidneys from deceased donors ^{50, 51}, so decreasing CS-associated injury would make progress toward improving outcomes of transplants with organs from deceased donors. It has been difficult to develop new therapies to support kidney transplants because few studies have investigated the CS-related mechanisms of organ damage. Accordingly, our goal was to evaluate CS-related immunopathogenic mechanisms of injury in kidney grafts after CS+Tx. We identified the upregulated *i*proteasome as a potential therapeutic target and subsequently tested ONX 0914—a specific, reversible inhibitor of the β 5i subunit ³¹ of the *i*proteasome—in the context of CS to mitigate injury and improve kidney function after transplantation. However, this inhibitor, when used during CS alone, did not prevent acute tubular injury/graft function following transplantation. On the other hand, systemic administration of ONX 0914 post-transplant in rats

ameliorated chronic antibody-mediated rejection⁵². Future studies designed to target $\beta 5i$ both during CS and after transplantation are warranted to conclude the role of $\beta 5i$ on acute and chronic kidney injury following transplantation.

In summary, we showed that CS increases the severity of graft dysfunction in a time dependent manner (0-h<4-h<18-h), and further showed that prolonged CS (18-h) decreases overall animal survival. These data further indicate a need for therapeutic intervention during renal CS. Our data also demonstrated that CS alone decreases the abundance of *i*proteasome ($\beta 5i$ subunit); in contrast, reperfusion (transplantation) increases *i*proteasome levels/assembly in renal allografts, which correlates with allograft dysfunction after transplantation. Mechanistically, we showed that IFN- γ is a potential inducer of the *i*proteasome ($\beta 5i$). Finally, inhibiting the *i*proteasome ($\beta 5i$) during CS alone did not confer renal protection after transplantation. Future studies designed to inhibit the *i*proteasome during CS—as well as after transplantation—should be tested to define the role of the *i*proteasome during CS+Tx.

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Table 1. Primer sequences for qRT-PCR (RealTimePrimers.com)

<i>Tnfa</i>:	Forward, 5'-CCCATTACTCTGACCCCTTT-3'	Reverse, 5'-TGAGCATCGTAGTTGTTGGA-3'
<i>Il18</i>:	Forward, 5'-CAGACCACTTTGGCAGACTT-3'	Reverse, 5'-ATCCTTCCATCCTTCACAG-3'
<i>Tbp</i>:	Forward, 5'-CGATAACCCAGAAAGTCGAA-3'	Reverse, 5'-AGATGGGAATTCCAGGAGTC-3'

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Figure Legends

Figure 1. Cold storage increases *iproteasome* ($\beta 5i$) level and function in rat kidneys after transplantation. Lewis rat kidneys were flushed with and stored in cold storage (CS) solution (4°C) for 18 h followed by transplantation to a recipient Lewis rat ($n=4/\text{group}$). Autotransplant (ATx, transplant with no CS) rats were used as transplant control and Sham (right nephrectomy, control kidney) rats were used as healthy controls. **(A)** Immunoblot of $\beta 5i$ (LMP7) proteins in rat kidney homogenates from untreated control, cold storage (CS), sham, autotransplantation (ATx), and cold storage plus transplant (CS+Tx) groups ($n=4/\text{group}$). β -actin served as loading control. Representative blots from 3 independent experiments shown. Graphs show densitometry (mean $\pm$ SEM) normalized to β -actin ($n=4/\text{group}$). **(B)** Immunoproteasome function ($\beta 5i$ peptidase activity) assayed in renal homogenates using a peptide substrate (Ac-ANW-AMC). Data are the mean $\pm$ SEM (bar graph, $n=4/\text{group}$). Differences between group means were compared with Student's t-test (2-group comparison) or ANOVA (3-group comparison). * $p < 0.05$ was considered statistically significant.

