

1 **Metformin Confers Cardiac and Renal Protection in Sudden Cardiac Arrest via AMPK Activation**

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26

27 **Abstract**

28 Sudden cardiac arrest (SCA) affects over 600,000 individuals annually in the United States and is
29 associated with substantial mortality. After resuscitation, multi-system organ damage is common and
30 largely attributable to ischemia-reperfusion injury. The anti-diabetic drug metformin improves cardiac
31 outcomes in models of myocardial ischemia and ischemia-reperfusion. In this study, we evaluated the
32 role of metformin pretreatment in a mouse model of SCA. We found that two weeks of metformin
33 pretreatment protects cardiac ejection fraction and reduces acute kidney injury post-SCA in non-diabetic
34 mice. Further, metformin pretreatment prior to SCA activates AMPK signaling and is associated with
35 altered mitochondrial dynamics and markers of autophagy following arrest. Direct AMPK activation and
36 inhibition studies demonstrate that activation is necessary and sufficient for metformin-mediated
37 protection of cardiac and renal tissues in this model. We were unable to demonstrate cardiac protection
38 with a single-dose metformin rescue therapy. Importantly, these findings translate into patients. We
39 retrospectively evaluated the extent of cardiac and kidney damage in diabetic patients resuscitated from
40 SCA. Metformin-treated patients have less evidence of heart and kidney damage after arrest than
41 diabetics who have not received metformin. Together, these data support AMPK activation as a
42 preventive mechanism in ischemia-reperfusion injury.

43

44 Introduction

45 Sudden cardiac arrest (SCA) refers to the abrupt cessation of cardiac function and affects over
46 600,000 patients annually in the United States (1,2). Patients with return of spontaneous circulation
47 after SCA experience systemic ischemia-reperfusion injury, typically resulting in multi system organ
48 damage. Common findings include cardiogenic shock, acute renal failure, liver damage, and neurologic
49 dysfunction (3–5). Previous observational studies in cardiac arrest patients have shown that low cardiac
50 ejection fraction (EF) (6) and reduced kidney function (7,8) are predictors of increased mortality. Despite
51 its prevalence, no pharmacologic therapy has been shown to improve overall survival in post-cardiac
52 arrest syndrome.

53 Metformin is an oral antihyperglycemic agent used as the first-line agent for type 2 diabetes
54 that has proven beneficial in a number of cardiovascular conditions(9,10). Metformin enhances insulin
55 sensitivity and normalizes glucose and lipid homeostasis (11–13). Beyond its role in controlling diabetes,
56 metformin has demonstrated clinical benefit across a wide variety of pathologies, including improved
57 mortality in the setting of coronary artery disease (10), congestive heart failure (14), acute kidney injury
58 (AKI)(15), chronic kidney disease (14), septic shock (16,17), and major surgical procedures (18).
59 Cardiovascular studies suggest that improved outcomes occur independently of the glucose-lowering
60 effects of metformin and may instead be attributable to metformin's other pleiotropic effects (19).
61 Several mechanisms beneficial to cardiovascular health have been implicated in metformin's numerous
62 effects, including reduced oxidative stress, anti-apoptotic activities, JNK inhibition, complex I inhibition,
63 and AMPK activation (20,21).

64 Importantly, AMPK activity and expression is induced in mice and humans by ischemic stress as
65 a compensatory response (22). AMPK activity limits endoplasmic reticulum (ER) stress (23) and AMPK
66 deficiency can be partially rescued through reduction of mitochondrial oxidative stress (24). While AMPK
67 activity is essential for survival after ischemic stress (23), it is unclear whether AMPK activity is the
68 mediator of metformin-mediated protection in ischemic heart disease. Furthermore, it is not clear
69 whether the adaptive upregulation of AMPK during cardiac stress is optimal, or whether further
70 activation could even more strongly impact recovery in cardiac injury models.

71 In this study, we sought to test metformin's potential benefit on heart and kidney protection
72 after SCA, both to clarify its relevance to the development of cardiac and peripheral tissue dysfunction,
73 and to determine the role of AMPK in post-arrest outcomes. Therefore, we evaluated outcomes
74 following SCA in mice with chemical activation of AMPK, via 5-aminoimidazole-4-carboxamide-1- β -D-
75 ribofuranoside (AICAR; an AMP-mimetic(25)) and with metformin treatment, and with inhibition of
76 AMPK via compound C (a reversible competitive inhibitor(26)) given concomitantly with metformin
77 treatment. We found that both metformin and AICAR pretreatment improved cardiac and renal
78 outcomes after resuscitation from SCA, and that metformin therapy is associated with altered markers
79 of autophagy and mitochondrial dynamics. We also showed that metformin's benefits are negated by
80 compound C, supporting an AMPK-dependent mechanism. Further, we performed a retrospective
81 analysis of clinical outcomes in diabetic cardiac arrest patients with and without metformin therapy
82 prior to arrest. We found that diabetic patients taking metformin prior to SCA had lower serum markers
83 of cardiac and renal damage 24 hours after arrest than non-metformin diabetic patients. Taken
84 together, we have identified AMPK activity as a protective mechanism invoked SCA-induced damage and

85 demonstrated benefit of metformin pretreatment on cardiac and renal outcomes in SCA, which provides
86 substantial support for metformin's use as a prophylactic in patients at risk for SCA.

