cGMP-Selective Phosphodiesterase Inhibitors Stimulate Mitochondrial Biogenesis and Promote Recovery from Acute Kidney Injury

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ABSTRACT

Recent studies demonstrate that mitochondrial dysfunction is a mediator of acute kidney injury (AKI). Consequently, restoration of mitochondrial function after AKI may be key to the recovery of renal function. Mitochondrial function can be restored through the generation of new, functional mitochondria in a process called mitochondrial biogenesis (MB). Despite its potential therapeutic significance, very few pharmacological agents have been identified to induce MB. To examine the efficacy of phosphodiesterase (PDE) inhibitors (PDE3: cAMP and gMP activity; and PDE4: cAMP activity) in stimulating MB, primary cultures of renal proximal tubular cells (RPTCs) were treated with a panel of inhibitors for 24 hours. PDE3, but not PDE4, inhibitors increased the FCCP-uncoupled OCR in primary cultures of renal proximal tubular cells (RPTCs) were treated with a panel of inhibitors for 24 hours. PDE3, but not PDE4, inhibitors increased the FCCP-uncoupled oxygen consumption rate (OCR), a marker of MB. Exposure of RPTCs to the PDE3 inhibitors, cilostamide and trequinsin, for 24 hours increased peroxisome proliferator-activated receptor γ coactivator-1α, and multiple mitochondrial electron transport chain genes. Cilostamide and trequinsin also increased mRNA expression of mitochondrial genes and mitochondrial DNA copy number in mice renal cortex. Consistent with these experiments, 8-Br-cGMP increased FCCP-uncoupled OCR and mitochondrial gene expression, whereas 8-Br-cAMP had no effect. The cGMP-specific PDE5 inhibitor sildenafil also induced MB in RPTCs and in vivo in mouse renal cortex. Treatment of mice with sildenafil after folic acid–induced AKI promoted restoration of MB and renal recovery. These data provide strong evidence that specific PDE inhibitors that increase cGMP are inducers of MB in vitro and in vivo, and suggest their potential efficacy in AKI and other diseases characterized by mitochondrial dysfunction and suppressed MB.

Introduction

Mitochondrial dysfunction is increasingly recognized as an important pathophysiological mediator of a variety of disease states, including neurodegeneration, cardiovascular disease, metabolic syndrome, and acute organ injury (Choumar et al., 2011; Pundik et al., 2012; Andreux et al., 2013; Bayeva et al., 2013; Cheng and Ristow, 2013; Cooper, 2013; Hwang, 2013; Yan et al., 2013). Mitochondrial dysfunction is an established component of the pathogenesis of acute kidney injury (AKI) and a cause of renal tubular dysfunction and cell death (Jassem et al., 2002; Jassem and Heaton, 2004; Hall and Unwin, 2007; Weinberg, 2011; Venkatachalam and Weinberg, 2012). Our group has demonstrated persistent disruption of mitochondrial homeostasis and inhibition of mitochondrial biogenesis (MB) after ischemia-reperfusion (I/R), rhabdomyolysis-induced AKI (Funk and Schnellmann, 2012), and folic acid (FA)–induced AKI (unpublished data).

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ABBREVIATIONS: AKI, acute kidney injury; ATP5β, ATP synthase subunit β; COX1, cytochrome c oxidase subunit 1; CREB, cAMP-response element-binding protein; DOI, 1-(2,5-dimethoxy-4-iodophenyl)-2-aminopropane hydrochloride; eNOS, endothelial nitric-oxide synthase; FA, folic acid; FCCP, carbonyl cyanide 4-(trifluoromethoxy)phenylhydrazone; I/R, ischemia reperfusion; KIM-1, kidney injury molecule-1; MB, mitochondrial biogenesis; mtDNA, mitochondrial DNA; ND1, NADH dehydrogenase 1; ND6, NADH dehydrogenase 6; NDUFβ, NADH dehydrogenase [ubiquinone] 1β subcomplex subunit 8; NO, nitric oxide; NRF1, nuclear respiratory factor 1; OCR, oxygen consumption rate; PDE, phosphodiesterase; PGC-1α, peroxisome proliferator-activated receptor γ coactivator-1α; PKA, protein kinase A; qPCR, quantitative real-time polymerase chain reaction; Ro 20-1724, 4-[3-butoxy-4-methoxyphenyl]methyl-2-imidazolidone; ROS, reactive oxygen species; RPTC, renal proximal tubular cell; SIRT1, silent mating type information regulation 2 homolog 1; SRT1720, N-[2-[3-(piperazin-1-ylmethyl)[imidazo][1,2-b][1,3]thiazol-6-yl][phenyl]quinoxaline-2-carboxamide; Tfam, mitochondrial transcription factor A.
Restoration of mitochondrial number and function is thought to be required for recovery from AKI due to the high energy requirements of tissue repair. These data provide support for development of pharmacological agents that induce MB for treatment of AKI and other pathologies characterized by mitochondrial dysfunction.