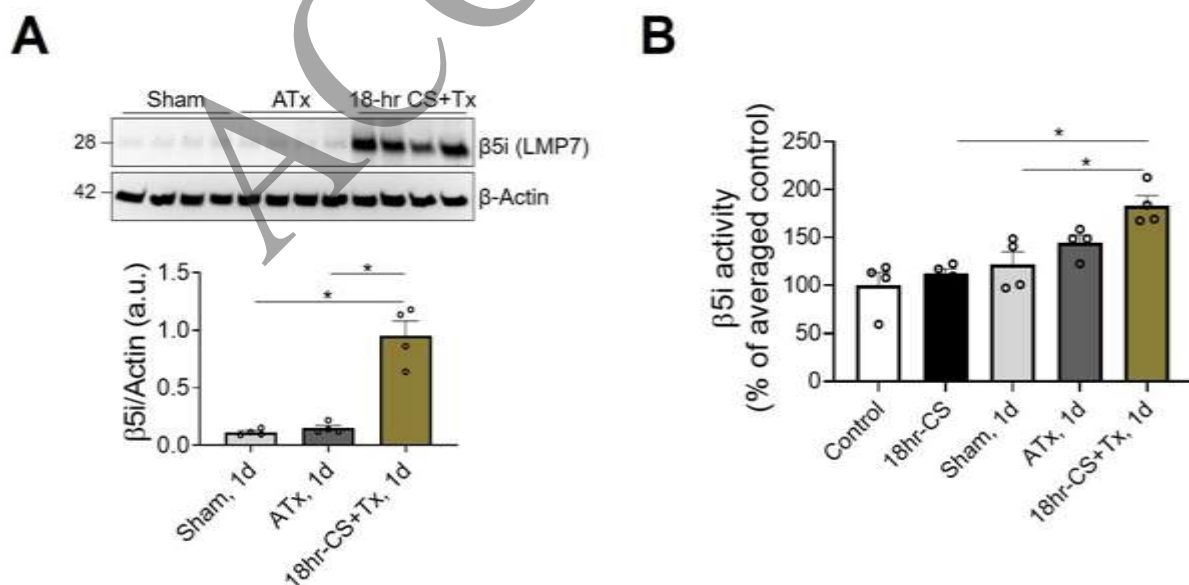


Figure 2. Cold storage injury increases mRNA abundance of tumor necrosis factor-alpha and interleukin-18 in rat kidneys after transplantation. Quantitative real-time PCR (SYBR Green) was performed with rat renal mRNA extracts to assay expression of TNF- α and IL-18 genes after kidney CS or CS+Tx (1 day post surgery). Sham kidneys were used as a control. Data are the mean $\pm$ SEM (bar graphs, n=3/group). Differences between group means were compared with ANOVA (3 groups comparison). *p < 0.05 was considered statistically significant.

