

1 **Elucidation of cGMP-Dependent Induction of Mitochondrial Biogenesis Through Protein**  
2 **Kinase G and p38 MAPK in the Kidney**

3 **Pallavi Bhargava<sup>1\*</sup>, Jaroslav Janda<sup>1\*</sup>, and Rick G. Schnellmann<sup>1,2</sup>**

4 \* These authors contributed equally to this work

5 <sup>1</sup>*Department of Pharmacology and Toxicology, College of Pharmacy, University of Arizona,*  
6 *Tucson, Arizona;* <sup>2</sup>*Southern Arizona Veterans Affairs Health Care System, Tucson, Arizona*

7 Running Title: cGMP Induced Mitochondrial Biogenesis in the Kidney

8 Corresponding Author: Dr. Rick G. Schnellmann, PhD

9 University of Arizona

10 Department of Pharmacology and Toxicology

11 College of Pharmacy

12 Roy P. Drachman Hall, Rm. B307B

13 1295 N. Martin Ave

14 Tucson, AZ 85721

15 [schnell@pharmacy.arizona.edu](mailto:schnell@pharmacy.arizona.edu)

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47 **Abstract**

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49 Previous studies have shown that cGMP increases mitochondrial biogenesis (MB). Our  
50 laboratory has determined that formoterol and LY344864, agonists of the  $\beta_2$ -adrenergic and 5-  
51 HT<sub>1F</sub> receptor, respectively, signal MB in a sGC-dependent manner. However, the pathway  
52 between cGMP and MB produced by these pharmacological agents in renal proximal tubule cells  
53 (RPTC) and the kidney, has not been determined. We show that treatment of RPTC with  
54 formoterol, LY344864, or riociguat, an sGC stimulator, induces MB through protein kinase G  
55 (PKG), a target of cGMP, and p38, an associated downstream target of PKG and a regulator of  
56 PGC-1 $\alpha$  expression in RPTC. We also examined if p38 plays a role in PGC-1 $\alpha$  phosphorylation  
57 *in vivo*. L-Skepinone, a potent and specific inhibitor of p38 $\alpha$  and p38 $\beta$  administration to naïve  
58 mice inhibited phosphorylated PGC-1 $\alpha$  localization in the nuclear fraction of the renal cortex.  
59 Taken together, we have demonstrated a pathway, sGC/cGMP/PKG/p38/PGC-1 $\alpha$ , for  
60 pharmacological induction of MB and the importance of p38 in this pathway.

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80 **Introduction**

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82 In the presence of nitric oxide (NO), soluble guanylyl cyclase (sGC) produces cGMP from GTP.

83 The resulting cGMP can bind to cGMP gated ion channels, phosphodiesterases, and protein  
84 kinase G (PKG)<sup>9</sup>. As such, cGMP plays a role in a variety of processes in the cell including  
85 mitochondrial biogenesis (MB)<sup>21,22</sup>.

86  
87 PKG is a serine/threonine kinase that exists in two forms, PKG1 and PKG2<sup>14</sup>. In renal tubular  
88 cells, PKG1 activity and expression decreased when exposed to cisplatin<sup>19</sup>. Increasing PKG1  
89 activity protected mitochondrial function and prevented cell apoptosis<sup>19</sup>. In brown adipose tissue,  
90 natriuretic peptides activate GC resulting in activated PKG<sup>1,5</sup> and the induction of MB<sup>12</sup>.  
91 Adipocytes exposed lipoamide also undergo MB through PKG<sup>28</sup>.

92  
93 Activated PKG leads to the phosphorylation of p38 in human adipocytes when stimulated with  
94 naturetic peptides<sup>1,13</sup>. Browning et al., showed the importance of activated PKG in NO-induced  
95 p38 phosphorylation in 293T fibroblasts<sup>3</sup>. It was also shown in human platelets stimulated by  
96 thrombin, that p38 activation is necessary for integrin activation and activated PKG plays an  
97 important role in this mechanism<sup>18</sup>. These studies show that activated PKG plays a role in  
98 activating p38 in certain cell types.

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100 PGC-1 $\alpha$  is thought to be the master regulator of MB<sup>15,25</sup>. Phosphorylated p38 can directly  
101 phosphorylate PGC-1 $\alpha$  at three sites: Threonine 298, Threonine 262, and Serine 265<sup>8</sup>.  
102 Phosphorylation at these sites can increase the stability of PGC-1 $\alpha$ , promote its translocation into

103 the nucleus and induce transcription of PGC-1 $\alpha$  target genes<sup>12</sup>. Puigserver et al, showed that  
104 cultured muscle cells treated with LPS resulted in p38-mediated PGC-1 $\alpha$  phosphorylation<sup>24</sup>.

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106 Our laboratory showed that maximal mitochondrial respiration (i.e. uncoupled respiration), a  
107 marker of MB, increases in renal proximal tubule cells (RPTC) treated with the membrane  
108 soluble cGMP analogue 8-Br-cGMP but not 8-Br-cAMP, suggesting that cGMP is responsible  
109 for inducing MB rather than cAMP in RPTC<sup>31</sup>. Moreover, we have shown that cGMP is a key  
110 player in inducing MB by two G-protein-coupled receptor (GPCR) agonists. Formoterol, a  $\beta_2$   
111 adrenergic receptor ( $\beta_2$ AR) agonist, and LY344864, a 5-HT<sub>1F</sub> receptor agonist, induced MB  
112 through the G $_{\beta/\gamma}$  subunit, Akt, sGC, and cGMP<sup>4,10</sup>. Our studies are consistent with reports for  
113 other GPCRs, such as cannabinoid type 1 receptor, inducing MB through increased NO  
114 production<sup>30</sup>.

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116 However, the pathway from cGMP to MB following pharmacological stimulation in highly  
117 oxidative renal epithelial cells is not clear. We propose that formoterol, LY344864, and  
118 riociguat, an sGC stimulator that increases cGMP, activates protein kinase G (PKG), which leads  
119 to the phosphorylation of p38. In turn, p38 phosphorylates PGC-1 $\alpha$  to facilitate translocation to  
120 the nucleus and produce MB.

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126 **Material and Methods**

127 *Reagents:*

128 L-Skepinone was purchased from Selleckchem (Houston, TX). LY344864 and KT5823 was  
129 purchased from Tocris (Minneapolis, MN). Riociguat was purchased from Biovision (Milpitas,  
130 CA). Formoterol fumarate was purchased from Sigma (St. Louis, MO).

131

132 *In vitro Studies:*

133 Renal Proximal tubule cells (RPTC) were isolated from female NZW rabbit (2kg) kidneys using  
134 the iron oxide perfusion method<sup>23</sup>. Cells were plated and grown on 35 mm tissue culture dishes  
135 in conditions that are similar to physiological conditions *in vivo*. Confluent RPTC were treated  
136 with riociguat (10  $\mu$ M), LY344864 (10 nM), formoterol (30 nM) or vehicle, consisting of DMSO  
137 (<0.5%). For the inhibitor studies, RPTC were pretreated with 100 nM KT5823 (KT) or 100 nM  
138 skepinone (SK) for 30 min. Riociguat, formoterol, LY344864, or DMSO was added to RPTC,  
139 incubated for 2 hr and harvested for further analysis. The concentrations of formoterol and  
140 LY344864 have previously be shown to induce MB in RPTC<sup>4,10</sup>.

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142 *In vivo Studies:*

143 Skepinone was dissolved in DMSO and diluted in sterile saline. Final DMSO concentration was  
144 2%. Eight to nine-week old male C57BL/6 mice (20–25 g from Charles River Laboratories) were  
145 injected intraperitoneally with skepinone at 1, 3, or 10 mg/kg or vehicle. After 6 hr the mice were  
146 euthanized and kidneys were removed. Part of the kidney was flash frozen or processed for  
147 subcellular fractionation. Animal studies and animal use was approved by the Institutional  
148 Animal Care and Use Committee at the University of Arizona.