Mitochondria are dynamic organelles that are continuously regenerated through the processes of biogenesis, mitophagy, fission, and fusion (Brooks et al., 2009; Shaw and Winge, 2009; Cho et al., 2010; Funk and Schnellmann, 2012; Kabli and Gustafsson, 2012). MB is the assembly of new mitochondria from existing mitochondria, occurring under basal conditions to replace damaged mitochondria, but is rapidly induced in response to both physiologic and pathophysiologic stimuli, including sepsis, exercise, fasting, hypoxia, and cellular injury (Puigserver and Spiegelman, 2003; Tran et al., 2011; Kang and Li Ji, 2012; Wenz, 2013). The primary regulator of MB is the transcriptional coactivator peroxisome proliferator-activated receptor γ coactivator 1α (PGC-1α). PGC-1α exerts its functions by activating the transcription factors, nuclear respiratory factors 1 and 2 (Nrf1 and Nrf2). Nrf1 controls the expression of mitochondrial transcription factor A (Tfam), which regulates the transcription of mitochondrial DNA (mtDNA) (Puigserver et al., 1998; Wu et al., 1999; Scarpulla, 2008; Scarpulla et al., 2012). PGC-1α is enriched in tissues with high metabolic demand, including heart, muscle, and kidneys (Liang and Ward, 2006). The ability of PGC-1α to respond to a variety of stimuli and its importance in cellular bioenergetics make it an ideal target for pharmacological intervention in disease states characterized by mitochondrial dysfunction.

Despite the promise of PGC-1α and MB as a therapeutic target, there is a paucity of pharmacological agents capable of stimulating PGC-1α expression and activity. Activators of silent mating type information regulation 2 homolog 1 (SIRT1)—including isoflavones, resveratrol, and N-[2-[3-(piperazin-1-ylmethyl)imidazo[2,1-b][1,3]thiazol-6-yl]phenyl]quinoxaline-2-carboxamide (SRT1720)—have been demonstrated to induce PGC-1α and promote increased mitochondrial number and function (Rasbach and Schnellmann, 2008; Funk et al., 2010; Menzies et al., 2013). Our laboratory also identified the 5-hydroxytryptamine type 2 agonist, 1-(2,5-dimethoxy-4-iodophenyl)-2-amino propane hydrochloride (DOI), and the β2-adrenergic receptor agonist, formoterol, as potent inducers of PGC-1α and MB (Rasbach et al., 2010; Wills et al., 2012). Stimulation of MB after injury accelerates recovery of cellular morphology and function (Rasbach and Schnellmann, 2007; Funk et al., 2010; Rasbach et al., 2010). These data demonstrate the importance of MB in recovery of renal tubular epithelial cells after injury and suggest that agents that stimulate MB could serve as viable therapies after AKI.

Because of the importance of the cAMP/protein kinase A (PKA/cAMP-response element-binding protein (CREB) axis in PGC-1α regulation, drugs that increase cellular cAMP levels may induce MB. The β2-adrenergic signaling cascade, which upon activation increases intracellular cAMP through Gs-mediated activation of adenylyl cyclase, was shown to regulate oxidative metabolism and energy expenditure (Tadaishi et al., 2011; Muller et al., 2013). Formoterol induces MB in renal proximal tubular cells (RPTCs), and mice treated with formoterol demonstrated increased mitochondrial gene expression and mtDNA copy number in renal cortex and heart (Wills et al., 2012). cGMP levels have also been shown to regulate PGC-1α expression and MB. Pharmacologically induced generation of nitric oxide (NO) via endothelial nitric oxide synthase (eNOS) and subsequent NO-dependent activation of guanylyl cyclase led to MB in U937, L6, and PC12 cells. (Nisoli et al., 2004).