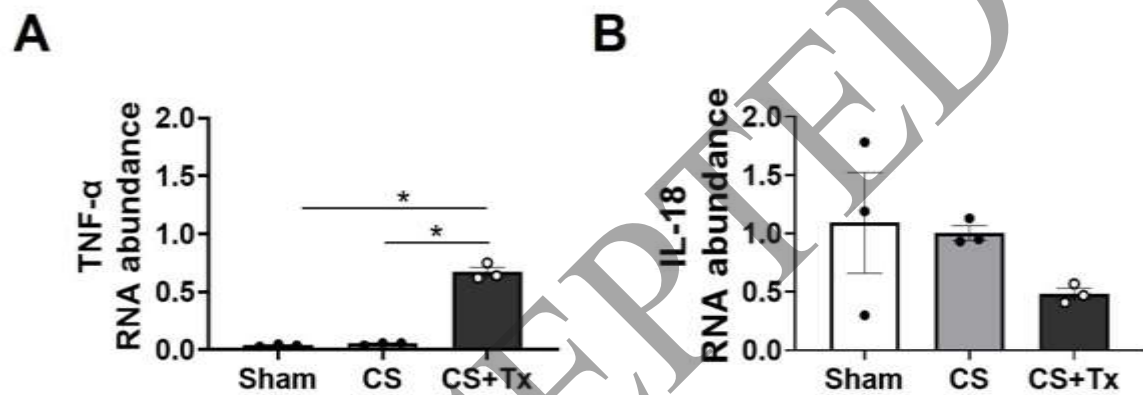


Figure 3. IFN- γ increases *i*proteasome ($\beta 5i$) level in rat proximal tubular cells. Rat proximal tubular (NRK) cells were treated with IFN- γ (10 ng/ml) or TNF- α (10 ng/ml) and solubilized with RIPA lysis buffer. Representative immunoblots (n=3) showing (A) *i*proteasome ($\beta 5i$ protein) and (B) constitutive proteasome ($\beta 5$ protein) levels in NRK cells after treatment with IFN- γ or TNF- α . β -actin served as loading control. Data are the mean $\pm$ SEM (dot plots, n=3). Differences between group means were compared with ANOVA (3-group comparison). *p < 0.05 was considered statistically significant.

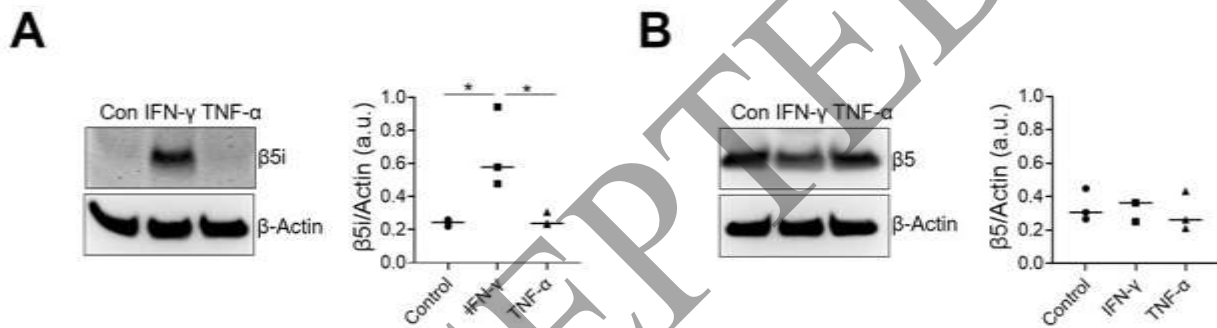


Figure 4. Prolonged cold storage of rat kidneys decreases organ function and rat survival after transplantation. Lewis rat kidneys were flushed with and stored in cold storage (CS) solution (4°C) for 4 h or 18 h followed by transplantation to a recipient Lewis rat (n=4/group). Autotransplant (ATx, transplant with no CS) and Sham rats were used as controls. Renal function was assayed in rat blood 1 and 3 days post-surgery by measuring **(A)** serum creatinine (SCr) and **(B)** blood urea nitrogen (BUN). Data are the mean $\pm$ SEM (dot plots, n=3/group). Differences between group means were compared with ANOVA (> 3-group comparison). *p < 0.05 was considered statistically significant. **(C)** Rat survival after surgery.

