

GSK3 β Promotes Apoptosis after Renal Ischemic Injury

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ABSTRACT

The mechanism by which the serine-threonine kinase glycogen synthase kinase-3 β (GSK3 β) affects survival of renal epithelial cells after acute stress is unknown. Using *in vitro* and *in vivo* models, we tested the hypothesis that GSK3 β promotes Bax-mediated apoptosis, contributing to tubular injury and organ dysfunction after acute renal ischemia. Exposure of renal epithelial cells to metabolic stress activated GSK3 β , Bax, and caspase 3 and induced apoptosis. Expression of a constitutively active GSK3 β mutant activated Bax and decreased cell survival after metabolic stress. In contrast, pharmacologic inhibition (4-benzyl-2-methyl-1,2,4-thiadiazolidine-3,5-dione [TDZD-8]) or RNA interference-mediated knockdown of GSK3 β promoted cell survival. Furthermore, RNA interference-mediated knockdown of Bax abrogated the cell death induced by constitutively active GSK3 β . In a cell-free assay, TDZD-8 inhibited the phosphorylation of a peptide containing the Bax serine¹⁶³ site targeted by stress-activated GSK3 β . In rats, TDZD-8 inhibited ischemia-induced activation of GSK3 β , Bax, and caspase 3; ameliorated tubular and epithelial cell damage; and significantly protected renal function. Taken together, GSK3 β -mediated Bax activation induces apoptosis and tubular damage that contribute to acute ischemic kidney injury.

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Acute ischemic renal failure remains a common cause of death that precipitates organ failure by causing apoptosis, necrosis, autophagy, and the desquamation of viable proximal tubule epithelial cells from the basement membrane.^{1–7} Although epithelial cell apoptosis has been detected in the intact kidney after ischemia^{8–10} and in cultured renal epithelial cells subjected to metabolic stress,^{11,12} it has been difficult to assess its contribution to human organ failure.² More importantly, efforts to improve outcome in ischemic organ failure have been limited by an incomplete understanding of the signal events that regulate renal epithelial cell injury.

Glycogen synthase kinase-3 β (GSK3 β) is a 47-kD serine-threonine kinase that was first observed to phosphorylate and inactivate glycogen synthase, a distal enzyme in the glycogen synthesis pathway.¹³ GSK3 β is an ideal “survival” enzyme, because it controls several extrametabolic processes that are perturbed by ischemia, including cytoskeletal dynamics, gene expression, proliferation, and apoptosis.^{14–22}

Normally, GSK3 β is suppressed by proliferative, pro-survival signals that increase serine⁹ phosphorylation, such as WNT ligands, EGF, IGF-I and -II, and fibroblast growth factor,^{16,19,23} as well as Akt.¹⁹ Conversely, GSK3 β is activated by noxious stimuli, including serum starvation,²³ hypertonic stress,²⁴ potassium deprivation,²⁵ hypoxia,²⁶ endotoxin exposure,²⁷ and tissue ischemia.^{21,28–31} In acute models of injury, GSK3 β promotes the systemic inflammatory response, increases the proinflammatory release of cytokines, induces apoptosis, and alters cell proliferation.^{32,33}

Although the mechanism by which GSK3 β reg-

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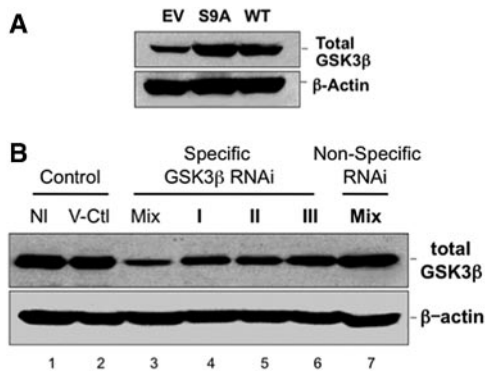


Figure 1. Effect of transfection and RNAi on GSK3 β expression is shown. (A) Cell GSK3 β content after transfection with empty vector (EV), constitutively active (S9A), or WT GSK3 β (top); β -actin loading control (bottom). (B) Total GSK3 β content in nontransfected cells (NI), in cells transfected with vector only (V-Ctl), a mixture of three (Mix; top), single specific RNAi directed against GSK3 β (I, II, or III) or a mixture of nonspecific RNAis (Mix-Non-Specific-RNAi); β -actin is loading control (bottom). Each lane contains 40 μ g of total protein; results are representative of at least three separate studies.

ulates epithelial cell survival after ischemia is poorly characterized, substantial evidence indicates that GSK3 β inhibition preserves organ function after ischemia of the brain,³⁴ heart,³⁵ and gut.³⁶ In the ischemic cell, GSK3 β targets multiple proteins with the potential to regulate survival; however, most are transcription factors such as β -catenin/TCF-Lef genes, cAMP response element binding protein, heat shock factor-1, nuclear factor of activated T cells, and NF- κ B as well as cyclin D1^{17,18,26,37,38} that are unlikely to mediate acute cell survival during or immediately after transient ischemia.

In contrast to these transcription factors and cell-cycle regulators, immediate cell fate after de-energization or ischemia is regulated by BCL2 proteins that target mitochondria, resulting in the activation of both caspase-dependent and -independent cell death cascades.^{12,39–41} Intriguingly, GSK3 β directly phosphorylates and activates Bax, a proapoptotic member of the BCL2 family.⁴² Both Bax and GSK3 β are constitutively expressed and are regulated by phosphorylation at specific serine sites. Phosphorylation of serine⁹ inhibits GSK3 β ,^{19,43} whereas dephosphorylation of this residue activates GSK3 β , thereby promoting downstream serine¹⁶³ phosphorylation and activation of Bax.⁴² GSK3 β has been linked to mitochondrial dysfunction after oxidative stress caused by renal ischemia or hypoxia-induced oxidative stress.⁴⁴ In addition, GSK3 β has been implicated in promoting mitochondrial permeabilization, a terminal event, by direct phosphorylation and destabilization of MCL-1, an antiapoptotic member of the BCL2 family.⁴⁵ By directly phosphorylating and activating Bax, we propose that GSK3 β mediates mitochondrial injury and determines early epithelial cell fate *in vitro* as well as the severity of tubular injury and organ dysfunction *in vivo* after an acute ischemic insult.