87

88 Results

89 SCA mice have increased AMPK signaling by pathway analysis

90 To gain insight into gene expression pathways affected in the heart *in vivo* after SCA, we
91 performed microarray analysis of left ventricles (LVs) 24 hours post-resuscitation. In these initial
92 discovery experiments, male and female mice were evenly divided into untreated sham and untreated
93 arrest groups, where the arrest group underwent ultrasound-guided direct LV injection of potassium
94 chloride (KCl) to cause SCA (Figure 1A). In brief, these mice sustained eight minutes of asystole followed
95 by up to three minutes of cardiopulmonary resuscitation (CPR) until return of spontaneous circulation
96 (ROSC) occurred. One day after surgery, LV tissue was collected for RNA expression analysis (Figure 1B).
97 From Ingenuity Pathway Analysis, *AMPK Signaling* pathway was the most prominent by ranked p-value,
98 while *Autophagy* was the sixth most significantly changed pathway between sham and untreated SCA
99 mice (Figure 1C, Supplemental Figure 1).

100 Metformin pretreatment protects cardiac EF and kidney function after SCA

101 AMPK signaling has been shown to be upregulated in myocardial ischemia/reperfusion injury as
102 a compensatory response (27). Similarly, loss of key AMPK subunits increases infarct size in experimental
103 systems (23,24). In *ex vivo* rat hearts, infarct size can be reduced by acute, transitory AMPK activation
104 (28,29). Furthermore, AMPK activation has been shown to delay the progression of heart failure in a
105 chronic pressure overload model (30). Although the etiology of cardiac dysfunction in myocardial
106 infarction and pressure overload is distinct from SCA, we reasoned that further enhancing AMPK activity
107 could show functional cardiac benefits in the *in vivo* SCA model. To test this hypothesis, male and female
108 mice were divided into sham and arrest groups, with and without metformin pretreatment. Metformin-
109 treated mice were given 1 mg/mL of metformin in water for two weeks. There were no significant
110 differences among baseline animal characteristics, including weight, ratio of female mice, or EF (Table 1,
111 Figure 2A). Importantly, there were no differences among groups for body weight in these non-diabetic
112 mice. Twenty-four hours after surgery, untreated arrest mice had significantly lower EF than untreated
113 sham mice (sham: 59.5±1.7%; untreated arrest: 41.1±2.7%, p<0.0001, Figure 2A), as expected based on
114 our previous description of this model(31). Importantly, metformin pretreatment significantly improved
115 EF 24 hours post-SCA (arrest metformin: 51.6±2.6%, p<0.01 vs. untreated arrest) to a level not
116 significantly different from sham mice (Figure 2A). There were no changes to post-operative body
117 temperature, time to resuscitation, or 24-hour glucose levels between arrest groups (Table 1). The lack
118 of detectable changes in body weight or in the 24 hour post-SCA glucose suggests that differences in
119 systematic glucose handling was inadequate to explain later phenotypes.

120 The effects of whole-body ischemia/reperfusion injury can be detected in peripheral tissues,
121 particularly in the kidney (31). Unsurprisingly, untreated arrest mice had significant kidney damage
122 when compared to untreated sham mice at one day post-SCA. Kidney damage was quantified by serum
123 creatinine (sham: 0.36±0.04 mg/dL; untreated arrest: 1.49±0.14 mg/dL, p<0.0001, Figure 2B-C) and
124 blood urea nitrogen (BUN) levels (sham: 26.6±5.2 mg/dL; untreated arrest: 153.9±28.4 mg/dL, p<0.001),
125 and tubular injury score by histological analysis (sham: 0.11±0.04; untreated arrest: 3.33±0.24,
126 p<0.0001). The 1 mg/mL metformin-pretreated arrest mice had some improvement over untreated
127 arrest mice, with significantly reduced creatinine levels (metformin arrest: 0.92±0.24 mg/dL, p<0.05 vs.
128 untreated arrest). However, metformin treatment did not fully protect kidneys from SCA-induced
129 damage because serum creatinine was still significantly elevated compared to untreated sham (p<0.05

130 vs. sham), as was BUN (121.2 ± 28.0 mg/dL, $p < 0.05$ vs. sham). Similarly, histological analysis revealed an
131 increase in tubular injury score in metformin arrest mice (2.26 ± 0.24 , $p < 0.0001$ vs. sham) that was
132 partially improved compared to untreated arrest ($p < 0.01$; Figure 2B-C). Metformin pretreatment in
133 sham mice did not significantly affect creatinine levels (0.38 ± 0.06 mg/dL), BUN (28.8 ± 9.4 mg/dL), or
134 tubular injury score (0.14 ± 0.04 ; Figure 1C) relative to untreated sham. The time of renal ischemia was
135 unchanged between untreated and metformin-pretreated arrest groups (Figure 2D; Table 1), suggesting
136 the reperfusion differences were not a component of kidney protection.