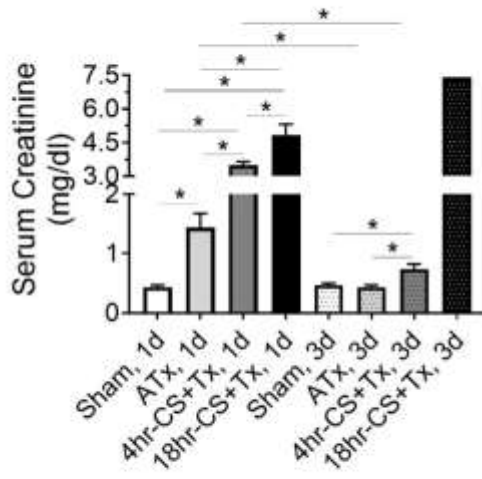
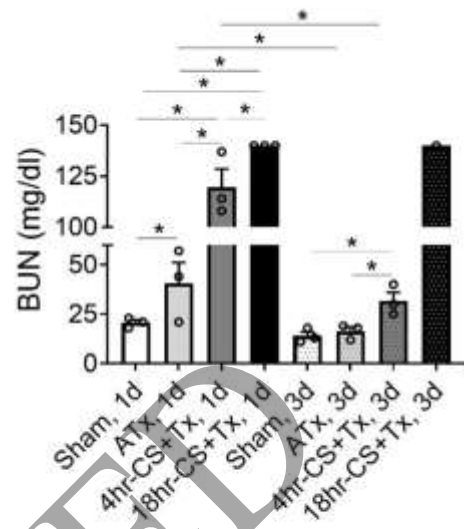
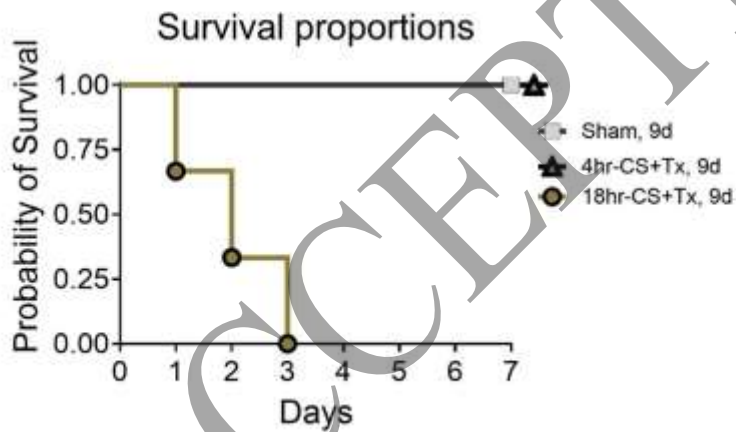
A**B****C**

Figure 5. CS+Tx persistently increases immunoproteasome ($\beta 5i$) level in rat kidney allografts. Fischer rat kidneys were flushed with and stored in cold storage (CS) solution (4°C) for 4 h or 18 h followed by transplantation to a recipient Lewis rat. Sham rats were used as a control. Immunoblot of $\beta 5i$ (LMP7) proteins in rat kidney homogenates from sham and cold storage and transplant (CS+Tx) groups (9 days post-surgery) (n = 4/group). GAPDH served as loading control. Representative blots from 3 independent experiments shown. Graphs show densitometry (mean $\pm$ SEM) normalized to GAPDH (n = 4/group). *p < 0.05 was considered statistically significant.