Using models of rotenone-induced ATP depletion *in vitro* and transient renal artery occlusion in rats *in vivo*, this study

shows that direct manipulation of GSK3 β regulates Bax activation, apoptosis, tubular injury, and organ dysfunction after metabolic stress or ischemia. Selective GSK3 β upregulation (constitutively active GSK3 β) or suppression (using RNA interference [RNAi] or 4-benzyl-2-methyl-1,2,4-thiadiazolidine-3,5-dione [TDZD-8], a selective GSK3 β inhibitor), causes parallel changes in Bax and caspase 3 activation, apoptosis, and tubular injury, indicating that GSK3 β mediates acute epithelial cell survival partly by disrupting the balance of pro- versus antiapoptotic BCL2 proteins in the renal epithelial cell. Specifically, GSK3 β primarily operates *via* a Bax-dependent mechanism to regulate epithelial cell survival.

RESULTS

Transfection of renal epithelial cells with constitutively active (S9A) or wild-type (WT) GSK3 β resulted in an expected increase in total GSK3 β content compared with empty vector (Figure 1A). In contrast, exposure of cells to one of three GSK3 β RNAi directed against distinct GSK3 β regions resulted in variable downregulation of GSK3 β expression compared with either normal (nontransfected) or vector-exposed cells or cells incubated with nonspecific RNAi (Figure 1B, lanes 1, 2, and 7, respectively). Exposure to a mixture of three RNAis resulted in a marked reduction in immunoreactive GSK3 β content compared with control (Figure 1B, lane 3).

Compared with control, transfection with constitutively active GSK3 β (GSK3 β -S9A) increased total GSK3 β content before, during, and after metabolic stress (Figure 2B), without marked differences in p-serine⁹ GSK3 β content between the two groups (Figure 2A) as detected with a phospho-serine⁹-specific GSK3 β antibody. The later observation is due to the fact that the constitutively active GSK3 β -S9A mutant cannot be inactivated by phosphorylation at the serine⁹ position. Although GSK3 β -S9A expres-

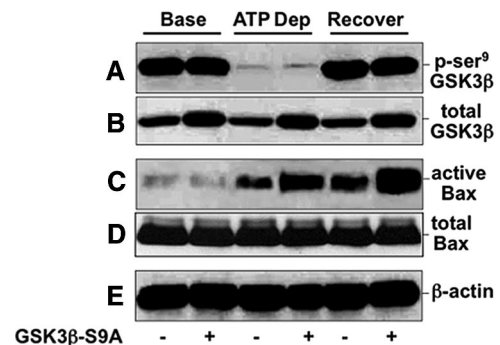


Figure 2. Effect of ATP depletion on GSK3 β , active Bax, caspase 3, and apoptosis is shown. Content of total and p-serine⁹ GSK3 β , active Bax (6A7 epitope), total Bax, and β -actin in cells at baseline ("Base"); after 60 minutes of metabolic stress (ATP Dep); and after 60 minutes of recovery (Recover) in cells that express empty vector (GSK3 β S9A -) or constitutively active GSK3 β (GSK3 β S9A +). Immunoblot results are representative of two to three independent studies; each lane contains 40 μ g of total protein.

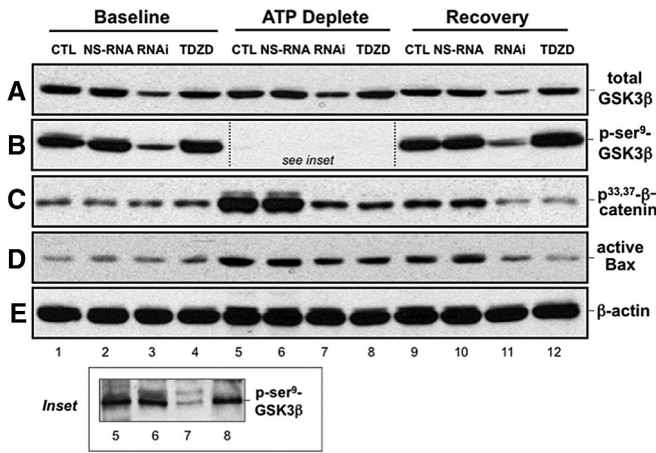


Figure 3. Effect of GSK3β regulation on active Bax after ATP depletion is shown. Content of p-serine⁹ GSK3β, total GSK3β, phospho^{33,37}-β-catenin (a GSK3β substrate), active Bax (6A7 epitope), and β-actin (loading control) in cells at baseline, after 90 min metabolic stress (ATP Deplete), and after 60 min recovery (Recovery) in cells that were not exposed to RNAi (CTL) or were exposed to nonspecific RNAi (NS-RNA), anti-GSK3β RNAi (RNAi), or 10 μM TDZD-8 (TDZD), a GSK3β kinase inhibitor. Inset shows the indicated region of immunoblot in panel B subjected to a longer exposure time to permit bands to be visualized. Immunoblot results are representative of two to three independent studies; each lane contains 40 μg of total protein.

sion did not alter the content of active Bax at baseline (Figure 2C), this GSK3β mutant promoted Bax activation (detected by a 6A7 epitope-specific antibody) during and after stress in the absence of changes in total Bax content (Figure 2D). These results show that constitutively active GSK3β expression promotes stress-induced Bax activation.

To confirm that GSK3β regulates Bax activation, we used both molecular and biochemical maneuvers to inhibit GSK3β. As expected, only specific RNAi downregulated total GSK3β expression at baseline (Figure 3A, lane 3), after 90 min stress (Figure 3A, lane 7), and during recovery (Figure 3A, lane 11).

The decrement in total GSK3β caused by specific RNAi also decreased p-serine⁹ GSK3β content at these same time points (Figure 3B). Because phospho-GSK3β content was markedly reduced during ATP depletion (Figure 3B, lanes 5 through 8), the exposure time of the immunoblot was extended to increase sensitivity (Figure 3B, inset). With prolonged exposure time, only specific RNAi decreased p-serine⁹ GSK3β during ATP depletion. At baseline, no difference in the content of phosphoserine^{33,37} β-catenin, a substrate that is phosphorylated at these sites by GSK3β, was detected in the presence of nonspecific or specific RNAi (Figure 3C, lane 1 versus lanes 2 and 3). Likewise, TDZD-8, an agent that decreases kinase activity without affecting p-serine⁹ GSK3β content, did not change phosphoserine^{33,37} β-catenin (Figure 3C, lane 4). After stress, however, both specific RNAi and TDZD-8 markedly inhibited β-catenin serine^{33,37} phosphorylation (Figure 3C, lanes 7 and 8 and lanes 11 and 12). Furthermore, both specific RNAi and TDZD-8 diminished stress-induced Bax activation (Figure 3D, lanes 7 and 8 versus 5 and 6 and lanes 11 and 12 versus lanes 9 and 10).