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150 *Subcellular Fractionation:*

151 RPTC were harvested in sucrose isolation buffer containing 250 mM sucrose, 1 mM EGTA, 10  
152 mM HEPES, and 1 mg/ml fatty acid free BSA at a pH of 7.4. Cells were homogenized using a  
153 dounce homogenizer and centrifuged at 700 g for 5 min. The cytosolic supernatant was stored in  
154 phosphatase inhibitors (1:100), 1mM sodium orthovanadate, and 1 mM sodium fluoride, and  
155 Triton X-100 and SDS at 4%. The pellet was washed twice in isolation buffer and centrifuged at  
156 1,000 g for 5 min. The pellet/nuclear fraction, was resuspended in RIPA buffer containing 50  
157 mM Tris-HCl, 150 mM NaCl, 0.1% SDS, 0.5% sodium deoxycholate, 1% Triton X-100, pH 7.4  
158 with phosphatase inhibitors (1:100), 1mM sodium orthovanadate, and 1 mM sodium fluoride.  
159 Purity of the cytosolic and nuclear fractions was determined using immunoblot analysis. Histone  
160 H3 and/or lamin B1 were used as nuclear markers, and  $\alpha$ -tubulin was used as a cytosolic marker.  
161 Antibodies for Histone H3 were purchased from Cell Signaling Technology (Danvers, MA).  
162 Antibodies for lamin B1 and  $\alpha$ -tubulin were purchased from Abcam (Cambridge, MA).

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164 *Immunoblot Analysis*

165 RPTC or snapped frozen mouse kidney cortex was added to RIPA buffer. Cells were sonicated  
166 for approximately 10 seconds and centrifuged at 7,500 g for 5 min at 4°C. Supernatants were  
167 removed and protein was measured using a BCA assay. Equal protein was loaded onto 4-15%  
168 SDS page gels and separated by gel electrophoresis. Protein was transferred onto nitrocellulose  
169 membranes and blocked in 5% milk or 5% bovine serum albumin dissolved in TBST.  
170 Membranes were incubated with primary antibodies overnight. Membranes were washed in  
171 TBST 3 times for 5 min, incubated with a horseradish peroxidase-conjugated secondary

172 antibody, and visualized using enhanced chemiluminescence (Thermo Scientific) and GE  
173 ImageQuant LAS4000 (GE Life Sciences). Optical density was determined using the ImageJ  
174 software from NIH. Primary antibodies p-vasodilator-stimulated phosphoprotein (VASP) Ser239  
175 (1:1,000), phospho-p38 MAPK (Thr180/Tyr182) (1:1000), and p38 MAPK (1:1000) were  
176 purchased from Cell Signaling Technology (Danvers, MA).

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### 178 *Immunoprecipitation*

179 Dynabeads<sup>TM</sup> Protein G Immunoprecipitation Kit (ThermoFisher) was used to immunoprecipitate  
180 PGC-1 $\alpha$  in the cytosolic and nuclear fractions. Protein (200  $\mu$ g) from cytosolic and nuclear  
181 fractions were precleared with Pierce Protein A/G Plus Agarose beads (ThermoFisher) for 2 hr  
182 and then centrifuged at 14,000g for 10 min at 4°C. Dynabeads<sup>TM</sup> magnetic beads were incubated  
183 with 10  $\mu$ g of PGC-1 $\alpha$  antibody for 4 hr at room temperature. Supernatants from the precleared  
184 cytosolic and nuclear fractions were added to the magnetic beads and PGC-1 $\alpha$  antibody and  
185 incubated with rotation overnight at 4° C. PGC-1 $\alpha$  was immunoprecipitated and eluted  
186 (denaturing elution) based on manufacturer's instructions. The resulting supernatant was directly  
187 loaded on to a 4-15% SDS page gel and separated by gel electrophoresis. After protein transfer,  
188 one nitrocellulose membrane was blocked in 5% milk for measuring total PGC-1 $\alpha$  and the other  
189 was blocked in 5% bovine serum albumin for measuring phosphorylated serine/threonine  
190 residues for 1 hr. Membranes were incubated with phosphoserine/threonine (1:1,000) antibody  
191 from Abcam (Cambridge, MA) or PGC-1 $\alpha$  antibody (1:1,000) from EMD Millipore (Billerica,  
192 MA) overnight at 4°C. After the membranes were washed in TBST 3 times, they were incubated  
193 in horseradish peroxidase-conjugated secondary antibodies from Abcam (Cambridge, MA) for 2  
194 hrs at room temperature. Membranes were visualized as described above.

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*Statistics*

Data are represented as mean  $\pm$  SEM. RPTC isolated from a single rabbit represent N=1. For a single comparison, Student's *t*-test was performed. Multiple comparisons were subjected to one-way ANOVA and Tukey's post hoc test with  $p < 0.05$  being statistically significant between means. Statistical differences between means are denoted by an \*, #, or \$ accordingly.

218 **Results**

219 **Riociguat activates PKG and is blocked by KT5823**

220 To elucidate the signaling pathway from sGC to nuclear PGC-1 $\alpha$  phosphorylation, RPTC were  
221 treated with riociguat (10 $\mu$ M), a sGC stimulator that targets the reduced form of sGC<sup>2,16,20,29</sup>. The  
222 reduced form of sGC predominates in naïve cell types and increases cGMP directly. cGMP  
223 production can activate PKG, and PKG activation was determined by measuring the  
224 phosphorylation of vasodilator-stimulated phosphoprotein (VASP) at site serine 239, a specific  
225 target of PKG<sup>6</sup>.

226

227 RPTC were treated with riociguat for 2 hr, resulting in a 1.3-fold increase in VASP  
228 phosphorylation compared to controls (Figure 1). Pretreatment of KT5823, a PKG inhibitor, for  
229 30 min and subsequent treatment with vehicle or riociguat for 2 hr resulted in a decrease in  
230 VASP phosphorylation to control levels, demonstrating that KT inhibits PKG. Pretreatment with  
231 KT5823 alone lowered phosphorylation of VASP in RPTC to below control levels.