Both cAMP and cGMP levels are tightly regulated through cleavage to AMP and GMP, respectively, by a class of enzymes called cyclic nucleotide phosphodiesterases (PDEs). The PDE superfamily consists of 11 families differing in tissue distribution, regulation, and substrate affinity (e.g., cAMP versus cGMP) (Francis et al., 2011). Potent, selective inhibitors of nearly all family members are available (Bender and Beavo, 2006). Inhibition of PDEs would serve as a novel and potentially efficacious drug target to induce MB. As such, we studied inhibitors of PDE3, PDE4, and PDE5 for their ability to induce MB in the kidney and promote recovery from FA-induced AKI.

Materials and Methods

Reagents. Cilostamide, trequinsin, 3-(4-[(iminomethylene)oxy]phenoxy)-N-(2-methylpropyl)aniline (0.5 M), allowing for the measurement of uncoupled OCR.

Isolation and Culture of Proximal Tubules. Female New Zealand white rabbits (1.5–2.0 kg) were purchased from Charles River Laboratories (Wilmington, MA). RPTCs were isolated using the iron oxide perfusion method previously described (Nowak and Schnellmann, 1995). For respirometry experiments, cells were plated on 100-mm culture-grade Petri dishes at 37°C in a 5% CO2/95% air environment. Dishes were continuously swirled on an orbital shaker at 80 rpm. Cell culture media consisted of a 1:1 mixture of Dulbecco’s modified Eagle’s essential medium and Ham’s F-12 (lacking glucose, phenol red, and sodium pyruvate; Invitrogen, Carlsbad, CA), supplemented with HEPES (15 mM), glutamine (2.5 mM), pyridoxine (1 μM), sodium bicarbonate (15 mM), and lactate (6 mM). Hydrocortisone (50 nM), selenium (5 ng/ml), human transferrin (5 μg/ml), bovine insulin (10 nM), and L-ascorbic acid 2-phosphate (50 μM) were added daily to fresh culture media. After 3 days of culture, dedifferentiated RPTCs were trypsinized and replated on XF-96 polystyrene cell culture microplates (Seahorse Bioscience, North Billerica, MA) at a concentration of 18,000 cells per well. Cells were maintained at 37°C for an additional 2 days before experimentation (Beeson et al., 2010). For all other RPTC experiments, cells were plated and cultured in 35-mm dishes in the above-described media. Experiments were performed on the sixth day after plating when cells had formed a confluent monolayer. RPTCs were treated with various compounds for 24 hours unless otherwise noted.

Oxygen Consumption. The oxygen consumption rate (OCR) of RPTCs was measured using the Seahorse Bioscience XF-96 Extracellular Flux Analyzer according to a previously described protocol (Beeson et al., 2010). Each assay plate was treated with vehicle control (dimethylsulfoxide <0.5%), and increasing concentrations of the compounds of interest. Basal OCR was measured before injection of carbonyl cyanide 4-(trifluoromethoxy)phenylhydrazone (FCCP) (0.5 μM), allowing for the measurement of uncoupled OCR.
Testing of Compounds in C57BL/6 Mice. Male C57BL/6 mice (aged 6–8 weeks) were obtained from the National Institutes of Health National Cancer Institute (Bethesda, MD). Mice were housed individually in a temperature-controlled room under a 12-hour light/dark cycle. Mice were randomly assigned to saline control, cisteamine (0.3 or 3 mg/kg), trequinsin (0.3 or 3 mg/kg), or sildenafil (0.3 or 3 mg/kg) treatment groups. Mice were given a single intraperitoneal injection of saline or compound at the above-described doses. Mice were euthanized by CO2 asphyxiation followed by cervical dislocation 24 hours after treatment. Kidneys were removed and preserved by flash-freezing in liquid nitrogen. Tissues were stored at −80°C for later analysis.