Expression of either WT GSK3β or constitutively active GSK3β significantly reduced cell survival after stress (Figure 4A). To limit cell death in control, we reduced the duration of ATP depletion to 60 min in this protocol. Interestingly, the lowest cell survival was observed in cells that express constitutively active GSK3β, a mutant that cannot be inactivated by phosphorylation (e.g., by Akt). In contrast, both specific RNAi directed against GSK3β or TDZD-8 significantly improved cell survival after stress (*P* < 0.05; Figure 4B). In fact, survival exceeded 90% during recovery from stress when GSK3β was inhibited with either of these molecular or pharmacologic manipulations; however, none of these maneuvers (GSK3β expression, specific or nonspecific RNAi or TDZD-8 exposure) caused significant changes in cell survival at baseline (data not shown).

To confirm that reduced survival was attributable to apoptosis, we stained cells with Hoechst dye. At baseline, exposure to empty vector or vehicle (DMSO) did not alter cell morphology or increase nuclear staining with Hoechst dye (Figure 5A,

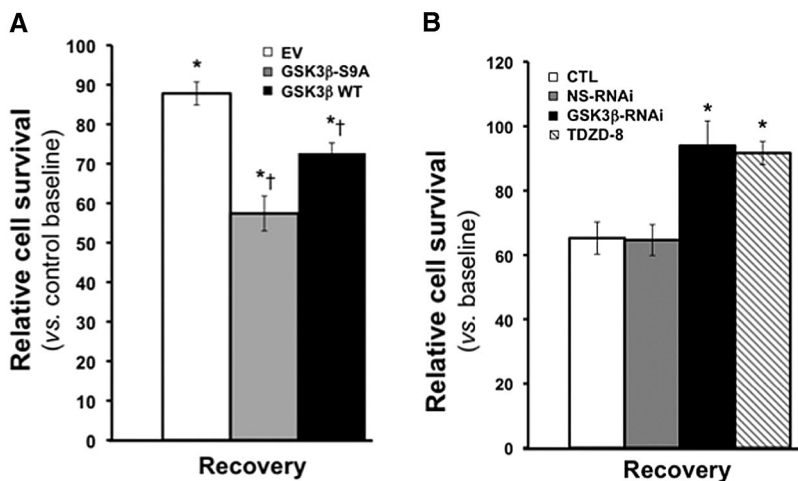


Figure 4. Effect of GSK3β expression on renal cell survival after ATP depletion is shown. (A) Cell survival [3-(4,5 dimethylthiazol)-2,5-diphenyl tetrazolium bromide (MTT) assay] after 60 minutes of stress with 6 hours of recovery in cells expressing empty vector (EV), constitutively active GSK3β (GSK3β-S9A), or WT GSK3β (GSK3β-WT) compared with control at baseline. **P* = 0.05 versus EV by ANOVA; *n* = 5. (B) Cell survival after 90 minutes of stress with 6 hours of recovery in control (CTL) or cells exposed to either nonspecific (NS RNAi) or specific (GSK3β RNAi) RNAi directed against GSK3β or to TDZD-8. **P* < 0.05 versus EV by ANOVA; *n* = 5. None of the maneuvers significantly altered cell survival at baseline (data not shown).

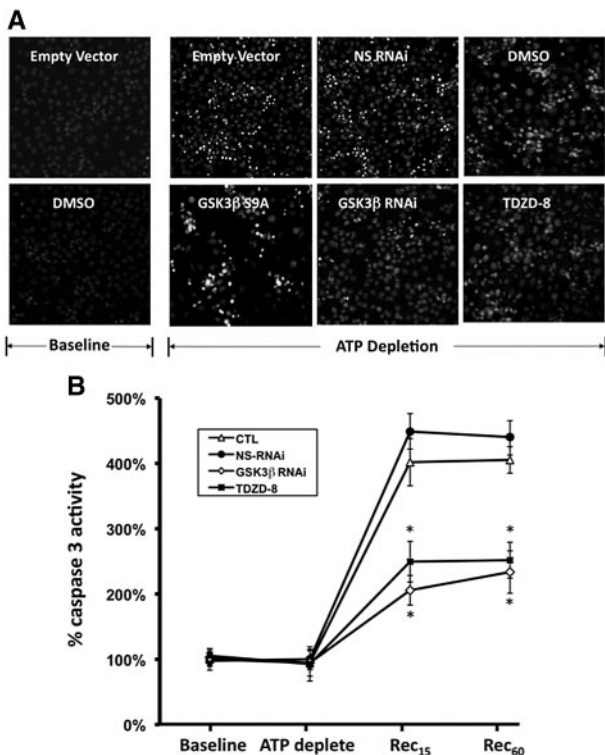


Figure 5. Effect of ATP depletion on apoptosis and caspase 3 activity is shown. (A) Apoptosis assessed by staining cells with Hoechst dye before stress (Baseline) and 3 hours after 60 minutes of metabolic stress (ATP Depletion) in cells that express empty vector (Empty Vector), constitutively active GSK3 β (GSK3 β -S9A), nonspecific RNAi (NS-RNAi), or GSK3 β -specific RNAi (GSK3 β RNAi) or are exposed either to vehicle alone (DMSO) or to TDZD-8; apoptotic cells appear round and smaller and have more brightly stained nuclei than healthy cells stained with Hoechst dye. (B) Caspase 3 enzyme activity at baseline, immediately after stress (ATP deplete), and 15 or 60 minutes of recovery from stress (Rec₁₅ and Rec₆₀) in control (CTL) versus cells exposed to NS-RNAi, specific RNAi directed against GSK3 β (GSK3 β RNAi), or TDZD-8. Values are normalized to the baseline for each group. * $P < 0.05$ versus either control or NS-RNAi.

left-hand panels). In contrast to normal-appearing, relatively large, faintly stained nuclei at baseline, ATP depletion caused marked changes in morphology and increased Hoechst staining in cells transfected with empty vector or exposed to either nonspecific RNAi or vehicle. In these groups, characteristic features of apoptosis were observed: many nuclei appeared smaller or rounded or were fragmented into apoptotic bodies that intensely stained with Hoechst (Figure 5A). Expression of constitutively active GSK3 β dramatically increased ATP depletion-induced apoptosis and the appearance of apoptotic bodies and also decreased the number of adherent cells (Figure 5A, bottom). In contrast, specific GSK3 β RNAi or TDZD-8 decreased the number of apoptotic cells and increased cell adherence after ATP depletion (Figure 5A, bottom).