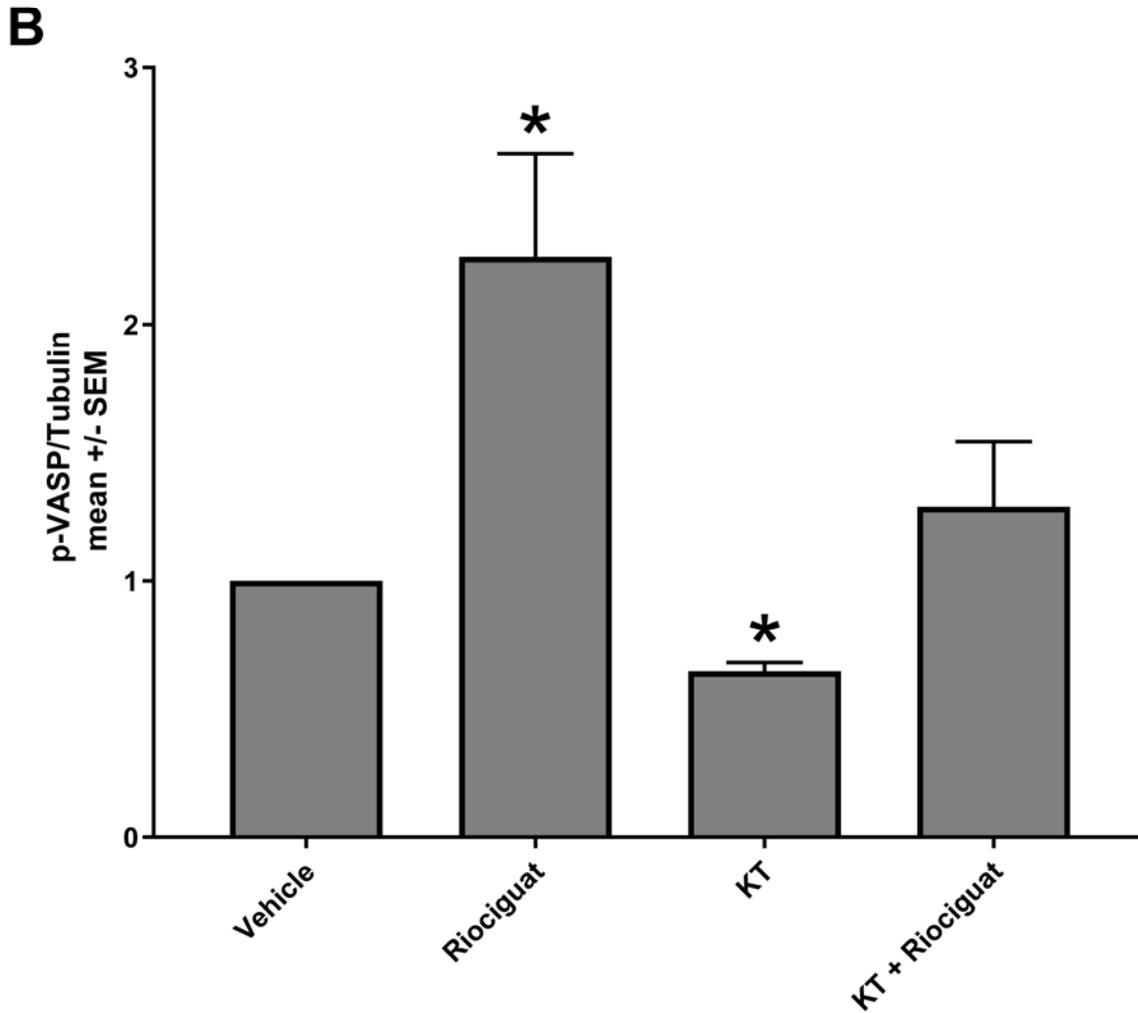
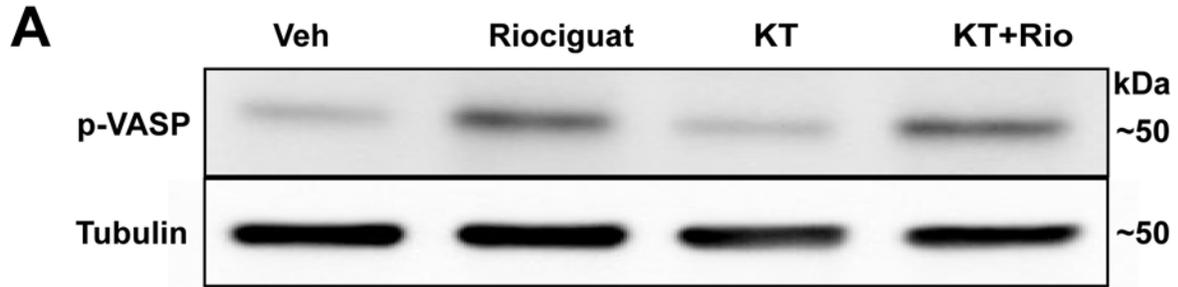
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240 **Figure 1: Riociguat phosphorylates VASP (p-VASP) and is blocked by KT5823 (KT)**

241 A) Representative immunoblot for p-VASP and tubulin after treatment. B) Densitometry analysis

242 for p-VASP protein. Data are represented as mean S.E., N=6-7. \* represents significance

243 compared to Vehicle ( $p < 0.05$ ).

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246 **Riociguat activates p38 through PKG**

247 We hypothesized that p38 is the mediator between PKG activation and PGC-1 $\alpha$  phosphorylation  
248 in RPTC. Riociguat treated RPTC increased p-p38 2.1-fold at 2 h. Pretreatment with KT5823 for  
249 30 min with subsequent treatment with vehicle or riociguat for 2 hr resulted in a decrease in p-  
250 p38 to control levels (Figure 2A-2B).

251 To validate skepinone inhibition of p38 phosphorylation at 2 h in the presence and  
252 absence of riociguat, RPTC were treated with skepinone for 30 min and then treated with vehicle  
253 or riociguat for 2 h. Skepinone inhibited p-p38 when RPTC were treated with riociguat (90%) or  
254 vehicle (80%) at 2 h (Figure 2C-D).