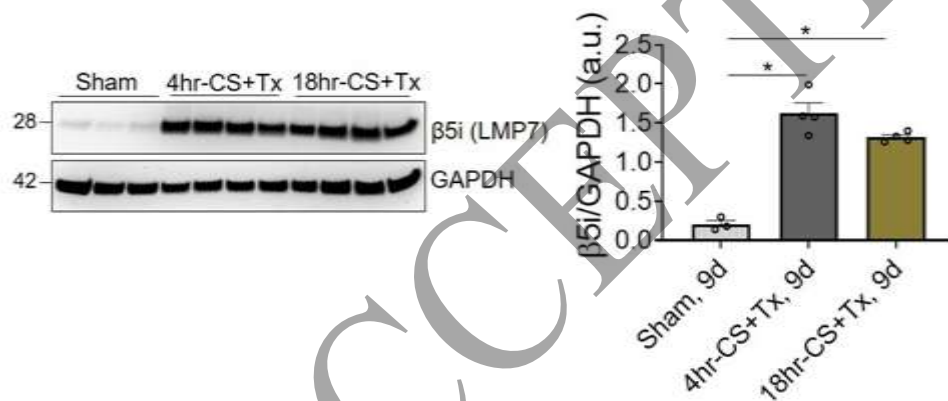


Figure 6. Abundance of *i*proteasome components changes during CS vs. CS+Tx. Fischer rat kidneys were flushed with and stored in cold storage (CS) solution (4°C) for 4 h or 18 h followed by transplantation to a recipient Lewis rat. Untreated (Control) or Sham rats were used as controls. **(A-B)** Non-denatured renal extracts (9 days post-surgery) prepared with digitonin lysis buffer were subjected to native gel electrophoresis followed by western blotting with anti- $\beta 5i$ antibody. Representative blots from 3 independent experiments shown. Graphs show densitometry (mean $\pm$ SEM) normalized to β -actin (n =4/group). *p < 0.05 was considered statistically significant.

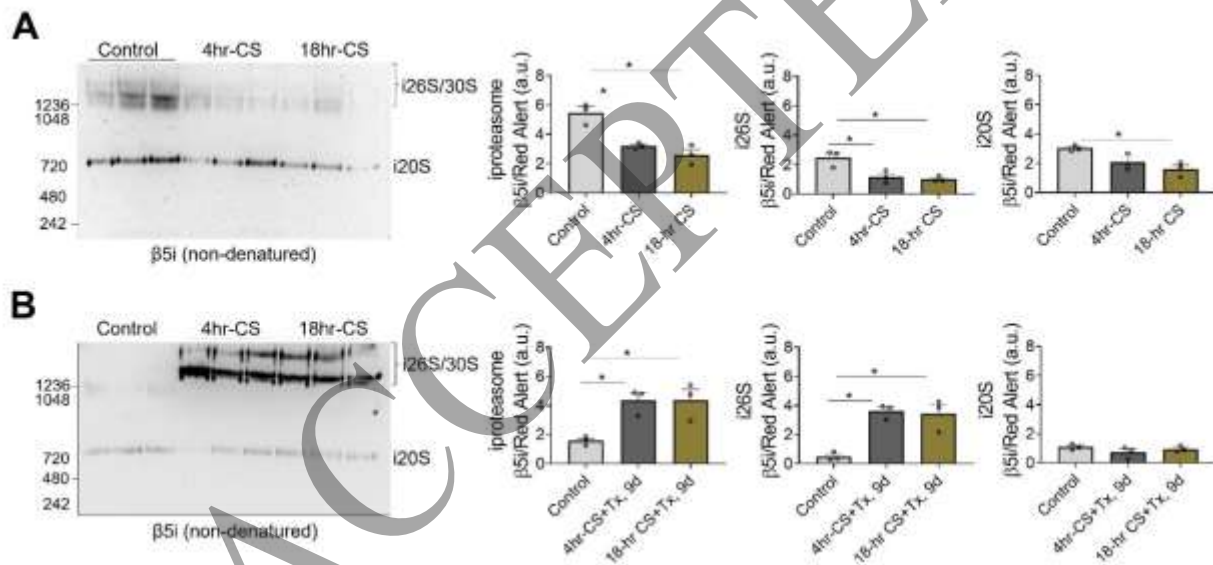


Figure 7. Treating rat kidneys with ONX 0914 in CS solution does not improve allograft function/injury after transplantation. Lewis rat kidneys were treated with the $\beta 5i$ inhibitor ONX 0914 (0 or 100 nM) in cold storage (CS) solution (4°C) for 4 h followed by transplantation to a recipient Fischer rat (n=4/group, 9-day post-surgery). Sham rats were used as a control. **(A-B)** Morphological study of rat kidney sections was performed using Periodic acid-Schiff (PAS) staining. **(A)** Representative images of Periodic acid-Schiff (PAS) stained renal sections (400 X magnification). **(B)** Tubular injury was assessed and scored in a blinded fashion using the PAS stained renal sections from A. Data are the mean $\pm$ SEM (bar graph, n=4/group). **(C-D)** Renal function was assayed in rat blood (9 days post-surgery) by measuring serum creatinine (SCr) **(C)** or blood urea nitrogen (BUN) **(D)**. Data are the mean $\pm$ SEM (violin plots, n=4/group). Differences between group means were compared with Student's t-test (2-group comparison) or ANOVA (3-group comparison). *p < 0.05 was considered statistically significant. **(E) & (F)** Representative images of KIM1 and NGAL immunohistochemistry of renal sections (400 X magnification; n=4/group, respectively). **(G)** Representative images of Masson's trichome staining of renal sections (400 X magnification; n=4/group). O Medulla: Outer Medulla; I Medulla: Inner Medulla