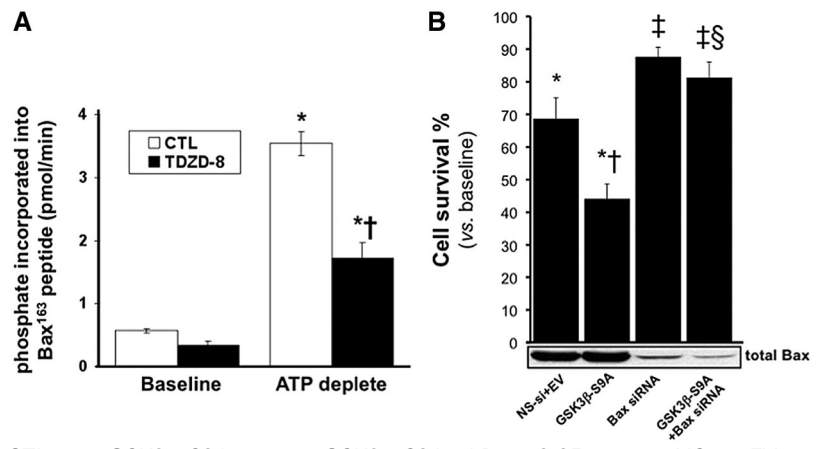
At baseline, caspase 3 enzyme activity, another surrogate measure of apoptosis, did not differ in control (nontrans-

ected, non-RNAi exposed cells) versus cells exposed to non-specific RNAi, GSK3 β -specific RNAi, or TDZD-8 (Figure 5B). Compared with control or nonspecific RNAi, however, both specific GSK3 β RNAi and TDZD-8 exposure significantly decreased stress-induced caspase 3 activation ($P < 0.05$).

To strengthen the causal relationship between the activation of GSK3 β and Bax during ATP depletion, we devised a cell-free assay that measures the phosphorylation of Bax serine¹⁶³, the residue targeted by GSK3 β (see the Concise Methods section). In this assay, stress caused a six-fold increase in the phosphorylation of Bax serine¹⁶³ (0.57 ± 0.03 versus 3.54 ± 0.19 pmol of phosphate incorporated into serine¹⁶³ of the Bax peptide, respectively; $P < 0.05$; Figure 6A). In contrast, 10 μ M TDZD-8 significantly reduced GSK3 β -specific Bax serine¹⁶³ phosphorylation during stress by $>50\%$ (3.54 ± 0.19 versus 1.72 ± 0.19 pmol of phosphate incorporated; $P < 0.05$ versus vehicle control). To support a pathogenic role of GSK3 β in mediating Bax activation and apoptosis after stress, we assessed survival in cells in which Bax was suppressed before expressing constitutively active GSK3 β (Figure 6B). Bax-specific RNAi (but not nonspecific RNAi) decreased Bax content (Figure 6B, immunoblot). Compared with control, nonspecific RNAi, Bax-specific RNAi, or GSK3 β -S9A did not alter baseline cell survival (data not shown). GSK3 β -S9A expression significantly decreased cell survival compared with control (44 ± 5 versus $69 \pm 6\%$, respectively, Figure 6B). In contrast, suppressing Bax with specific RNAi resulted in almost 90% cell survival after stress. Interestingly, Bax suppression also significantly reduced cell death associated with GSK3 β -S9A expression. In fact, Bax suppression almost completely prevented the loss of cell viability associated with overexpression of the constitutively active GSK3 β mutant (81 ± 5 versus $44 \pm 5\%$ cell survival, respectively; $P > 0.05$).

In renal cortical homogenates, transient renal ischemia *in vivo* caused virtually identical biochemical perturbations as were caused by metabolic stress *in vitro*. Specifically, ischemia activated GSK3 β as indicated by a decrease in serine⁹ phosphorylation (Figure 7A, panel 1) but not in renal cortices harvested from either a sham-operated kidney with its renal artery encircled with a nonocclusive ligature or the nonischemic, contralateral kidney. Total GSK3 β content did not change with any experimental maneuver (Figure 7, panel 2). Reperfusion for 30 minutes partially inactivated, whereas 60 minutes of reperfusion completely inactivated GSK3 β (*i.e.*, restored phospho-serine GSK3 β content to the preischemic baseline level) in the sham kidney (data not shown). Administration of TDZD-8 (1 mg/kg intravenously) as a single dose 1 hour before ischemia blocked GSK3 β kinase activity as indicated by a partial inhibition of stress-induced serine^{33,37} β -catenin phosphorylation (Figure 7, panel 3) but without altering total β -catenin content (Figure 7, panel 4). Ischemia also caused a dramatic increase in the accumulation of the 17-kD active caspase 3 cleavage product (Figure 7, panel 5) as well as marked Bax activation (Figure 7, panel 6). In contrast, TDZD-8 not only reduced Bax activation (Figure 7, panel 7) without alter-

Figure 6. Effect of metabolic stress on Bax serine¹⁶³ phosphorylation is shown. (A) ³²P incorporation into a Bax-specific peptide containing serine¹⁶³, the residue targeted by GSK3β in a cell-free system (see the Concise Methods section), at baseline, immediately after 60 minutes of ATP depletion, and in presence and absence of 10 μM TDZD-8. **P* < 0.05 TDZD-8 versus vehicle; *n* = 5. (B) Survival (MTT assay) 8 hours after 60 minutes of ATP depletion in cells exposed to nonspecific RNAi plus empty vector adenovirus (NS-si+EV), constitutively active GSK3β (GSK3β-S9A), Bax-specific RNAi (Bax-siRNA), or constitutively active GSK3β plus Bax specific RNAi (GSK3β-S9A+Bax siRNA). **P* < 0.05 versus no ATP depletion; †*P* < 0.05 versus ATP-depleted CTL, no GSK3β-S9A versus GSK3β-S9A; ‡*P* < 0.05 versus NS-si+EV or GSK3β-S9A; §*P* > 0.05 versus Bax siRNA alone; *n* = 5.



ing total Bax content but also prevented the loss of immunoreactive Bcl2 after renal ischemia (Figure 7, panel 8). These results were confirmed by statistical analysis of densitometric data pooled from several immunoblot studies (Figure 7B).