137 Metformin dosage has been a concern for patients with preexisting kidney damage, especially
138 for those with chronic kidney disease, but lower doses have been found safe and avoid hyperlactatemia
139 (32,33). Furthermore, data have suggested that differences in dosage alter metformin targets, with low
140 doses activating AMPK, but higher doses causing mitochondrial Complex I inhibition(34). We therefore
141 treated an additional mouse cohort with a lower dose of metformin (0.2 mg/mL in drinking water as
142 opposed to 1 mg/mL in Figure 2) to reduce potential complications of decreased metformin clearance
143 after arrest. We found that the low-dose metformin cohort did not have significant cardio-protection
144 when compared to untreated arrest mice (EF of low-dose metformin: $50.7 \pm 3.3\%$, $p = 0.08$ vs. untreated
145 arrest) (Supplemental Figure 3A), though this cohort may not have been powered to detect cardiac
146 changes. However, the low-dose metformin arrest cohort did have lower serum creatinine (0.40 ± 0.05
147 mg/dL, $p < 0.001$ vs. untreated arrest) and BUN levels (64.83 ± 8.24 mg/dL, $p < 0.05$ vs. untreated arrest),
148 but unchanged tubular injury when compared to untreated arrest mice (2.04 ± 0.71 , $p = 0.08$ vs. untreated
149 arrest) (Supplemental Figure 3). These results suggest superior protection against kidney damage with
150 low-dose metformin over the higher dose metformin pretreatment, and protection against
151 ischemia/reperfusion injury by metformin need not be through mitochondrial respiration dysfunction.

152 **Metformin promotes AMPK activation in the LV following SCA**

153 As a marker of total AMPK activation, we assessed phosphorylation of threonine-172 of the
154 AMPK α subunit (35). One day after surgery, LVs from sham and arrest groups with and without
155 metformin pretreatment ($n = 6$ /group) were assessed for p-AMPK, total AMPK, and glyceraldehyde 3-
156 phosphate dehydrogenase (GAPDH) protein expression (Figure 2E). Surprisingly, metformin treatment
157 did not appear to cause activation of p-AMPK/AMPK in the sham mice (untreated sham: 1.00 ± 0.13 ;
158 metformin sham: 0.80 ± 0.11). However, we found significantly elevated p-AMPK/AMPK in metformin-
159 pretreated arrest mice (1.52 ± 0.14 AU) when compared to untreated sham (1.00 ± 0.13 AU, $p < 0.05$),
160 untreated arrest (0.70 ± 0.12 , $p < 0.001$), and metformin-pretreated sham mice (0.80 ± 0.11 , $p < 0.01$). p-
161 AMPK/GAPDH was similarly elevated in the LVs of metformin arrest mice (1.47 ± 0.08 AU) when
162 compared to untreated sham (1.00 ± 0.10 AU, $p < 0.05$), untreated arrest (0.90 ± 0.14 AU, $p < 0.01$), and
163 metformin-pretreated sham (0.98 ± 0.09 AU, $p < 0.05$). Total AMPK/GAPDH was not significantly changed
164 between groups. These data suggest that cardiac injury with metformin-pretreatment increases the
165 potential for AMPK activation.

166 **AMPK activation causes cardiac and renal protection after SCA and is necessary for metformin's** 167 **protection of EF**

168 Because metformin has multiple potential modes of action, we tested whether direct AMPK
169 activation was sufficient for the observed enhanced EF, and whether metformin benefit in SCA
170 depended on AMPK activity. To that end, male and female mice were divided into two groups: 1) AICAR

171 pretreatment, which activates AMPK (36), and 2) metformin-pretreatment combined with compound C,
172 an established AMPK inhibitor (13). Both groups underwent two weeks of intraperitoneal (IP) injections
173 prior to SCA and subsequent evaluation 24-hours after arrest. These groups were compared to the
174 untreated arrest and metformin pretreated arrest mouse results described above. One day after
175 surgery, AICAR pretreated arrest mice had significantly improved EF when compared untreated arrest
176 mice (AICAR arrest: $52.0 \pm 2.4\%$, $p < 0.05$ vs. untreated arrest, Figure 3A). Compound C not only prevented
177 the beneficial effects of metformin on post-SCA EF, but also caused significantly reduced EF (metformin
178 + C arrest: $30.0 \pm 2.9\%$) when compared to untreated arrest ($p < 0.05$). There was no change to body
179 weight, ratio of female mice, or baseline EF (Table 1 and Supplemental Figure 2) prior to arrest. The
180 effects of AICAR pretreatment phenocopies metformin pretreatment, and compound C blocks the
181 benefit of metformin strongly suggest that AMPK activation is necessary and sufficient for the
182 metformin-mediated protection of cardiac function after SCA. We cannot rule out the involvement of
183 involvement of glucose homeostasis and insulin sensitivity as contributing factors in this process. We
184 have not observed differences in random glucose and insulin levels, suggesting that such an effect would
185 be modest.