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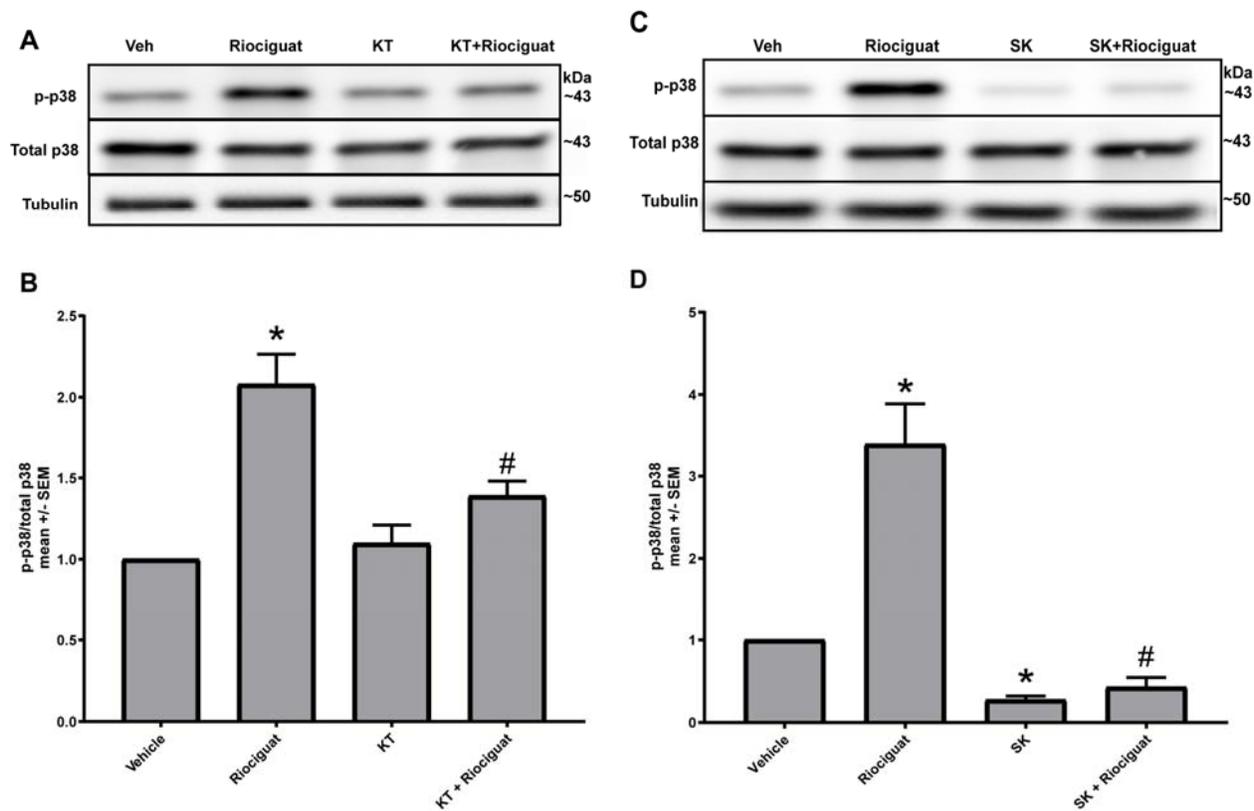
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266 **Figure 2: KT5823 and skepinone inhibit phosphorylation of p38**  
 267 A and C) Representative blot for p-p38, total p38, and tubulin after treatment. B and D)  
 268 Densitometry analysis for p-p38 protein. Data are represented as mean S.E., N=6-7. \* represents  
 269 significance compared to Vehicle ( $p < 0.05$ ). # represents significance compared to riociguat  
 270 ( $p < 0.05$ ).  
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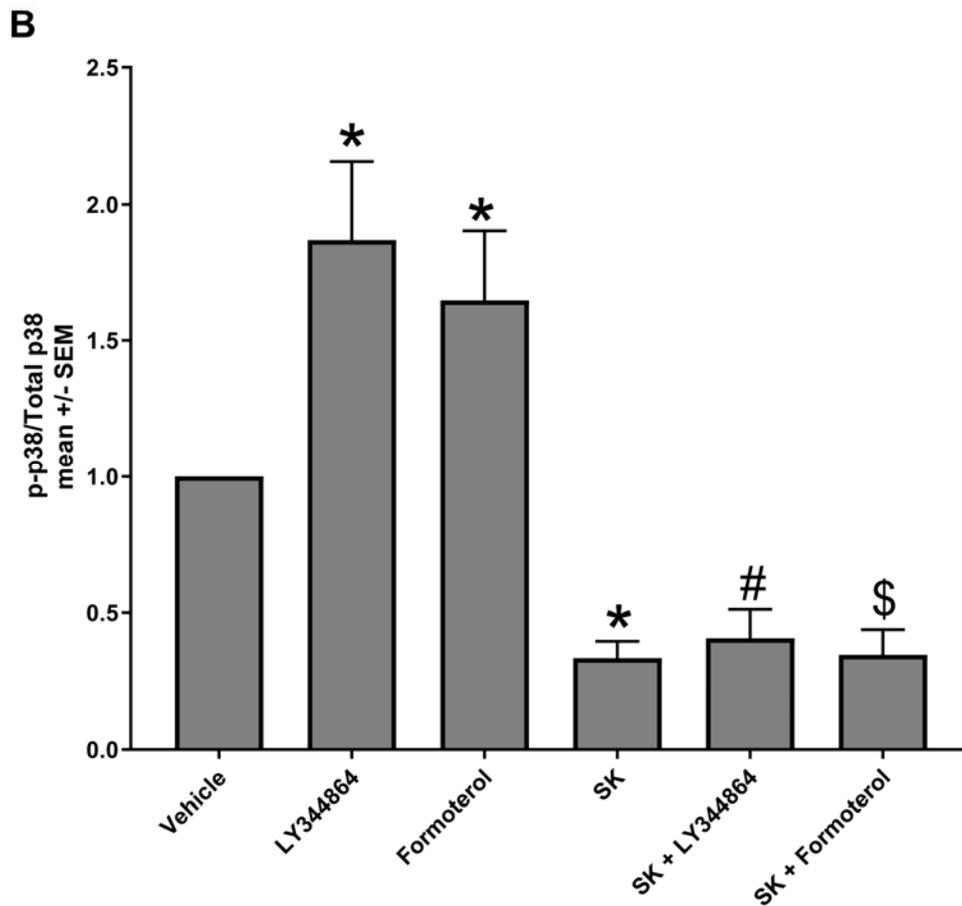
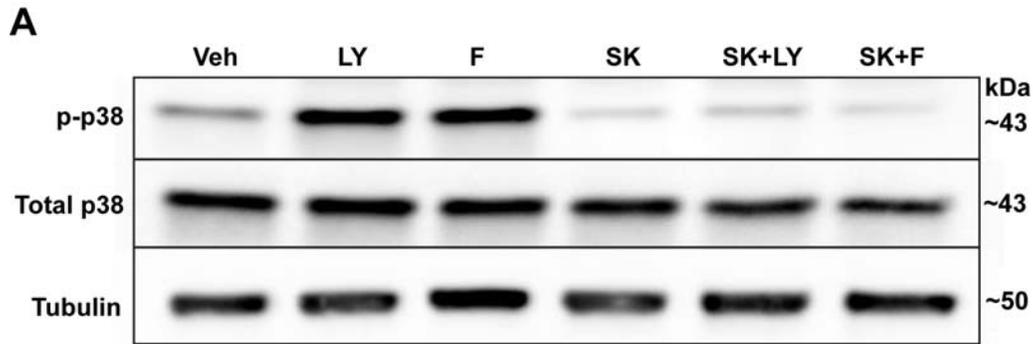
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**LY344864 and formoterol activate p38**

Based on the above results, we examined if p38 plays a role in the signaling pathways of formoterol and LY344864. We treated RPTC with vehicle, formoterol, or LY344864 for 2 hr. p38 phosphorylation increased 1.6-fold and 1.9-fold when exposed to formoterol and LY344864, respectively (Figure 3). RPTC were pretreated with skepinone for 30 min and exposed to vehicle, formoterol, and LY344864 for 2 h. Skepinone alone inhibited p-p38 by 70% compared to vehicle control. Skepinone pretreatment inhibited p-p38 by 80% after formoterol and LY344864 treatment.



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297 **Figure 3: Skepinone inhibits LY344864 and formoterol mediated p38 phosphorylation**

298 A) Representative blot for p-p38, total p38, and tubulin after treatment. B) Densitometry analysis  
 299 for p-p38 protein. Data are represented as mean S.E., N=8. \* represents significance compared to  
 300 Vehicle ( $p < 0.05$ ). # represents significance compared to LY344864. \$ represents significance  
 301 compared to formoterol.

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303 **Pretreatment with KT5823 or skepinone decreases nuclear phosphorylated PGC-1 $\alpha$  at**  
304 **serine and threonine residues in the presence of riociguat**

305 Phosphorylation of PGC-1 $\alpha$  at serine and threonine sites by p38 can increase the stability of  
306 PGC-1 $\alpha$ , resulting in its translocation to the nucleus and transcription of PGC-1 $\alpha$ . RPTC were  
307 pretreated with KT5823 for 30 min and then exposed to vehicle or riociguat for 2 h. RPTC were  
308 subjected to subcellular fractionation and tested for purity of nuclear and cytosolic fractions by  
309 measuring  $\alpha$ -tubulin, a cytosolic marker, and Lamin B, a nuclear marker<sup>7,11</sup>. PGC-1 $\alpha$  was  
310 immunoprecipitated from the nuclear and cytosolic fractions, and immunoblotted for  
311 phosphorylated serine and threonine residues (Figure 4A,C). Phosphorylated PGC-1 $\alpha$  in the  
312 nuclear fraction was increased by 1.64-fold following riociguat treatment (Figure 4B), and  
313 decreased to control levels in the presence of KT5823. There were no changes in PGC-1 $\alpha$   
314 phosphorylation in the cytosolic fraction compared to vehicle control (Figure 4C-D).

315 RPTC were pretreated with skepinone for 30 min and then exposed to vehicle or riociguat  
316 for 2 h. Phosphorylated PGC-1 $\alpha$  was measured in nuclear and cytosolic fractions. Skepinone  
317 alone had no effect on phosphorylated PGC-1 $\alpha$  in the nucleus (Figure 5A,B). Skepinone  
318 decreased phosphorylated PGC-1 $\alpha$  in the nucleus to control levels when exposed to riociguat.  
319 Phosphorylated PGC-1 $\alpha$  did not change in the cytosolic fraction compared to vehicle control  
320 (Figure 5C,D).