**FA Animal Model.** Male C57BL/6 mice (aged 8–10 weeks) were given a single intraperitoneal injection of 250 μg/kg FA dissolved in 250 mM sodium bicarbonate or saline control based on previous literature (Doi et al., 2006). Mice were injected with sildenafil (0.3 mg/kg) or diluent every 24 hours beginning 1 day after FA injection. Mice were euthanized at 7 days via isoflurane asphyxiation and cervical dislocation. Kidneys were removed and preserved by flash-freezing.

**Quantitative Real-Time Polymerase Chain Reaction.** Total RNA was extracted from RPTCs or renal cortex samples using TRIzol reagent (Invitrogen) according to the manufacturer’s protocol. cDNA was synthesized via reverse transcription using the iScript Advanced cDNA synthesis kit (Bio-Rad, Hercules, CA) with 2 μg of RNA. Quantitative real-time polymerase chain reaction (qPCR) analysis was performed with cDNA using SsoAdvanced SYBR Green Supermix (Bio-Rad, Hercules, CA) with 2

**mtDNA Content.** Real-time PCR was used to determine the relative quantity of mtDNA in both RPTC and mouse renal cortical tissue samples. After treatment, DNA was extracted from cells or tissue using the DNeasy Blood and Tissue Kit (QIAGEN, Valencia, CA) and 5 ng of cellular DNA was used for qPCR. For RPTC samples, mitochondrial-encoded ND6 was used to measure mitochondrial copy number and was normalized to nuclear-encoded tubulin expression. For renal cortex, ND1 was used as the mitochondrial gene and expression was normalized to nuclear-encoded β-actin expression.

**cAMP and cGMP Enzyme-Linked Immunosorbent Assay (ELISA).** RPTCs in 35-mm dishes were treated with vehicle control (dimethylsulfoxide) or the compound of interest for 20 minutes. RPTCs were then harvested according to the manufacturer’s protocol and cAMP or cGMP levels were measured using a commercially available enzyme-linked immunosorbent assay kit (Cayman Chemical, Ann Arbor, MI).

**Tissue ATP Levels.** ATP was isolated from renal cortical tissue via phenol-Tris-EDTA extraction as previously described (Chida et al., 2012). In brief, freshly prepared tissue was homogenized in 3.0 ml ice-cold Tris-EDTA saturated phenol. One milliliter of the homogenate was combined with 200 μl chloroform and 150 μl of deionized water and vortexed and centrifuged at 10,000g for 5 minutes at 4°C. An aliquot from the supernatant was diluted 200-fold in deionized water, and ATP levels were measured using a luciferin-luciferase–based ATP determination kit (Invitrogen).

**Statistical Analysis.** Data are presented as the mean ± S.E.M. Simple comparisons were performed using the t test. Multiple comparisons were subjected to one-way analysis of variance followed by the Newman–Keuls test, with P < 0.05 considered to be a statistically significant difference between means. RPTCs isolated from a single rabbit represented an individual experiment (n = 1) and were repeated until n ≥ 4 was obtained. Mouse studies were repeated until n ≥ 3 was obtained.

**Results**

**PDE3 Inhibitors, but not PDE4 Inhibitors, Increase FCCP-Uncoupled OCR in RPTCs.** We treated RPTCs in XF-96 plate cultures with the PDE3 inhibitors cisteamine or trequinsin, the PDE4 inhibitors (R)-(−)-rolipram or Ro 20-1724, or vehicle control for 24 hours. PDE3 hydrolyzes both cAMP and cGMP to their noncyclic forms, AMP and GMP, whereas PDE4 specifically hydrolyzes cAMP to AMP (Francis et al., 2011). FCCP-OCR increased in RPTCs compared with vehicle control after 24-hour exposure to cisteamine (25–100 nM) and trequinsin (30–100 nM) (Fig. 1, A and B), but no significant changes were observed in FCCP-OCR after treatment with (R)-(−)-rolipram (0.5–50 μM) or Ro 20-1724 (5–20 μM) (Fig. 1, C and D). These data suggest a functional selectivity for the MB response between PDE3 and PDE4 inhibition in RPTCs.