186 In the same cohort of mice, measures of kidney damage were significantly lower in AICAR-
187 pretreated mice than untreated arrest mice. Both creatinine (AICAR arrest: 0.67 ± 0.25 mg/dL, $p < 0.05$,
188 Supplemental Figure 4A) and tubular injury score (AICAR arrest: 1.45 ± 0.42 , $p < 0.01$) were significantly
189 improved relative to untreated arrest mice (creatinine 1.49 ± 0.14 mg/dL, tubular injury score 3.33 ± 0.24).
190 In contrast, there was no significant change when comparing renal outcomes in metformin-treated and
191 metformin + compound C-treated arrest mice (Supplemental Figure 4A). AICAR phenocopied the
192 metformin benefit in measures of kidney function and damage, suggesting a role for AMPK activation in
193 those processes. The limited change in these measures in the metformin + compound C cohort vs.
194 untreated arrest mice may suggest that at baseline AMPK activation is limited and cannot be further
195 reduced but would require further experimentation.

196 **AICAR promotes AMPK activation in the LV following SCA**

197 Twenty-four hours after surgery, LVs from arrest groups ($n=6$) were assessed for AMPK and p-
198 AMPK expression (Figure 3B). As expected, p-AMPK/AMPK was significantly elevated in AICAR-
199 pretreated arrest mice (AICAR arrest: 2.54 ± 0.59 AU) when compared to untreated arrest (1.00 ± 0.17 AU,
200 $p < 0.05$). p-AMPK/GAPDH and AMPK/GAPDH were not significantly changed in the AICAR group.
201 Metformin + compound C pretreated mice had lower p-AMPK/AMPK than the AICAR group (arrest
202 metformin + C: 0.52 ± 0.13 AU, $p < 0.05$) and p-AMPK/GAPDH (0.54 ± 0.11 AU, $p < 0.05$) but no significant
203 change from the untreated arrest group.

204 **Metformin affects mitochondrial morphology and markers of mitochondrial dynamics and autophagy**

205 Since ischemia/reperfusion is well known to induce mitochondrial damage, electron microscopy
206 was used to identify differences in mitochondrial perimeter, area, and circularity among in hearts of
207 untreated sham, untreated arrest, and metformin-pretreated arrest mice (Figure 4A). The untreated
208 arrest mice showed a decrease in mitochondrial perimeter and area (perimeter: 3.06 ± 0.06 μm ; area:
209 0.60 ± 0.02 μm^2 ; Figure 4B) when compared to sham (perimeter: 3.81 ± 0.09 μm ; area: 0.87 ± 0.04 μm^2 ,
210 $p < 0.0001$ for both measures). Metformin-pretreated arrest mice showed a modest but significant
211 increase of perimeter and area (perimeter: 3.33 ± 0.07 μm ; area: 0.70 ± 0.03 μm^2) when compared to
212 untreated arrest mice ($p < 0.05$ for both measures). Both the untreated arrest mice (0.76 ± 0.001 μm ,

213 $p < 0.0001$) and metformin-pretreated arrest mice ($0.75 \pm 0.01 \mu\text{m}$; $p < 0.0001$) had more circular
214 mitochondria than the untreated sham mouse mitochondria ($0.69 \pm 0.01 \mu\text{m}$). These data demonstrate
215 changes in mitochondrial morphology at 24 hours post-SCA, with improvements in mitochondrial area
216 and perimeter in metformin-pretreated arrest mice compared to untreated arrest mice.

217 We also looked at markers of mitochondrial abundance in the LVs. First, representative proteins
218 of mitochondrial respiratory complexes were assessed from LV extracts taken 24 hours after arrest from
219 untreated sham, untreated arrest, metformin-pretreated sham, and metformin-pretreated arrest mice.
220 There was a mild but significant elevation in relative complex II expression in the metformin-pretreated
221 arrest mice (1.29 ± 0.10 AU) compared to untreated sham mice (0.98 ± 0.04 AU, $p < 0.01$), but otherwise,
222 relative expression of representative proteins was unchanged across the groups (Figure 4C). Second, we
223 examined mtDNA relative abundance and damage in the cardiac tissue from these groups. Unlike in
224 failed hearts (37), the arrest group did not show a statistically significant decrease in mtDNA levels at 24
225 hours post-SCA. The metformin-pretreated arrest mice had slightly higher relative mitochondrial DNA
226 (mtDNA) levels (1.16 ± 0.06 AU) than untreated arrest mice (0.89 ± 0.05 AU, $p < 0.05$, Figure 4D), but were
227 not significantly different from the sham group (1.00 ± 0.05 AU). Interestingly, the metformin-pretreated
228 arrest mice had significantly less mtDNA damage (0.03 ± 0.1 lesions/10kb mtDNA) than untreated arrest
229 mice (0.59 ± 0.16 lesions, $p < 0.05$) as measured by long extension PCR assays. The small but significant
230 increases in mtDNA and complex II in the metformin-pretreated arrest mice are consistent with an
231 increase in mitochondrial biogenesis, but the significant improvement in mtDNA integrity suggests an
232 improvement in overall mitochondrial quality.