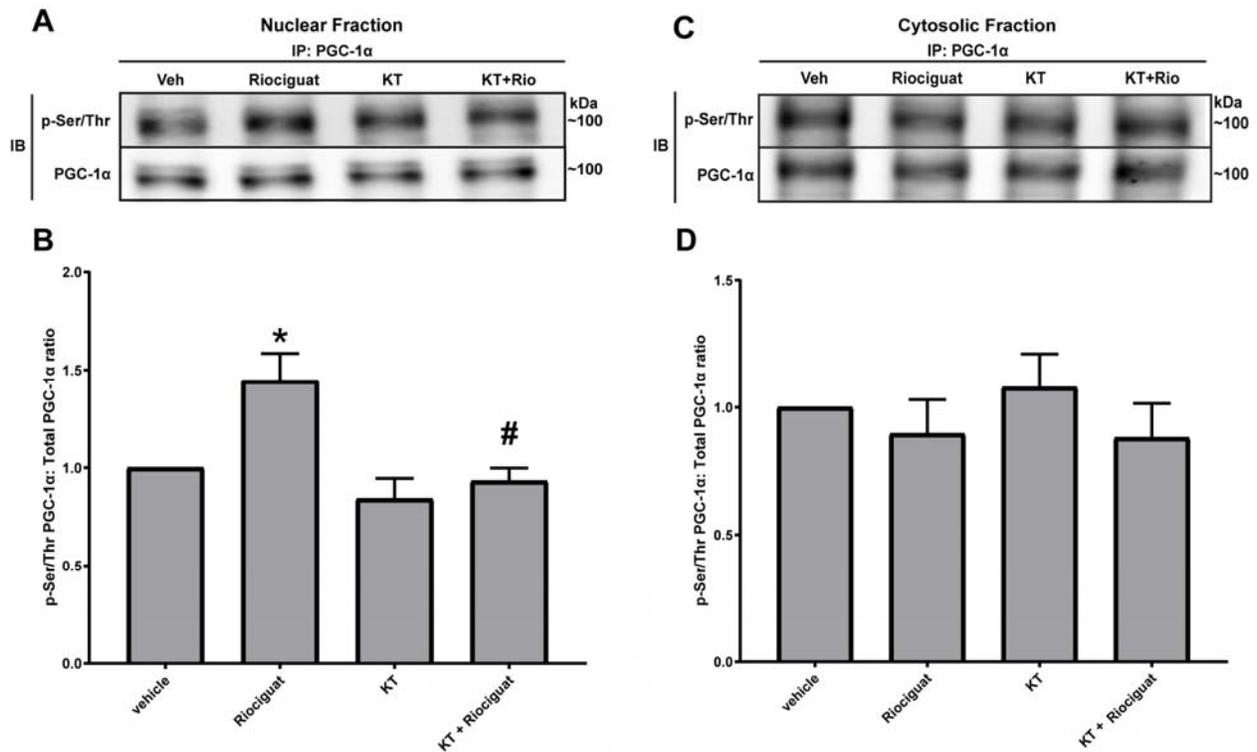
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327 **Figure 4: Pretreatment with KT5823 decreases nuclear phosphorylated PGC-1α in the**  
 328 **presence of riociguat.** A and C) Phosphorylated serine and threonine residues were measured  
 329 following immunoprecipitation of PGC-1α by immunoblot analysis in nuclear and cytosol  
 330 fractions after 30 min pretreatment with Vehicle or KT5823 (KT) followed by exposure to  
 331 DMSO and riociguat for 2 hr. Total PGC-1α expression was measured after  
 332 immunoprecipitation. B and D) Densitometry analysis for phosphorylated serine and threonine  
 333 residue in the nuclear and cytosolic fractions. Data are represented as mean S.E., N=6. \*  
 334 represents significance compared to Vehicle control (p<0.05). # represents significance  
 335 compared to riociguat.

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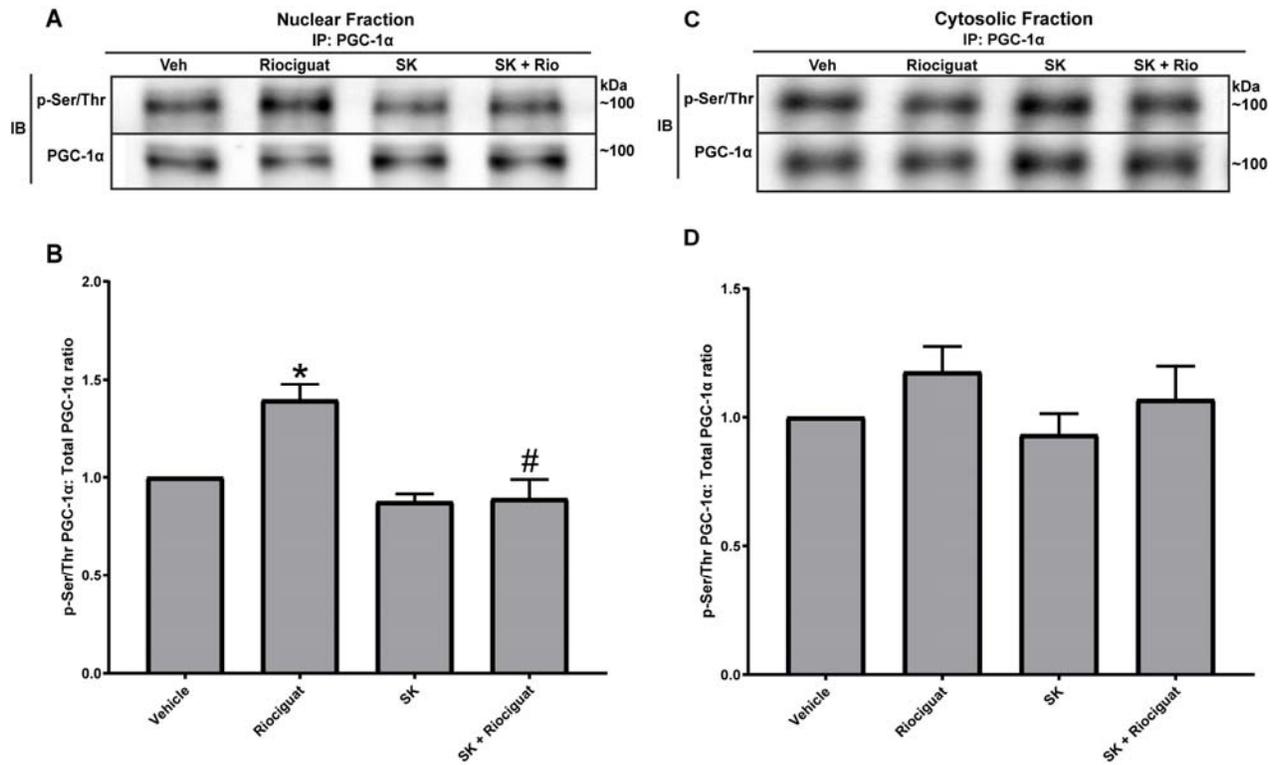
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348 **Figure 5: Pretreatment with skepinone decreases nuclear phosphorylated PGC-1α at serine**  
 349 **and threonine sites in the nucleus**

350 A and C) Phosphorylated serine and threonine residues were measured following  
 351 immunoprecipitation of PGC-1α by immunoblot analysis after 30 min treatment with DMSO or  
 352 100nM SK followed by exposure to DMSO and riociguat for 2 hr, in cytosol and nuclear  
 353 fractions. Total PGC-1α expression was measured after immunoprecipitation. B and D)  
 354 Densitometry analysis for phosphorylated serine and threonine residues in the cytosol and  
 355 nuclear fractions. Data are represented as mean S.E., N=6-7. \* represents significance compared  
 356 to Vehicle control (p<0.05). # represents significance compared to riociguat.