**PDE3 Inhibitors Induce MB in RPTCs.** To validate that the increased FCCP-OCR observed in RPTCs after treatment with PDE3 inhibitors was due to MB, mRNA levels for PGC-1α, the mitochondrial-encoded complex I protein ND6, and the nuclear-encoded complex I protein NDUF8 were measured via qPCR. Gene expression was normalized to tubulin. PGC-1α levels increased versus control after treatment with cisteamine (1.8-fold) or trequinsin (2.5-fold) (Fig. 2). In addition, mRNA expression of mitochondrial-encoded ND6 and the nuclear-encoded NDUF8 mitochondrial proteins were increased versus control with cisteamine (1.5- and 2.2-fold, respectively) and trequinsin (1.8- and 2.4-fold, respectively). These data provide strong evidence that inhibition of PDE3 causes functional MB in RPTCs.

**Increased cGMP, but Not cAMP, Induces MB in RPTCs.** To examine the functional selectivity of PDE3 and PDE4 inhibitors under conditions that induce MB, RPTCs were treated with the PDE3 inhibitors cisteamine and trequinsin, the PDE4 inhibitor (R)-(−)-rolipram, or vehicle control for a period of 20 minutes. Sildenafil, a specific inhibitor of PDE5 (cGMP-specific PDE), was included as a control. Both cAMP and cGMP levels increased in response to cisteamine and trequinsin compared with vehicle control, whereas cAMP only increased in RPTCs treated with rolipram (Fig. 3, A and B). RPTCs treated with sildenafil resulted in increased cGMP, but not cAMP. These data agree with the classic mechanisms of PDE3 (hydrolyzes both cAMP and cGMP), PDE4 (hydrolyzes only cAMP), and PDE5 (hydrolyzes only cGMP) (Bender and Beavo, 2006). The inability of rolipram and other PDE4 inhibitors tested to induce MB suggests that cGMP may be the primary mediator of MB in RPTCs.

**To test this hypothesis, we treated RPTCs with the cell-permeable cyclic nucleotide analogs, 8-Br-cAMP and 8-Br-cGMP for a 24-hour period. RPTCs treated with 8-Br-cGMP (10–300 μM) showed an approximately 20% increase in FCCP-uncoupled OCR at all concentrations tested, whereas treatment with 8-Br-cAMP resulted in no change (Fig. 3C). To validate that this increase in FCCP-OCR is due to stimulation of MB, mRNA expression of PGC-1α, ND6, and NDUF8 was measured by qPCR. RPTCs treated with 8-Br-cGMP had elevated mRNA levels of PGC-1α (2.2-fold), ND6 (1.7-fold),
and NDUF8 (1.9-fold). 8-Br-cAMP had no effect on mitochondrial gene expression (Fig. 3D). Furthermore, to test the ability of a PDE5 inhibitor to induce MB in vitro, RPTCs were treated with sildenafil for 24 hours (1 nM–100 nM) and FCCP-OCR was measured using the Seahorse XF96. RPTCs treated with sildenafil showed an approximately 20% increase in FCCP-uncoupled OCR versus controls (Fig. 4A) at 10 and 100 nM. To validate that the increase in respiration was due to MB, mRNA levels of PGC-1α, ND6, and NDUF8 were measured and were found to increase 1.8-, 2.0-, and 1.5-fold, respectively (Fig. 4B).

PDE3 Inhibitors Induce MB in Mouse Renal Cortex. In kidneys of cilostamide-treated mice, PGC-1α was induced 2- and 2.7-fold at doses of 0.3 and 3 mg/kg, respectively. Trequinsin induced PGC-1α 2.7- and 2.8-fold in the kidney at doses of 0.3 and 3 mg/kg. mRNA expression of the nuclear-encoded mitochondrial genes NDUF8 and ATPSβ both increased greater than 2-fold in kidneys of mice treated with either cilostamide or trequinsin at 0.3 or 3 mg/kg (Fig. 5, A and B). The mitochondrial-encoded mitochondrial genes ND1 and cytochrome c oxidase subunit I (COX1) increased in the kidneys of these mice. The mtDNA copy number was also increased in the kidneys of mice treated with cilostamide at 0.3 mg/kg, whereas mice treated with 3 mg/kg cilostamide had no effect (Fig. 5C). Trequinsin increased mtDNA copy number in the kidneys 1.6- and 2-fold at doses of 0.3 and 3 mg/kg, respectively (Fig. 5D). These data provide strong evidence that pharmacological inhibition of PDE3 induces MB in the kidney of naïve mice.