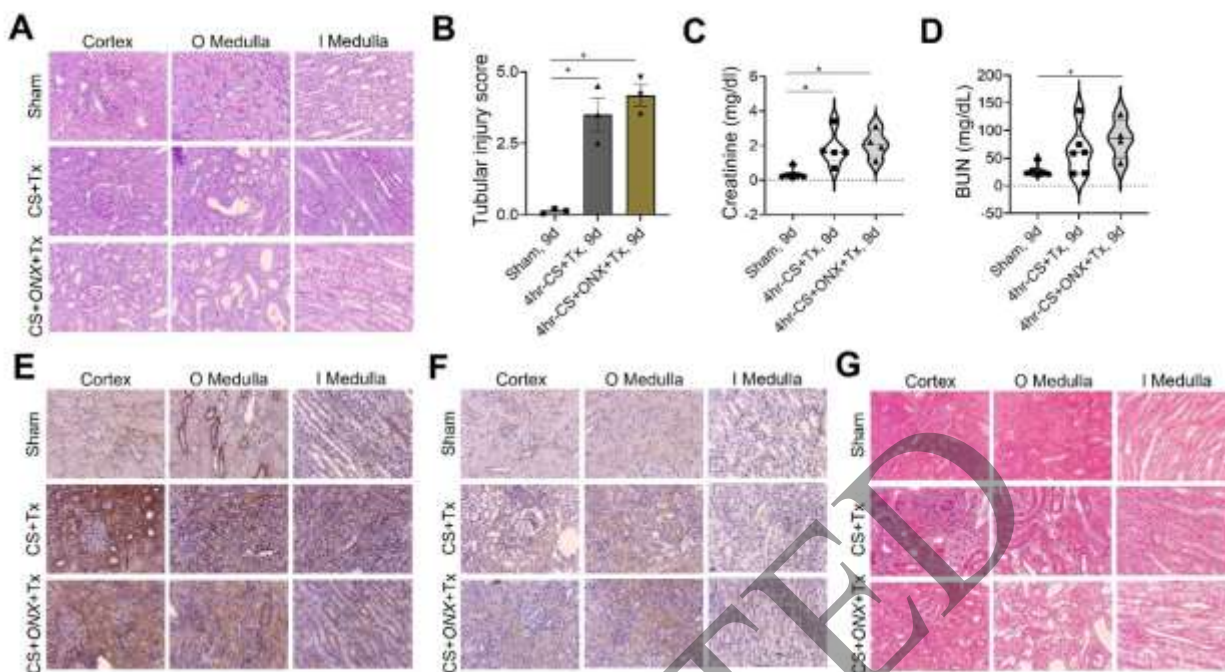


Figure 8. Effect of treating rat kidneys with ONX 0914 during CS on blood chemistry after CS+Tx. Lewis rat kidneys were treated with the β 5i inhibitor ONX 0914 (0 or 100 nM) in cold storage (CS) solution (4°C) for 4 h followed by transplantation to a recipient Lewis rat (n=4/group). Sham rats were used as a control. **(A-I)** Blood chemistry measured with i-STAT Chem8⁺ cartridge and blood chemistry analyzer. Data are the mean $\pm$ SEM (bar graph, n=4). Differences between group means were compared with Student's t-test (2-group comparison) or ANOVA (3-group comparison). *p < 0.05 was considered statistically significant.

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