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370 **Skepinone decreases nuclear phosphorylated PGC-1 $\alpha$  in the renal cortex**

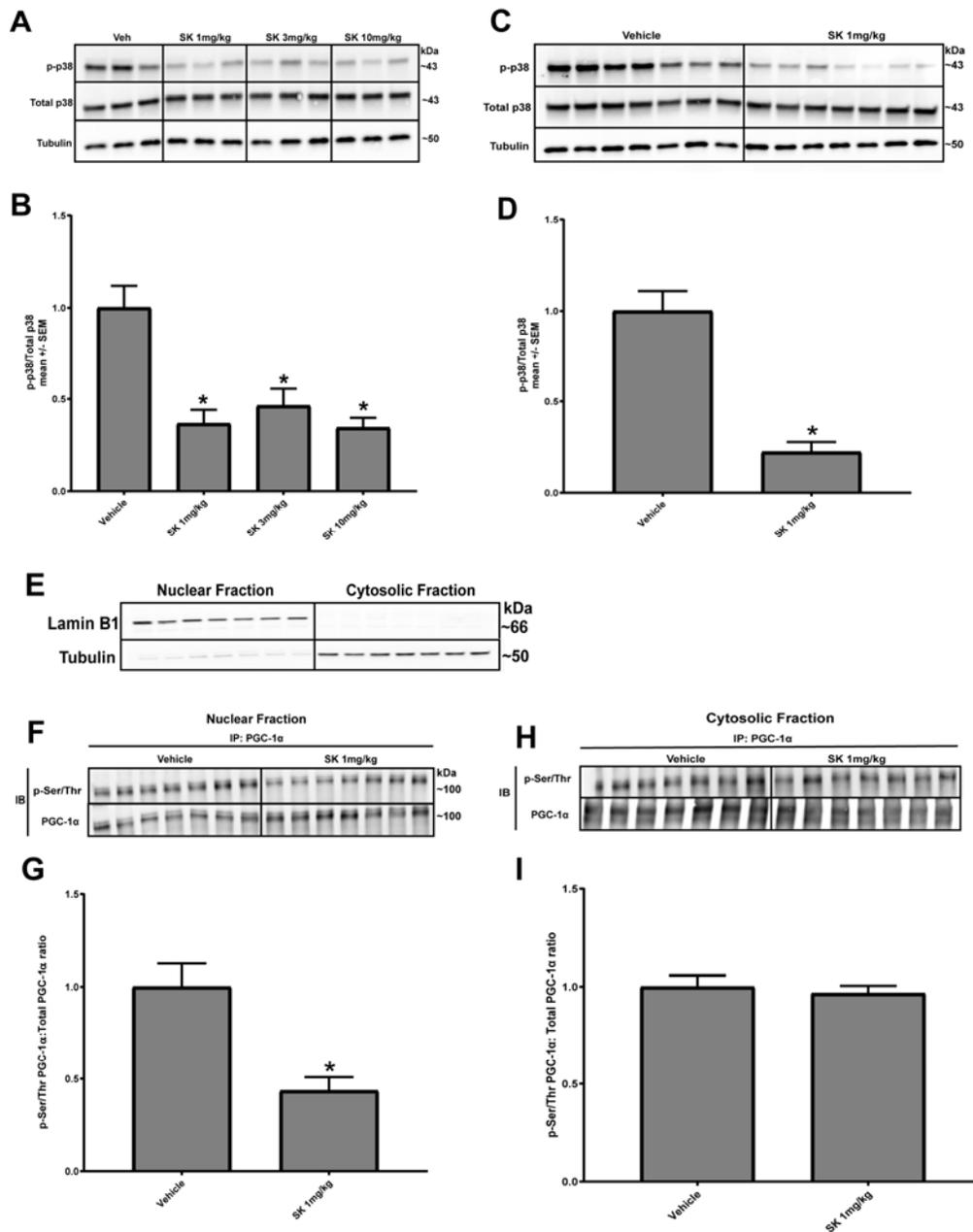
371 Based on the above cellular data, we determined if p38 inhibition alters phosphorylated PGC-1 $\alpha$   
372 in the kidney. We performed a dose response experiment using skepinone and measured p38  
373 phosphorylation in the renal cortex. Naïve mice were injected with 1 mg/kg, 3 mg/kg, and  
374 10mg/kg of skepinone and then euthanized 6 h after injection. At all three doses, skepinone  
375 equally blocked p38 phosphorylation by more than 50% (Figure 6A,B). Using the lowest dose of  
376 skepinone (1 mg/kg), the number of mice was increased (N=6-7) (Figure 6C) and skepinone  
377 decreased p38 phosphorylation by 75% (Figure 6D).

378 The next experiment measured phosphorylated PGC-1 $\alpha$  in cytosolic and nuclear fractions  
379 of the renal cortex (Figure 6F and H). Mice were treated with skepinone (1 mg/kg) for 6 h and  
380 renal cortex was subcellular fractionated. The purity of cytosolic and nuclear fractions was tested  
381 (Figure 6E). In the cytosolic fraction, skepinone had no affect on phosphorylated PGC-1 $\alpha$ ;  
382 however, in the nuclear fraction, skepinone decreased phosphorylated PGC-1 $\alpha$  by 50% (Figure  
383 6G, I).

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389 **Figure 6: Skepinone (SK) decreases nuclear phosphorylated PGC-1α in the renal cortex**

390 Representative blot for p-p38, total p38, and tubulin in naïve mice, 6 hr after treatment with 1  
 391 mg/kg, 3 mg/kg, 10 mg/kg of SK. C) Representative blot for p-p38, total p38, and tubulin with 1  
 392 mg/kg SK. B-D) Densitometry analysis for p38 protein. Data are represented as mean S.E., N=3-  
 393 7. E) Representative blot for tubulin and lamin B1 in nuclear and cytosol fractions. F and H)  
 394 Phosphorylated serine and threonine residues were measured following immunoprecipitation of  
 395 PGC-1α in cytosol and nuclear fractions by immunoblot analysis. G and I) Densitometry analysis  
 396 for phosphorylated serine and threonine residue in the cytosol and nuclear fractions. Data are  
 397 represented as mean S.E., N=7 \* represents significance compared to Vehicle (p<0.05). Black  
 398 lines in the center of blots are used for dividing experimental groups only and do not alter  
 399 information contained therein.

400 **Discussion**

401 Previous studies from our laboratory have shown that cGMP, not cAMP, induces MB in  
402 RPTC<sup>31</sup>. PDE3 inhibitors such as cilostamide and trequinsin, compounds that prevent the  
403 degradation of cGMP, also induce MB in RPTC<sup>31</sup>. *In vivo*, sildenafil, a PDE5 inhibitor, also  
404 induces MB in the renal cortex of naïve mice<sup>31</sup>. However, the mechanisms by which cGMP can  
405 induce MB are still under investigation for different cell types.

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407 We showed the induction of MB in RPTC treated with formoterol, a  $\beta_2$ AR agonist, and  
408 LY344864, a 5-HT<sub>1F</sub> agonist. Moreover, we reported that sGC and cGMP are important  
409 components for signaling MB in RPTC<sup>4,10</sup>. In this study, we proposed a signaling pathway  
410 between sGC and MB, sGC/cGMP/PKG/p38/PGC-1 $\alpha$  in RTPC. In addition, we  
411 pharmacologically targeted sGC by administering riociguat, an sGC stimulator that targets the  
412 reduced/heme-dependent form<sup>2,16,20,29</sup>.