Fig. 1. PDE3 inhibitors, but not PDE4 inhibitors, increase FCCP-induced uncoupled mitochondrial respiration in RPTCs. RPTCs were treated with cilostamide (A), trequinsin (B), (R)-(-)-rolipram (C), or Ro 20-1724 (D) for 24 hours. FCCP-uncoupled mitochondrial respiration was measured using the Seahorse XF-96 instrument. Data are presented as the mean ± S.E.M. (n ≥ 3). *P < 0.05 vs. vehicle control.

Fig. 2. PDE3 inhibitors cilostamide or trequinsin induce mitochondrial protein gene expression in RPTCs. RPTCs were exposed to cilostamide (25 nM) or trequinsin (30 nM) for 24 hours and evaluated for changes in mRNA expression of PGC-1α, ND6, and NDUF8 relative to dimethylsulfoxide controls. Data are presented as the mean ± S.E.M. (n = 4). *P < 0.05 vs. vehicle control.
Sildenafil Induces MB in Mouse Renal Cortex. The selectivity of the MB response for cGMP in RPTCs indicates that inhibitors of cGMP-specific PDEs, such as PDE5, may in fact be a better therapeutic target and could eliminate off-target effects due to the accumulation of cAMP. PDE5 inhibitors also have a much more favorable safety protocol than PDE3 inhibitors, particularly for extended administration (Cruickshank, 1993).

To determine whether PDE5 inhibition is capable of inducing MB in the kidney in vivo, mice were given a single intraperitoneal injection of sildenafil (0.3 or 3 mg/kg) or saline control. Mice were euthanized and kidneys were harvested.

Fig. 3. PDE inhibitor–induced increases in cGMP, but not cAMP, stimulate MB in RPTCs. cAMP (A) and cGMP (B) levels were measured in RPTCs by enzyme-linked immunosorbent assay 20 minutes after treatment with dimethylsulfoxide, cilostamide (25 nM), trequinsin (30 nM), rolipram (0.5 μM), or sildenafil (10 nM). (C) FCCP-uncoupled mitochondrial respiration was measured using the Seahorse XF-96 instrument after 24-hour treatment with 8-Br-cAMP or 8-Br-cGMP. (D) RPTCs were exposed to 8-Br-cAMP (10 μM) or 8-Br-cGMP (10 μM) for 24 hours and evaluated for changes in mRNA expression of PGC-1α, ND6, and NDUFβ8 relative to dimethylsulfoxide controls. Data are presented as the mean ± S.E.M. (n = 3). *P < 0.05 vs. vehicle control.

Fig. 4. The PDE5 inhibitor sildenafil stimulates MB in RPTCs. Sildenafil increases FCCP-uncoupled mitochondrial respiration at various doses (A) and mitochondrial gene expression at 10 nM (B) in RPTCs. mRNA expression of PGC-1α, ND6, and NDUFβ8 is presented as the mean ± S.E.M. of at least three biologic replicates. *P < 0.05 vs. vehicle control.
24 hours after treatment. mRNA levels of PGC-1α, NDUFB8, ND1, ATPβ, and COX1 were measured by qPCR. All mitochondrial genes, except for COX1 and ATPSβ, were increased in mice treated with 3 mg/kg sildenafil versus saline-treated animals (Fig. 6A). mtDNA copy number was assessed by qPCR in kidneys of sildenafil-treated mice and was found to increase 1.6-fold in mice treated with 0.3 mg/kg sildenafil. No change in mtDNA copy number was observed in mice treated with 3 mg/kg sildenafil (Fig. 6B).

To assess whether sildenafil-induced MB increased mitochondrial function in the kidney cortex, we measured ATP levels. ATP levels increased 32% in mice treated with 0.3 mg/kg sildenafil compared with control mice (Fig. 6C). These data strongly support our hypothesis that PDE5 inhibitors induce MB and mitochondrial function in vitro and in vivo.