Compared with nonischemic control, transient ischemia caused marked tubular injury, loss of brush border, tubular dilation, and intratubular cast formation (Figure 8A). In contrast, a single dose of TDZD-8 preserved tubular morphology after ischemic stress. Higher power magnification confirmed that TDZD-8 exposure *per se* did not alter the appearance of the cortical tubules at baseline but improved tubular morphology after ischemia (Figure 8B). Blinded analysis of multiple sections of renal cortex revealed that inhibition of GSK3β resulted in a marked decrease in several parameters of proximal tubule injury, including dilation, vacuolization, brush border loss, epithelial cell detachment, and intratubular cast formation (Table 1). A modest reduction in leukocyte infiltration and capillary edema were also detected in TDZD-8-treated animals. Importantly, inhibition of the apoptosis cascade and improvement in tubular morphology after ischemia resulted in a significant decrement in serum creatinine on days 1 through 6 after injury compared with vehicle-treated control (*P* < 0.03; *n* = 6; Figure 9). Creatinine peaked at 3.9 mg/dl on day 2 in vehicle-exposed *versus* 2.6 mg/dl in TDZD-8-treated

animals. On the final day of observation, creatinine was 60% higher in the vehicle-exposed group. All 12 animals survived renal ischemia.

DISCUSSION

A central feature of our *in vitro* and *in vivo* studies is the observation that stress transiently activates GSK3β, a multifunctional protein kinase that resides at the “biochemical intersection” between normal cell metabolism and cell survival. Although GSK3β affects inflammation and cell proliferation, it has recently been linked to apoptosis.^{27,33} Specifically, GSK3β affects BCL2 family proteins that regulate mitochondrial membrane permeabilization,⁴⁵ a primary event in the apoptotic cell death pathway. In addition, Plotnikov *et al.*⁴⁴ showed that renal ischemia or hypoxia increase reactive oxygen species formation, especially within renal tubules, reduced both mitochondrial membrane potential and respiration, and linked reactive oxygen species generation to GSK3β activation. This study extends those observations by identifying early, GSK3β-mediated events in the cell death pathway that link Bax-mediated mitochondrial injury to renal epithelial cell apoptosis and organ dysfunction.

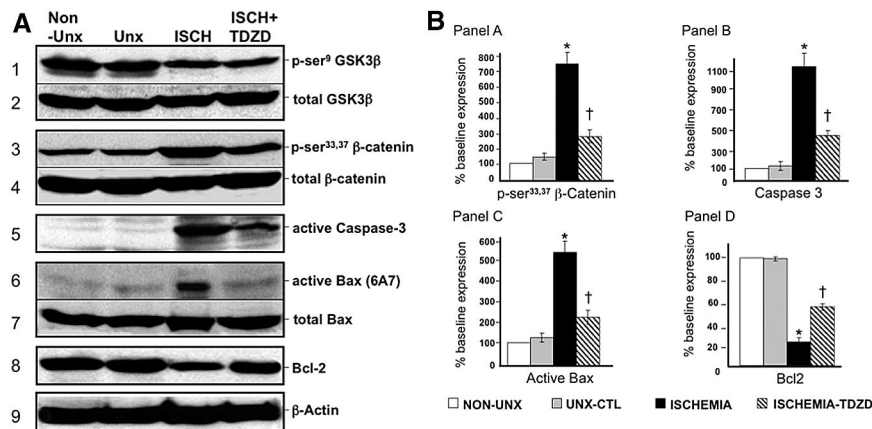


Figure 7. Effect of unilateral renal ischemia in the presence and absence of a GSK3β inhibitor on the apoptosis pathway is shown. (A) Effect of 30 minutes of renal ischemia on p-serine⁹ and total GSK3β content, p-serine^{33,37} and total β-catenin (a GSK3β substrate), active caspase 3, active and total Bax and Bcl-2; β-actin is a loading control (bottom). (B) Densitometric analysis of studies shown in A. **P* < 0.05 versus non-uninephrectomy (Non-Unx); †*P* < 0.05 versus Unx control; *n* = 4.

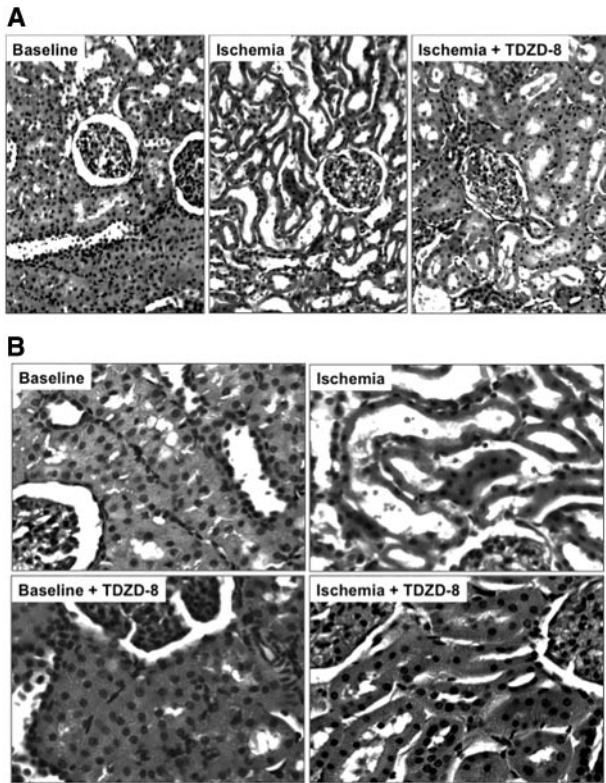


Figure 8. Effect of renal ischemia in the presence and absence of a GSK3 β inhibitor on renal histology is shown. (A) Hematoxylin- and eosin-stained sections harvested from the renal cortices of rats at baseline or 1 hour after 30 minutes of renal ischemia in the presence of vehicle (Ischemia) or TDZD-8 (Ischemia + TDZD-8). (B) Hematoxylin- and eosin-stained sections harvested from the renal cortices of rats at baseline in the presence of vehicle (Baseline) or TDZD-8 (Baseline + TDZD-8) and 1 hour after 30 minutes of renal ischemia (Ischemia versus Ischemia + TDZD-8). Magnifications: $\times 20$ in A; $\times 40$ in B.