233 To better understand the alterations in mitochondrial morphology, we examined the relative
234 levels of several proteins involved in establishing mitochondrial shape. Mitofusin 2 (MFN2), a
235 mitochondrial outer membrane GTPase involved in fusion, had reduced expression in metformin-
236 pretreated arrest mice (0.53 ± 0.07 AU) compared to all other groups (sham: 1.00 ± 0.10 AU, $p < 0.01$;
237 arrest: 1.06 ± 0.07 AU, $p < 0.001$; sham metformin: 0.89 ± 0.04 AU, $p < 0.05$; Figure 5A). OPA1, also a
238 dynamin-related GTPase, resides in the inner membrane to perform fusogenic functions(38), and
239 followed a similar overall trend with significantly reduced levels in metformin-pretreated arrest mice
240 (0.75 ± 0.04 AU) when compared to untreated arrest mice (1.13 ± 0.10 AU, $p < 0.05$). In contrast, dynamin-
241 related protein 1 (DRP1), whose activity is regulated by its phosphorylation at Ser-616(39), showed no
242 significant change in p-DRP/DRP between sham and arrest groups. Metformin pretreatment appeared to
243 modestly improve p-DRP/DRP post-arrest but the effect was not statistically significant ($p = 0.08$). While
244 mitochondrial perimeter and area are increasing with metformin-pretreatment in SCA hearts, markers
245 of fusion decrease and fission is unchanged from sham or arrest. The MFN2 decrease may instead
246 suggest that other mitochondrial quality control processes contribute to the improved mitochondrial
247 morphology.

248 Interestingly, MFN2 decrease is consistent with the activation of mitophagy (40,41), due to its
249 ubiquitination and targeted proteolysis during autophagy. Metformin has been reported to impact heart
250 autophagy through AMPK signaling (42,43). To test engagement of autophagy in the SCA model, LVs
251 collected 24 hours after SCA from untreated sham, untreated arrest, metformin-pretreated sham, and
252 metformin-pretreated arrest mice were evaluated for markers of autophagy by western blot analysis.
253 First, we evaluated the mTOR signaling pathway, a crucial negative regulator of autophagy (44).
254 Consistent with prior findings (45), we found the marker of mTOR activity, p-mTOR (Ser-2448), to be
255 significantly reduced in untreated arrest (0.58 ± 0.08 AU, $p < 0.05$), metformin-pretreated sham (0.55 ± 0.08

256 AU, $p < 0.05$), and metformin-pretreated arrest mice (0.53 ± 0.07 AU, $p < 0.05$) when compared to sham
257 mice (1.0 ± 0.15 AU, Figure 5A). Total mTOR protein (mTOR/GAPDH) was only reduced in the metformin-
258 pretreated arrest group (0.71 ± 0.08 AU) when compared to sham (1.00 ± 0.05 AU, $p < 0.05$) and untreated
259 arrest (0.96 ± 0.06 AU, $p < 0.05$) groups. The ratio of p-mTOR to total mTOR was unchanged between
260 groups, but we focus on p-mTOR/GAPDH as a measure of activity. As positive downstream markers of
261 mTOR activity, S6 ribosomal protein (S6) total expression and phosphorylation at Ser-240/244 (p-S6)
262 were assessed. We found that p-S6 (pS6/GAPDH) was reduced in untreated arrest (0.41 ± 0.05 AU,
263 $p < 0.01$) and metformin-pretreated arrest mice (0.52 ± 0.07 AU, $p < 0.05$) when compared to sham
264 (1.00 ± 0.17 AU; Figure 5B). Total S6 was increased in the metformin sham mice compared to untreated
265 sham (1.40 ± 0.08 AU, $p < 0.05$) and decreased in metformin-pretreated arrest mice (0.80 ± 0.05 AU,
266 $p < 0.001$) when compared to untreated sham mice (1.00 ± 0.10 AU). p-S6/S6 was significantly reduced in
267 the untreated arrest mice (0.49 ± 0.08 AU) when compared to untreated sham mice (1.00 ± 0.17 AU,
268 $p < 0.05$). p-S6 levels are increased with metformin in sham mice, consistent with the metformin
269 activation of mTOR observed in other contexts (43). However, because mTOR activity based on p-mTOR
270 and pS6 are not significantly different between arrest and metformin arrest mice, the data suggest that
271 decreased mTOR activation alone is insufficient to explain the survival benefit of metformin.