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414 To elucidate the role of p38 in the sGC-dependent induction of MB, skepinone, a potent  
415 and selective ATP-competitive inhibitor for p38 $\alpha$  and p38 $\beta$  inhibitor, was used<sup>17</sup>. Previous  
416 studies have shown that p38 can directly phosphorylate PGC-1 $\alpha$  at Threonine 298, Threonine  
417 262, and Serine 265 in the cytosol, causing its translocation to the nucleus to sustain PGC-1 $\alpha$   
418 transcription and therefore MB<sup>24</sup>. We show that inhibition of PKG activation and p38  
419 phosphorylation, decreases nuclear phosphorylated PGC-1 $\alpha$  when RPTC are exposed to  
420 pharmacological compounds riociguat, formoterol and LY344864, demonstrating that PKG  
421 activation and p38 are needed in the pathway from sGC to MB. Further support for this pathway  
422 is that administration of skepinone *in vivo*, decreased nuclear phosphorylated PGC-1 $\alpha$  in the  
423 renal cortex.

424 Previous literature shows that PKG activation can lead to p38 phosphorylation and that  
425 p38 is a downstream associated target of PKG<sup>1,3,18,26</sup>. Based on this literature, we did not explore  
426 the effect of skepinone on PKG activation in the presence of riociguat. Although the literature  
427 provides evidence for other downstream effectors such as ERK1/2, GSK3 $\beta$ , and Akt, we chose  
428 to focus on p38 for its role in NO signaling, cGMP/PKG dependent signaling, and regulation of  
429 PGC-1 $\alpha$ <sup>3,8,18,26,27</sup>.

430 In summary, we have elucidated a pathway for sGC-dependent induction of MB by  
431 investigating the importance of PKG and subsequent activation of p38 leading to an increase in  
432 nuclear phosphorylated PGC-1 $\alpha$  by pharmacological compounds. This study supports our  
433 previous work with potent inducers of MB and show the importance of sGC/cGMP in the  
434 signaling pathways leading to the induction of MB in the kidney.