How does GSK3 β induce mitochondrial injury and apoptosis? Evidence shows that GSK3 β directly phosphorylates and activates Bax. Using a site-directed mutation of Bax serine¹⁶³, Linseman *et al.*⁴² showed that this residue is critical for activating Bax and for translocating it to mitochondria. In this study, both metabolic stress and pharmacologic GSK3 β inhibition altered the ability of GSK3 β to phosphorylate Bax at the serine¹⁶³ site in our cell-free assay (Figure 4), suggesting that GSK3 β and Bax activation are causally linked. Interestingly, AKT has been shown to phosphorylate serine¹⁸⁴, inactivating Bax.⁴⁶ By simultaneously inactivating AKT and activating GSK3 β , both ATP depletion and ischemia induce a perfect *ménage a trois* relationship that involves both of these kinases and Bax in a manner that induces mitochondrial membrane injury and apoptosis. That expression of constitutively active GSK3 β alone and in the absence of metabolic stress does not activate Bax (Figure 2) or alter cell survival suggests that maximal Bax activation requires additional events. This observation is consistent with reports that distinct Bax domains (and likely Bax phosphorylation events⁴⁷) regulate the signals that

retain Bax in the cytosol or mediate its translocation to mitochondria.^{48,49}

The first committed step in the Bax activation pathway is conformational change, followed by Bax oligomerization and translocation to mitochondria.⁵⁰ Each of these three steps was recently confirmed in epithelial cells subjected to metabolic stress.⁵¹ After translocation, Bax permeabilizes the outer mitochondrial membrane by either forming *de novo pores* or opening existing ones.^{52,53} Considerable controversy surrounds the mechanism by which “Bax attack” alters membrane permeability. Similarly, the specific membrane site targeted by Bax is debated.^{53,54} Interaction between Bax and the voltage-dependent anion channel (VDAC) has received considerable attention, because VDAC is a key component of the membrane pore transition (MPT) complex responsible for releasing both cytochrome c and apoptosis-inducing factor (and others) after an apoptogenic stress.⁵⁵ Although the role of VDAC in MPT has been questioned using knockout of VDAC 1 and 3 and knockdown of VDAC 2,⁵⁶ substantial evidence suggests that GSK3 β regulates MPT. By phosphorylating VDAC and displacing hexokinase, a second MPT complex component, GSK3 β could “prepare” the VDAC-binding site for attack by Bax.^{26,57}

Regardless of the mechanism, activated Bax is clearly mitotoxic, particularly when the balance between pro- and anti-apoptotic proteins is perturbed.^{39,53} The degradation of Bcl2 during metabolic stress *in vitro*¹² or renal ischemia *in vivo* (Figure 7) by caspase 3 not only eliminates an anti-Bax protein⁵⁸ but also generates a potent “Bax-like” fragment.^{59,60} In the renal cortex, ischemia induced a >20 -fold increase in the active Bax-Bcl2 ratio (Figure 7), a critical determinant of the “apoptotic set point.”³⁹ GSK3 β inhibition *in vivo* partially restores the Bax-Bcl2 ratio by inhibiting Bax activation as well as by preserving Bcl2.

Although renal failure after ischemia has traditionally been attributed to acute tubular necrosis, it is now clear that necrosis fails to account for the severity of organ impairment, especially in humans.^{2,3,61} This information increases the likelihood that other forms of cell injury mediate organ dysfunction.^{2,5} The contribution of apoptosis to acute organ injury is confounded by the evanescent nature of apoptotic cells and the stochastic nature of the apoptotic process. Despite these limitations, caspase inhibitors reduce renal impairment⁶² and selective caspase knockout improves organ function in the posts ischemic mouse,⁶³ indicating that apoptosis contributes to ischemic renal failure. In this study, ischemia caused tubular dilation, accumulation of detached epithelial cells into the lumen, intratubular cast formation, and mild inflammation in the relative absence of frank tubular necrosis (Figure 8). As well, a GSK3 β inhibitor afforded a greater reduction in the indices of tubular injury than it did to reduce inflammation (Table 1). The modest reduction in the inflammatory response observed in this study may be because GSK3 β activation promotes the systemic inflammatory response, increases cytokine release, and alters cell proliferation required for recovery after diverse renal insults.^{27,32,33,64} In a murine model of LPS-induced acute renal

Table 1. Histologic injury score in renal cortical tissue sections obtained before ischemia (baseline), immediately after 30 minutes of renal artery occlusion (ischemia), and after 30 minutes of recovery

Parameter	TDZD-8					
	Baseline		Ischemia		Recovery	
	Sham -	Sham +	-	+	-	+
Proximal tubule dilation	0	0	1	0	2 to 3	1
vacuolar changes	0 to 1	0	0	0	1 to 2	0
brush border loss	0 to 1	0	1	1	3 to 4	1 to 2
cell detachment	0	0	1	0	1 to 2	0 to 1
nuclear condensation	0	0	0	0	0 to 1	0
Intratubular casts	0	0	1 to 3	0	2 to 4	0
Interstitial edema	0	0	0	0	1	1
Leukocyte infiltration	0	0	0	0	1 to 2	0
Capillary edema	0	0 to 1	1	1	1 to 2	0

Injury was scored in 10 randomly selected fields in each tissue section; n = 6 animals.

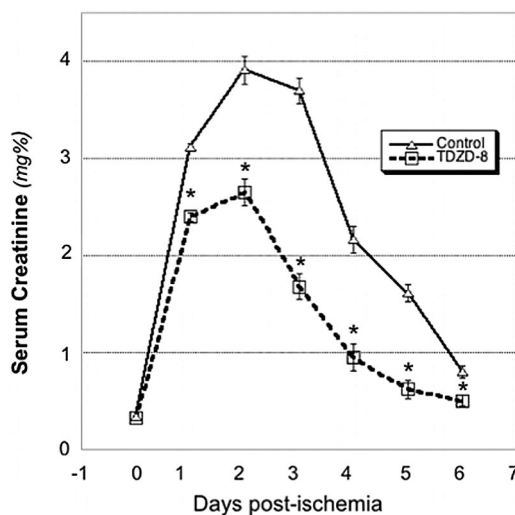


Figure 9. Effect of a GSK3β inhibitor on renal function after ischemia is shown. Serial serum creatinine levels in rats at baseline (Time 0) versus days 1 through 6 after 30 minutes of ischemia. *P < 0.03, vehicle versus TDZD-8; n = 6.