**Sildenafil Promotes Recovery of MB and Renal Function after FA-Induced AKI.** To test the hypothesis that sildenafil-induced MB will accelerate recovery of mitochondrial and renal function after AKI, we induced AKI by injecting FA and then treated these mice with sildenafil or vehicle once daily starting at 24 hours after injury for 6 days. mRNA expression of COX1 and Tfam were reduced to 27 and 36% of control, respectively, in FA-treated mice receiving vehicle control at 6 days. Sildenafil-treated FA mice demonstrated a 1.6-fold increase in mRNA COX1 expression to 43% of control mice, and a 1.4-fold increase in Tfam expression to 50% of control (Fig. 7A). mtDNA copy number was reduced to 36% of animals receiving FA alone, and treatment with sildenafil caused an approximately 2-fold induction to 63% of control (Fig. 7B). These data demonstrate that sildenafil can induce MB in a model of AKI.

To examine whether MB promoted renal recovery, kidney injury molecule-1 (KIM-1), a specific marker of tubular injury, was measured in renal cortex. KIM-1 levels were markedly increased (approximately 6-fold) in FA-treated animals compared with control animals and treatment of FA mice with sildenafil restored KIM-1 expression to control levels (Fig. 7, C and D). These data demonstrate that sildenafil promotes renal recovery with its induction of mitochondrial gene expression and mtDNA copy number.

**Discussion**

Mitochondria are highly regulated organelles whose function is tightly linked to the metabolic demands and health of a cell (Brooks et al., 2009; Shaw and Winge, 2009; Funk and Schnellmann, 2012; Kubli and Gustafsson, 2012). Mitochondrial function is necessary for normal cell and tissue function, and is critical in energy-dependent repair processes. A wide array of disease states are characterized by mitochondrial dysfunction, including diabetes, neurodegenerative disease, traumatic brain injury, and acute organ injury (Lifshitz et al., 2004; Pundik et al., 2012; Cheng and Ristow, 2013; Hwang, 2013; Yan et al., 2013). I/R and drug/toxicant-induced renal
injury demonstrate mitochondrial dysfunction and suppression of MB, and recovery of renal function is tightly linked to the restoration of mitochondrial number and function (Funk and Schnellmann, 2012). This suggests that development of therapies capable of inducing MB may have great potential in the treatment of a broad range of disease states.

Despite strong evidence supporting mitochondria as a therapeutic target, there are very few drugs/chemicals available that promote mitochondrial function or MB. Many of the agents that are available suffer from lack of specificity, low potency, or poor toxicity profiles. There is a clinical need to develop new pharmacological agents or identify existing therapeutics that induce MB. Because of the role of cyclic nucleotides as regulators of PGC-1α, in this study, we sought to determine the efficacy of various classes of PDE inhibitors at stimulating MB.

The cAMP/PKA/CREB signaling cascade is a well characterized regulator of PGC-1α expression and activity (Fernandez-Marcos and Auwerx, 2011). Increases in intracellular cAMP levels cause activation of PKA and subsequent phosphorylation and activation of CREB, an important transcriptional regulator of PGC-1α. Induction of cAMP levels in the cell occurs after activation of various G protein-coupled receptors. Our laboratory recently identified the β2-adrenergic agonist, formoterol, as a potent inducer of MB in the kidney and heart (Wills et al., 2012). β-agonism was previously shown to induce PGC-1α in skeletal muscle of treated mice (Miura et al., 2007). In addition, exercise-induced MB can be blocked by treatment with β-receptor antagonists, propranolol and ICI-118,551. cAMP levels in the cell are controlled both by the rate of synthesis and the rate of turnover by cyclic nucleotide PDEs. Therefore, inhibition of PDEs that hydrolyze cAMP may serve as a viable intervention to induce MB.