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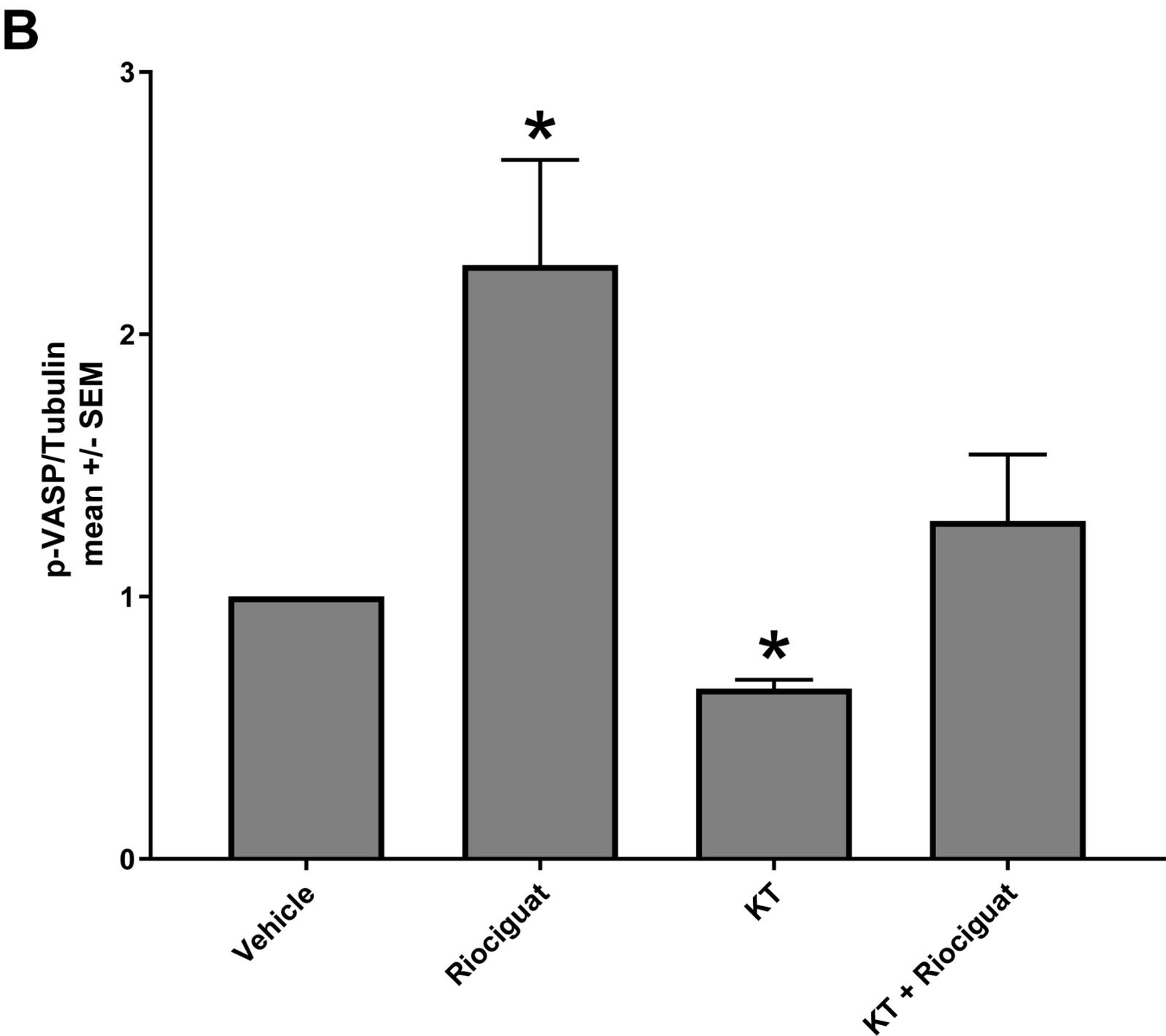
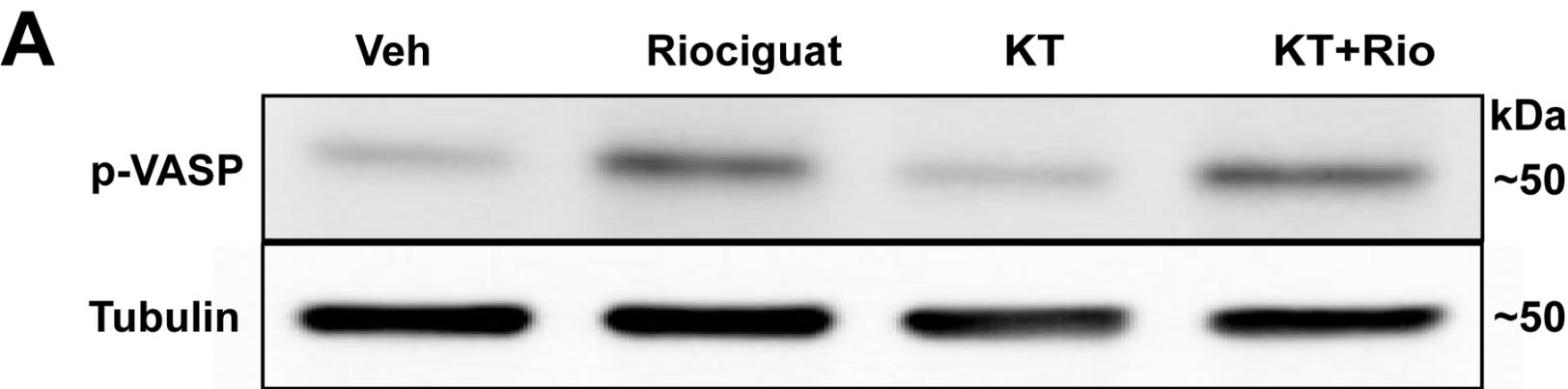
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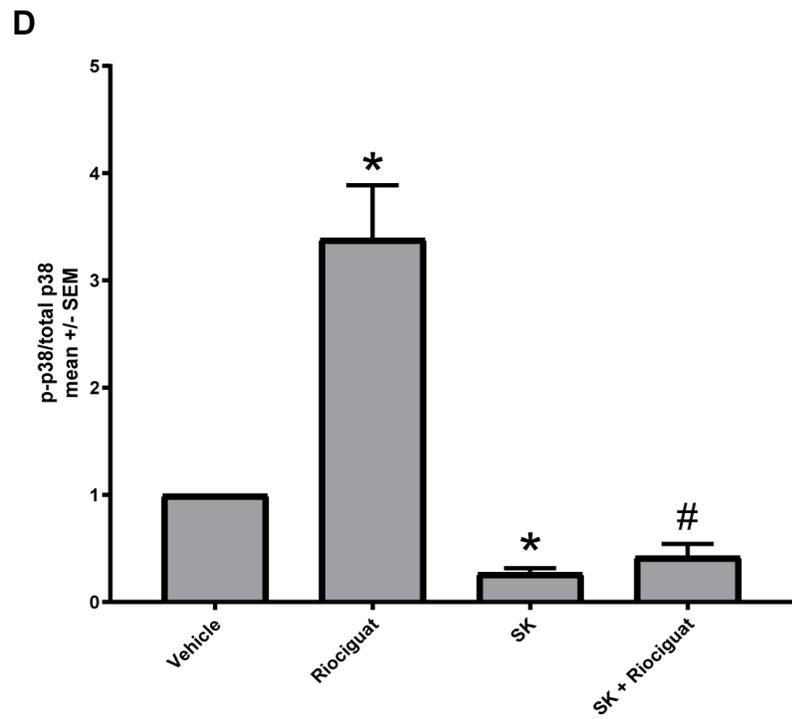
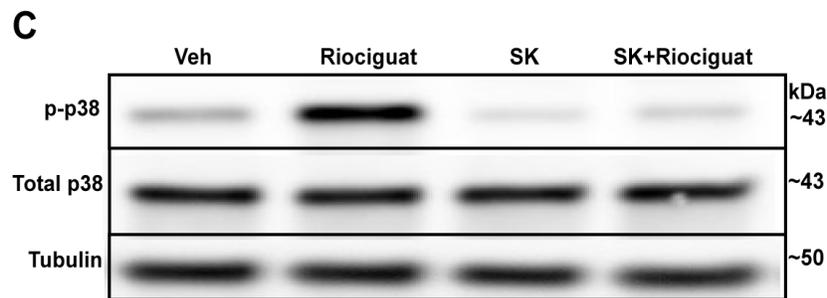
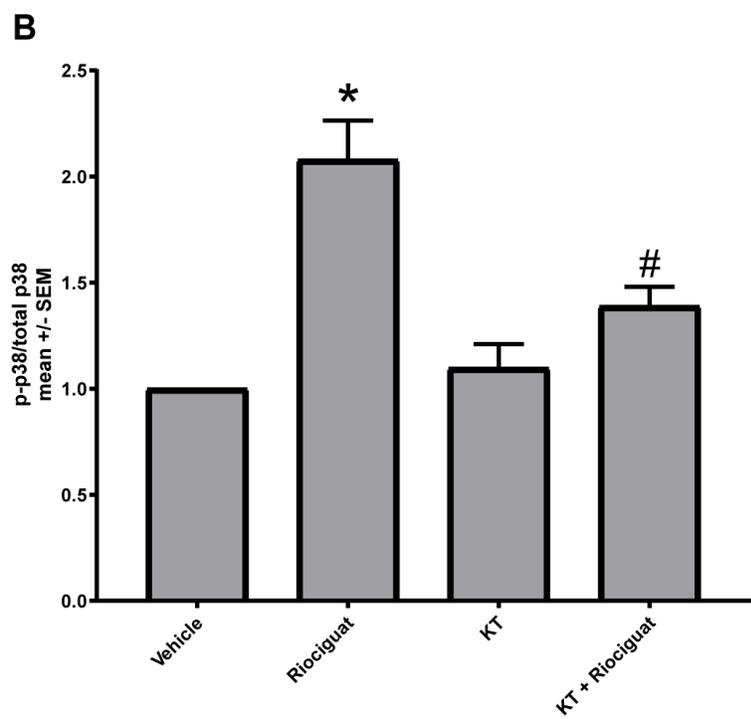
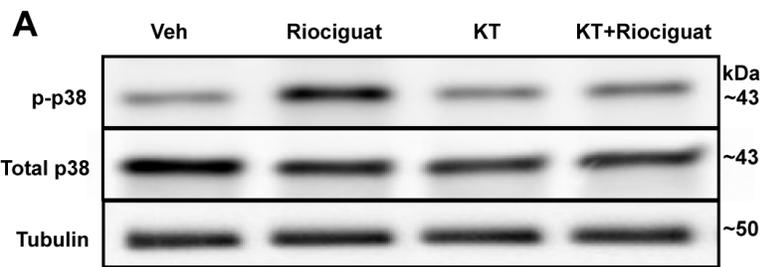
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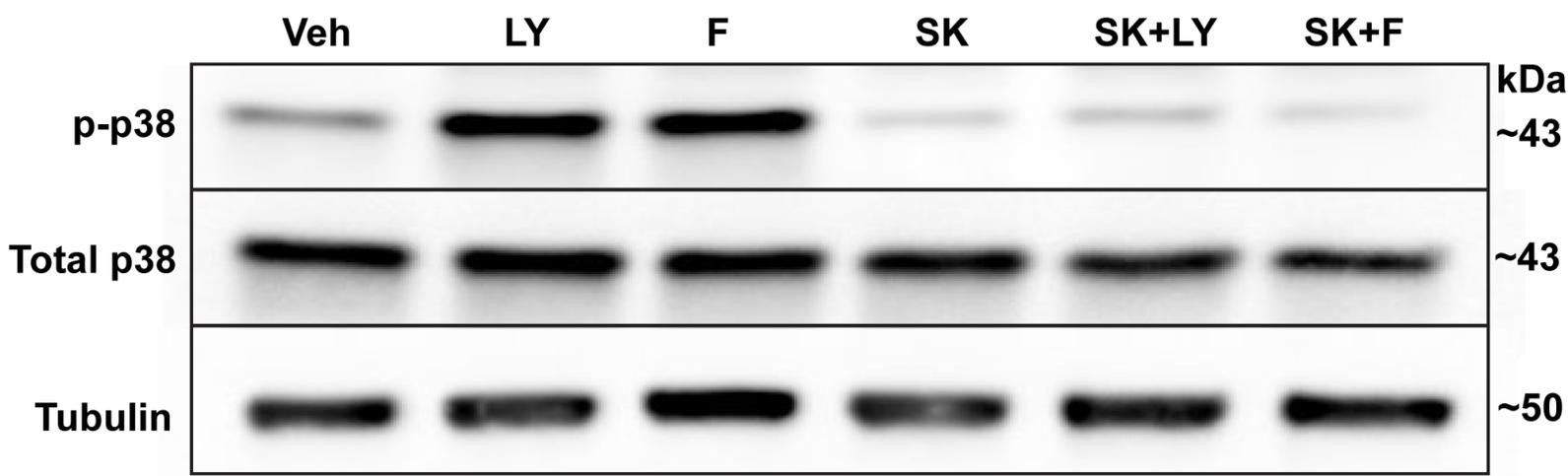
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