272 Because metformin has been reported to increase cardiac mitophagy in cardiomyopathy
273 (42,43), we evaluated protein expression for markers associated with autophagosome formation,
274 including p62/Sequestosome 1, a cargo receptor associated with degradation of ubiquitinated proteins
275 (46), and LC3 processing. p62 expression (normalized to GAPDH) was significantly lower in the
276 metformin arrest mice (0.67 ± 0.09 AU) when compared to the untreated arrest group (1.22 ± 0.10 AU,
277 $p < 0.05$, Figure 5B). The relative levels of microtubule-associated protein light chain 3 (LC3), specifically
278 levels of uncleaved (LC3-I) and cleaved (LC3-II) forms, were also monitored as an indicator of changes in
279 autophagy initiation(47). Interestingly, the LC3-II to LC3-I ratio was significantly increased in untreated
280 arrest mice (1.91 ± 0.15 AU) compared to sham (1.00 ± 0.18 AU, $p < 0.01$), whereas the metformin-
281 pretreated arrest mice had significantly reduced LC3-II/LC3-I (1.17 ± 0.17 AU) when compared to
282 untreated arrest mice ($p < 0.05$, Figure 5B). LC3-I and LC3-II expression levels (normalized to GAPDH)
283 were not significantly changed between groups (Supplemental Figure 5). The restoration of p62 and LC3-
284 II/I to untreated sham levels in metformin-pretreated LVs after SCA appears to associate with the
285 improvements in the mitochondrial area and perimeter in the metformin arrest hearts vs. the sham
286 arrest hearts.

287 Taken together, these data suggest that increase autophagic flux contributes to the improved
288 mitochondrial network in metformin-pretreated arrest mice. Metformin increases the mitochondrial
289 perimeter and area despite the decrease in MFN2 and OPA1 fusogenic proteins. Further, there is an
290 increase in p62 and LC3-II in arrest hearts, which is consistent with impaired autophagic flux observed by
291 others(48). Notably, metformin pretreatment reduced p62 and LC3-II levels and significantly improved
292 mtDNA integrity after SCA, suggesting that autophagy in those hearts was also improved, which is
293 consistent with the observed lower level of mitochondrial fragmentation. Additional studies are
294 required to analyze flux in real-time.

295 **Metformin does not improve outcomes as a rescue therapy**

296 In *ex vivo* studies of ischemia-reperfusion, acute metformin administration just prior to stop-
297 flow mediated injury improved developed pressures during recovery(44). In ligation-mediated

298 ischemia/reperfusion experiments *in vivo*, 125 $\mu\text{g}/\text{kg}$ metformin injection into the LV lumen at the time
299 of reperfusion resulted in a decreased infarct area and improved EF in non-diabetic mice(45). In contrast
300 to coronary artery ligation experiments, which generally rely on >30 minute ischemic times and
301 generate significant cardiomyocyte death, our SCA model is 8 minutes of ischemia and lacks overt cell
302 death(29). In our SCA model, when metformin was given directly into the LV at resuscitation as a rescue
303 therapy (1,250 $\mu\text{g}/\text{kg}$), there was no change to EF at one day after SCA (arrest rescue metformin:
304 $42.1\pm 2.4\%$) compared to untreated arrest mice (Figure 6). Similarly, there was no change in creatinine
305 (arrest rescue metformin: 1.2 ± 0.25), BUN (arrest rescue metformin: 155.3 ± 14.2), or tubular injury score
306 (arrest rescue metformin: 2.45 ± 0.55) when compared to untreated arrest mice (Supplemental Figure
307 4B). Baseline EF was not significantly different between groups (Supplemental Figure 2). These data are
308 consistent with the notion that metabolic adaptation is required for the metformin protection after SCA.

309 **Metformin improves markers of cardiac and renal damage in humans one day after SCA**

310 Data supporting metformin use in cardiovascular disease are conflicting, often varying with the
311 particular condition or disease (21). To mirror our observations in mice, we focused in this retrospective
312 study on diabetic patients who survived to hospital care after resuscitation from SCA. Some of these
313 patients were taking metformin prior to SCA. We started with clinical data from 2,692 patients treated
314 at a single academic medical center, of whom 692 (26%) had a history of diabetes. We excluded 268
315 patients with chronic kidney disease, 20 who were transferred to our facility more than 24 hours post-
316 arrest, 56 for whom home medications were unknown, 7 who rearrested prior to labs being drawn, and
317 1 who was resuscitated with extracorporeal membrane oxygenation, leaving 341 patients in our final
318 analysis. Mean age was 65 ± 13 years, and 148 (43%) were female (Table 2). Overall, 140 (41%) patients
319 were prescribed metformin prior to arrest, 153 (45%) were prescribed insulin, and 92 (27%) were
320 prescribed other oral hyperglycemic medications. Serum troponin and creatinine were measured as part
321 of routine clinical care, typically at least once daily, and used to quantify heart and kidney injury,
322 respectively. We did not find reliable echocardiography data in this cohort. Median peak troponin in the
323 first 24 hours post-arrest was 1.4 (interquartile range, IQR: 1.0-1.7) and median peak creatinine was 1.4
324 (IQR: 1.0-2.0). We used generalized linear models (gamma distribution, log link) to test the independent
325 association of pre-arrest metformin use with peak troponin and creatinine within 24 hours after SCA,
326 adjusting for clinically relevant confounders including age, sex, arrest location (in- vs out-of-hospital),
327 witnessed collapse, layperson CPR, number of epinephrine doses administered, cardiac etiology of
328 arrest, and Charlson comorbidity index, as well as the use of insulin or other oral hyperglycemic
329 medications (Table 2). We handled the 2% missing data using multiple imputations. Metformin
330 prescription at the time of SCA was independently associated with lower 24-hour peak serum troponin
331 and lower 24-hour peak serum creatinine (Table 3). Without A1C levels, we cannot rule out whether
332 there are differences in glycemic control that associate with the improved cardiac and renal measures in
333 these patients.