To test this hypothesis, we screened a panel of PDE3, PDE4, and PDE5 inhibitors using a phenotypic respirometric assay. FCCP-uncoupled OCR was used as a marker of increased energetic capacity and MB. Interestingly, PDE3 and PDE5 inhibitors increased FCCP-uncoupled OCR in RPTCs, whereas none of the PDE4 inhibitors tested caused an increase (Figs. 1 and 4). To further probe the functional selectivity of PDE3, PDE4, and PDE5 inhibition in promoting MB, cAMP and cGMP levels were measured in RPTCs after treatment with the PDE3 inhibitors cilostamide and trequinsin, the PDE4 inhibitor rolipram, or the PDE5 inhibitor sildenafil. PDE3 inhibition led to increases in levels of both cAMP and cGMP in RPTCs, whereas rolipram increased only cAMP levels and sildenafil increased only cGMP levels (Fig. 3). These data correspond with the classic substrate affinities of the various PDE family members: PDE3 hydrolyzes both cAMP and cGMP with nearly equal affinity, PDE4 specifically hydrolyzes cAMP, and PDE5 specifically hydrolyzes cGMP (Bender and Beavo, 2006; Francis et al., 2011). Finally, 8-Br-cGMP increased FCCP-uncoupled OCR in the respirometric assay and increased mRNA expression of mitochondrial genes after 24-hour treatment. 8-Br-cAMP had no effect on respiration of mitochondrial gene expression in RPTCs. This multipronged approach strongly supports our hypothesis that cGMP, rather than cAMP, is important for regulation of MB in renal tubules.

cGMP was previously demonstrated to induce MB through the eNOS/NO soluble guanylate cyclase/cGMP signaling cascade. Nisoli et al. (2004) showed that long-term administration of NO mimetics, guanyyl cyclase activators, or 8-Br-cGMP...
increased mRNA expression of mitochondrial genes, mtDNA copy number, mitochondrial respiration, and ATP levels in multiple cell lines. eNOS-deficient mice have a reduction in metabolic rate and accelerated weight gain, which has been correlated with reduced mitochondrial content and function (Nisoli et al., 2003).

Both the PDE3 inhibitors cilostamide or trequinsin (0.3–3 mg/kg) and the PDE5 inhibitor sildenafil (0.3–3 mg/kg) when administered to naïve mice induced renal cortical mRNA expression of PGC-1α, nuclear-encoded mitochondrial genes (NDUFB8 and ATPSβ), and mitochondrial-encoded mitochondrial genes (ND1 and COX1). mtDNA copy number was also increased in the renal cortex of these mice (Figs. 5 and 6). Sildenafil increased the number of functional mitochondria in the renal cortex as evidenced by a significant increase in tissue ATP levels (Fig. 6). These data confirm that PDE3 and PDE5 inhibitors are capable of inducing MB both in vitro in RPTCs and in vivo in mouse kidney.

Cyclic nucleotides including both cAMP and cGMP were shown to be activators of signaling pathways promoting MB in various model systems (Nisoli et al., 2003, 2004; Tadaishi et al., 2011; Muller et al., 2013). Acute ex vivo administration of the PDE5 inhibitor vardenafil to human skeletal muscle stimulated MB as evidenced by increases in mitochondrial gene expression and mtDNA copy number (De Toni et al., 2011). This is the first report of pharmacological induction of MB in vivo by inhibition of either PDE3 or PDE5, and could represent a novel use for these classes of compounds. Despite the evidence of their role in MB, these compounds have yet to be evaluated as potential therapies for mitochondrial damage and dysfunction.

Previous studies reported the ability of various classes of PDE inhibitors to protect against AKI. Pretreatment with rolipram, a specific PDE4 inhibitor, blunted I/R-induced renal dysfunction in rat kidney and reduced oxidative damage (Mammadov et al., 2012). Sildenafil was shown to be protective in cisplatin-induced AKI, whereas tadalafl, a long-acting PDE5 inhibitor, protected against early I/R injury in rats (Lee et al., 2009; Sohotnik et al., 2013). However, limitations of these studies have been the lack of a clear mechanism for the renoprotective effects and the use of pretreatment protocols. To address these issues, we examined the ability of sildenafil to promote recovery from FA-induced AKI by administering the drug 24 hours after induction of injury, and examined the effects of FA and sildenafil on both renal and mitochondrial function. Sildenafil promoted recovery mitochondrial gene expression (i.e., COX1 and Tfam) and mtDNA copy number. In addition, renal KIM-1 expression was reduced in sildenafil-treated mice, indicating an enhanced recovery from the renal injury. These results demonstrate that sildenafil accelerates recovery from AKI by activating MB pathways.

Our results indicate that PDE inhibitors that are capable of increasing tissue levels of cGMP, including sildenafil, are promising treatments for diseases characterized by mitochondrial dysfunction and suppression of MB, including AKI.