injury, GSK3β activation stimulated NF-κB followed by chemokine (C-C motif) ligand 5 known as CCL5 or RANTES expression, inflammation, and epithelial cell apoptosis.²⁷ Their report suggested that GSK3β exerts potent proinflammatory effects that contribute to organ injury. In addition, caspase activation itself promotes inflammation.⁶²

Given that GSK3β manipulation regulates apoptosis in renal epithelial cells *in vitro*, we propose that GSK3β-mediated apoptosis is an important cause of renal tubular injury and organ dysfunction after an ischemic insult (Figure 9). The observation that TDZD-8 inhibits ischemia-induced GSK3β kinase activity and Bax and caspase 3 activation, reduces tissue injury, and improves organ function is consistent with this interpretation. Furthermore, a single dose of TDZD-8, with a half-life of only a few hours,⁶⁵ not only inhibited premitochondrial events in the apoptosis cascade but also preserved organ function during prolonged recovery, suggesting that these

events are intimately linked. However, we cannot exclude the possibility that TDZD-8 affords protection by means other than its effect on GSK3β or that it alters survival in other cell types within the kidney. Importantly, we show that GSK3β is a readily modifiable cell survival signal that determines renal function after an acute ischemic insult.

CONCISE METHODS

Reagents

All reagents were purchased from Sigma (St. Louis, MO) unless otherwise indicated.

Antibodies

Rabbit polyclonal antibodies directed against phospho-serine⁹ GSK3β (inactive), total GSK3β, and active caspase-3 (Cell Signaling Technology, Beverly, MA), as well as Bcl2 (Calbiochem, San Diego, CA) were obtained. MAbs directed against fibrillarin (Abcam, Cambridge, MA) and β-actin were also obtained. Both total (5B7) and active (6A7) Bax antibodies were used (Trevingen, Gaithersburg, MD). Secondary antibodies conjugated to horseradish peroxidase (Jackson ImmunoResearch Laboratories, West Grove, PA) were used in combination with an enhanced chemiluminescence detection method (Pierce, Rockford, IL).

Cell Culture

A conditionally immortalized renal epithelial cell line derived from the mouse proximal tubule (BUMPT) was cultured as described previously.⁶⁶ These cells were maintained for up to 40 passages in DMEM with high-glucose medium (Life Technologies BRL, Carlsbad, CA) containing 10% bovine serum and 10% penicillin-streptomycin at 37°C in an incubator containing 5% CO₂.

GSK3β Overexpression

A WT and an established, constitutively active GSK3β construct (GSK3β-S9A) with a serine residue at position 9 was mutated to alanine^{21,42} and introduced into BUMPT cells using Lipofectamine 2000

(Invitrogen, Carlsbad, CA). The expression level of each mutant in cell lysates was confirmed by immunoblot analysis.

Inhibition of GSK3 β

Both molecular and pharmacologic maneuvers were used to inhibit GSK3 β . RNAi was achieved with short hairpin RNAs (shRNAs) expressed from the feline immunodeficiency virus (FIV)-based single-promoter pSIF1-H1-copGFP vector that produces shRNAs under control of H1 promoter and to express copGFP under the control of hCMV promoter (System Biosciences, Mountain View, CA). Each shRNA duplex was designed with a CTCCTGTCAGA loop. Three GSK3 β shRNA constructs were targeted to the following sequences: (1) 5'-GCTAGACCACTGTAACATAGT-3', (2) 5'-GCTGTGTGTTGGCTGAATTGT-3', and (3) 5'-GCGGGACCCAAATGTCAAAC-3'. Control shRNAs included the missense sequence, the nonspecific sequence, and the pSIF1-H1-copGFP vector. All constructs were sequenced before FIV packaging. HEK293TN cells were co-transfected with the lentiviral FIV-GSK3 β -shRNA constructs and packaging plasmids using Lipofectamine Plus (Invitrogen). Forty-eight hours after transfection, culture supernatant containing viral particles was harvested, clarified by filtration through a 0.45- μ m membrane filter (Nalge Nunc, Rochester, NY), and then concentrated by PEG precipitation. Fifteen hours after infection, the IU of each shRNA virus was estimated from the number of copGFP-expressing BUMPT cells. CopGFP-positive cells were then visualized and counted in 10 randomly selected fields using fluorescence microscopy. Cells were infected with virus (5 IU/cell) for 4 hours in the presence of 5 μ g/ml polybrene, and the medium containing virus particles was then exchanged for fresh medium. For infection with multiple viruses, cells were sequentially infected with a single virus for 4 hours followed by 4 hours of recovery, and the cycle was then repeated. Infected cells were cultured for 48 hours, and >90% of the population were GFP expression as assessed by direct visualization. The effect of each construct on total cell GSK3 β content was assessed by immunoblot analysis. TDZD-8, with an IC₅₀ of 4 μ M, inhibits GSK3 β by binding to its active site without altering the degree of serine⁹ phosphorylation.^{33,65} TDZD-8 dissolved in DMSO has been used to adequately and selectively inhibit GSK3 β in *in vitro* (10 μ M) as well and *in vivo* (1 mg/kg body wt, intravenously) studies.

Metabolic Stress

ATP content was reduced by incubating BUMPT cells in glucose-free medium that contained rotenone (10 μ M), a complex 1 electron transport inhibitor.^{67,68} This maneuver reduces ATP content to <2% of control within 10 minutes, a response that is comparable to that of cyanide⁶⁶ or antimycin.¹¹ For initiation of recovery, rotenone-free medium containing dextrose (5 mM) and heptanoic acid (2 mM) was added to bypass the rotenone-induced block of oxidative phosphorylation by supplying reducing equivalents to complex 2 in the electron transport chain.⁶⁸ As recently reported by our laboratory, metabolic stress for 1 to 2 hours causes apoptosis without appreciable epithelial cell necrosis.⁵¹

Hoechst Assay

For quantification of apoptosis, cells were grown on coverslips and then stained with Hoechst dye 33342 for 10 minutes at 37°C and fixed

with 2% paraformaldehyde for 30 minutes at 25°C.¹² The cells were mounted and then evaluated by ultraviolet immunofluorescence microscopy using established criteria for apoptosis.¹²

Cell Viability

Cell viability was assayed using a modified colorimetric technique that is based on the ability of live cells to convert 3-(4,5 dimethylthiazol)-2,5-diphenyl tetrazolium bromide (Promega, Madison WI), a tetrazolium compound, into purple formazan crystals as previously reported.²³ The number of surviving cells is expressed as a percentage of viable control cells detected at baseline. This assay accurately reflects cell survival in a model in which apoptosis is the primary cause of cell death.^{12,23,51,69}

Caspase 3 Activity

Enzyme activity was measured in cell lysates with a commercial kit (Enzolyte AFC Caspase-3 Assay Kit 71114; San Jose, CA) according to the manufacturer's instructions.