334 **Discussion**

335 SCA is a common cardiac event for which there are extremely poor outcomes and no current
336 course-altering interventions. Metformin therapy, a first-line diabetes treatment, is beneficial in a
337 number of cardiovascular disorders(9,10)making it a candidate approach for SCA. Metformin impacts
338 many metabolic processes, including activation of the AMPK signaling pathway, decreased ER stress and
339 ROS, improved autophagy and mitochondrial biogenesis, and inhibition of the mitochondrial
340 permeability transport pore (mPTP) (49–53). Studies of metformin’s effects on cardiac function have
341 largely utilized ischemia-reperfusion injury models with ischemic periods of 25-30 minutes, both *in vivo*
342 and *ex vivo*. Metformin, applied as a therapy, decreases scar size in reversible coronary artery ligation
343 (21,28,53–56) or whole-heart ischemia-reperfusion (21,57), respectively. The long ischemic duration in
344 these models involves substantial cardiomyocyte necrosis, and much of metformin’s beneficial effect
345 has been attributed to reduction of mPTP-mediated cell death in the infarct border zone (49). In
346 contrast, our data demonstrates metformin’s protection of *in vivo* EF in a SCA model that features 8
347 minute ischemia period without evidence of cardiac cell death (31). Therefore, with a shorter ischemic
348 duration and no evidence of apoptosis, we have identified a unique pathway of metformin’s cardiac
349 protection.

350 The requirement of AMPK signaling for metformin’s cardiac benefits was confirmed by
351 downstream inhibition and rescue experiments. Specifically, metformin’s effects were negated by
352 concomitant treatment with the AMPK inhibitor compound C and were replicated by treatment with the
353 direct AMPK activator AICAR alone. Taken together, these data rigorously support that AMPK activation
354 is sufficient and necessary for cardiac protection.

355 Metformin pretreatment also protects against kidney injury in our model of SCA, reducing
356 creatinine levels, BUN, and tubular injury scores (Figure 2). As there is no change to ischemic duration in
357 the kidneys between untreated arrest and metformin-pretreated arrest mice, as measured by renal
358 artery doppler ultrasound (Table 1), this renal protection is likely intrinsic to the kidney rather than
359 secondary to the improved cardiac function. Previous studies have demonstrated beneficial effects of
360 metformin on renal performance after primary renal ischemic injury (58–60). Our model extends the
361 known impact of metformin on renal function by demonstrating comparable beneficial effects in a
362 model featuring renal ischemia secondary to cardiac arrest, more analogous to clinically observed renal
363 injury.

364 There exists concern about metformin therapy in the presence of kidney injury, as metformin
365 accumulation can lead to potentially lethal lactic acidosis (33). While we did not assess lactate build-up
366 or metformin accumulation in our model, we did treat an additional cohort of mice with a lower dose of
367 metformin (Supplemental Figure 3), comparable to concentrations of human doses, to address concerns
368 about supra-pharmacologic metformin concentrations (61). We found that low-dose metformin
369 conferred better renal protection than standard-dose metformin, as measured by creatinine levels after
370 arrest (Supplemental Figure 3B), though cardiac EF improvement did not reach significance in this group.
371 There has been some concern about the mechanism of action in high-dose vs. low-dose metformin
372 therapy; recent work has demonstrated that supra-pharmacologic doses of metformin effect change
373 through inhibition of mitochondrial ATP production, whereas pharmacological doses of metformin
374 increase mitochondrial respiration and fission (62). The literature would suggest that our high-dose
375 metformin (1 mg/mL) is a Complex I inhibitor, whereas low-dose metformin (0.2 mg/mL) activates

376 AMPK. Our work did not confirm or explore the mechanistic differences between these two doses,
377 which would be a natural extension of this work in the SCA mouse model.

378 Metformin therapy has pleiotropic effects and a large number of molecular pathways are
379 implicated in metformin's physiological effects, including reducing oxidative stress, inhibiting apoptosis,
380 complex I inhibition, and AMPK activation (19,21,34,63). Pathway analysis of microarray data from SCA
381 mice hearts compared to sham mice implicated AMPK signaling as the most significantly changed
382 pathway in our model (Figure 1C). The rescue of cardiac dysfunction by metformin and AICAR suggests
383 that additional AMPK activation is both possible and beneficial. Reciprocally, mice that were
384 concomitantly treated with metformin and the AMPK inhibitor compound C had significantly lower EF
385 than untreated arrest mice (Figure 3A). Unexpectedly, metformin + compound C mice had lower serum
386 creatinine than the arrest mice, though these studies will need to be confirmed in larger cohorts
387 (Supplemental Figure 4A). Despite this paradox, these data demonstrate that AMPK activation alone is
388 sufficient to cause at least cardioprotection in our SCA model, and that metformin's cardioprotective
389 benefits are negated by AMPK inhibition.