Immunoblot Analysis

Cells were harvested on ice in a lysis buffer containing 20 mM Tris-HCl, 140 mM NaCl, 1 mM sodium orthovanadate, 1 mM NaF, 1 mM dithiothreitol, 1 mM PMSF, 1 mM Na₄P₂O₇, 0.5% sodium deoxycholate; 0.4% NP-40, and a protease inhibitor cocktail (Boehringer Mannheim, Indianapolis, IN) at pH 7.5. Cell lysates were incubated for 15 min on ice and then centrifuged at 14,000 \times g for 15 minutes at 4°C for collection of the supernatant. The protein concentration of the supernatant was estimated with the Bio-Rad assay (Bio-Rad, Hercules, CA). Samples (20 to 50 μ g) were boiled in 1 \times SDS sample buffer (Boston Bioproducts, Ashland, MA) and separated using SDS-PAGE. Total and phosphorylated and kinase content were assessed in the same samples using immunoblots performed in parallel. β -Actin, total GSK3 β , β -catenin, and/or Bax was used as a loading control when appropriate. Data from each immunoblot were digitally acquired, and individual band densities were automatically reported.

Bax Knockdown

RNAi was used to silence Bax gene expression. Cells were plated in six- or 48-well culture plates with complete medium and allowed to grow for 24 hours to achieve 70% confluence. Cells were then transfected with an siRNA set composed of three target-specific 20- to 25-nt siRNAs that target Bax mRNA (Santa Cruz Biotechnology, Santa Cruz, CA). A nonspecific siRNA (Santa Cruz Biotechnology) that does not lead to the specific degradation of any known cellular mRNA served as a control. A mixture of OptiMEM and LipofectAMINE was incubated for 5 minutes at 25°C and was then incubated with the siRNA (100 pmol/well in six-well plates; 10 pmol/well in 48-well plates) for 20 minutes at 25°C. The siRNA mixture was then added to each well according to the manufacturer's protocol. The medium was changed 16 hours after transfection and incubated for 6 hours before infection with GSK3 β -S9A lentivirus (as detailed already). Bax gene silencing was confirmed by immunoblot analysis.

GSK3 β -Mediated Bax Serine¹⁶³ Phosphorylation

GSK3 was immunoprecipitated from 600 μ g of protein harvested from cell lysates as described already by constant mixing with 5 μ g of

anti-GSK3 β mAb and protein A–Sepharose for 2 hours at 4°C. The resulting immunoprecipitates were washed three times in kinase assay wash buffer (20 mM HEPES, 20 mM β -glycerophosphate, and 1 mM EDTA at pH 7.5) and then resuspended in 35 μ l of kinase assay mixture containing 4 mM MOPS, 2.5 mM β -glycerophosphate, 1 mM EGTA, 0.4 mM EDTA, 4 mM MgCl₂, 0.05 mM dithiothreitol, 40 ng/ μ l BSA, 50 μ M ATP, 2.5 μ M cAMP-dependent protein kinase inhibitor peptide (IP₂₀), and 100 μ M [γ -³²P]ATP (PerkinElmer Life Sciences, Boston, MA) at pH 7.2 and a synthetic Bax-specific peptide that flanks serine¹⁶³ substrate WEGLLSYFGTP (400 ng/ μ l). For each assay, 30 μ l of this mixture was used in the presence or absence of TDZD-8 (10 μ M). The assay was terminated after 30 minutes of incubation at 30°C by spotting onto P81 ion-exchange paper. The paper was washed four times in 0.6% phosphoric acid, and bound radioactivity was quantified by scintillation counting. A commercially available GSK3 β phosphorylation substrate (Upstate Biotech, Lake Placid, NY) was used to optimize the reaction conditions.

In Vivo Renal Ischemia

Male Sprague-Dawley rats that weighed 250 to 270 g each were anesthetized with thiopental sodium (55 mg/kg, intraperitoneally), a mid-line laparotomy was performed, and the left kidney was removed. A nontraumatic vascular clamp was placed on the right renal artery for 30 minutes, an insult that produces severe, reversible acute renal failure.⁷⁰ Controls were subjected to either uni-nephrectomy (Unx or Unx CTL) or sham renal ischemia using a nonocclusive ligature without nephrectomy (Non-Unx). After transient ischemia, the clamp was removed and perfusion was confirmed by visual inspection. At serial time points after ischemia, the right kidney was harvested; the capsule was removed; the renal cortex was bluntly dissected into cortex and outer and inner medullas; and the tissues were prepared for immunoblot analysis to assess the content of Bcl2, active and total Bax, phospho- and total GSK3 β , phospho and total β -catenin, and active caspase 3. Renal histology was examined in stained in 5- μ m tissue sections. In a subsequent study, animals received either a single intravenous dose of 1 mg/kg TDZD-8 dissolved in DMSO 1 hour before ischemia³² or vehicle alone (control). These observations provided a physiologic basis for administering TDZD-8, a GSK3 β inhibitor with an IC₅₀ of 4 μ M,³³ as a single dose (1 mg/kg, intravenously) before ischemia, as reported by Dugo *et al.*³² in a rodent sepsis model. An observer who was blinded to the experimental conditions (R.B.) assessed the degree of tissue injury using a standardized, semiquantitative scoring scheme with four levels of severity for each of nine distinct categories on kidney sections stained with hematoxylin and eosin.^{71,72} Serum creatinine in tail-vein samples was measured at baseline and daily for 6 days after ischemia using a commercial kit (BioAssay Systems, Hayward, CA). All procedures were performed in adherence with the National Institutes of Health Guide for the Care and Use of Laboratory Animals.

Statistical Analysis

Comparison of two groups was performed using a two-tailed *t* test. Differences among three or more groups were assessed using two-tailed ANOVA (Excel, Microsoft Corp., Redmond, WA). An experimental result was considered significant at *P* < 0.05.

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DISCLOSURES

None.

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See related editorial, "GSK3β Plays Dirty in Acute Kidney Injury," on pages 199–200.