390 The downstream mechanisms underlying metformin's benefits in our model are difficult to fully
391 elucidate, as both metformin and AMPK signaling have been implicated in a wide number of signaling
392 cascades (21,27,64). However, we observed that the mitochondria in LVs of SCA mice are smaller and
393 rounder than sham cohorts by electron microscopy, and that metformin-pretreated arrest mice have
394 significantly larger mitochondria than untreated arrest mice (Figure 4), which suggest a possible role for
395 alterations in either mitochondrial dynamics or mitophagy as regulators of mitochondrial morphology.
396 Electron transport chain complex abundance estimates were largely unchanged between treatment
397 groups, except for a mild elevation in complex II expression in the metformin-pretreated arrest mice
398 over the sham cohort (Figure 4B). mtDNA copy number and damage were not significantly elevated in
399 the untreated arrest mice when compared to sham, but metformin pretreatment did drive an increase
400 in mtDNA copy number and a decrease in mtDNA damage when compared to the untreated arrest mice
401 (Figure 4D). These data could support an increase in mitochondrial biogenesis or a change in
402 mitochondrial turnover.

403 There were notable changes to markers of autophagy and mitochondrial dynamics, both of
404 which have been implicated as altered pathways related to AMPK signaling (64–67). We found evidence
405 of decreased expression levels of p-mTOR and p-S6 in untreated arrest, metformin sham, and metformin
406 arrest groups when compared to sham mice as indicated (Figure 5B), which could indicate increased
407 autophagic poise in the case of sham metformin hearts, and either poise or activation of autophagy in
408 the arrest and metformin-pretreated arrest samples. Importantly, an increase in LC3-II/LC3-I occurred in
409 arrest samples but was completely restored to baseline in metformin-pretreated arrest samples. P62,
410 another marker of the autophagosome, was not significantly changed in the untreated arrest vs sham
411 samples but was reduced in metformin-pretreated arrest mice compared to untreated arrest (Figure
412 5B). These findings are consistent with an increase in autophagy with arrest, which was improved by
413 either decreased mtDNA damage or increase autophagic clearance in the metformin arrest hearts.
414 Because mtDNA damage levels increased in metformin-pretreated arrest hearts beyond control levels,
415 the data suggest increased clearance of damaged mitochondria. SCA causes increased relative fission to
416 untreated sham in heart, but metformin, which improves mitochondrial morphology, does not increase
417 markers of fusion such as p-DRP1. It is unlikely that metformin pretreatment restores fusion, as MFN2
418 and OPA1 are both further depleted in metformin arrest mice relative to the arrest mice. Importantly,

419 depletion of MFN2 and OPA1 is associated with the induction of autophagy (68), suggesting that
420 clearance is enhanced in the metformin-treated SCA mice. Although this study lacks directed measures
421 of mitochondrial flux, the data are consistent with that metformin pretreatment decreased proteotoxic
422 stress in hearts and more efficient turnover of mitochondria and removal of damaged mtDNA, which
423 increase the overall quality of the mitochondria at 24 hours post-arrest.

424 While metformin pretreatment has been shown to be cardioprotective, it is thus far not a
425 clinically relevant model of therapeutic (post-arrest) intervention for SCA. There exists controversy
426 about the benefits of acute metformin therapy in cardiac ischemia and metformin's role as a rescue
427 therapy. Mouse studies of ischemia-reperfusion have demonstrated reduced infarct size when treated
428 with metformin at the time of coronary artery ligation(52) or after ischemic insult(69). However, these
429 benefits were not shown in a swine model of coronary artery ligation treated with metformin at the
430 time of reperfusion(70). We attempted to treat our SCA mice with acute metformin via left-ventricular
431 injection given at the time of reperfusion but found no evidence of cardiac (Figure 6) or renal
432 (Supplemental Figure 4B) protection in this model (Figure 6). Our data do not prove, but are supportive
433 of the idea, that metabolic adaptation is necessary to convey protection in a preventative setting.

434 To evaluate the clinical relevance of the cardiac and renal protection demonstrated with
435 metformin pretreatment in our SCA mouse model, we performed a retrospective analysis of diabetic
436 patients resuscitated from cardiac arrest. We divided these patients into metformin-treated and non-
437 metformin-treated patients. Our primary endpoints were peak serum troponin within 24 hours post-
438 arrest as a marker of cardiac damage and peak serum creatinine within the first 24 hours as a marker of
439 kidney damage. Multiple regression analyses demonstrated a significant association between metformin
440 pretreatment and decreased peak serum troponin and peak serum creatinine levels (Table 3). This
441 suggests that metformin is driving cardiac and renal protection independently of other baseline
442 characteristics. Previous studies have noted that improved renal function after arrest is predictive of
443 long-term outcomes (7,8). It is not clear, however, whether or not early cardiac outcomes are predictive
444 of survival (6,69). Our analysis was not powered to look for survival benefit in human subjects, but such
445 a study is warranted.

446 In summary, we provide evidence in both a mouse model and retrospective clinical study that
447 metformin pretreatment offers significant cardiac and renal protection acutely after SCA. Our SCA
448 mouse involves cardiac injury without cardiac cell death, making it a more relevant system in which to
449 study metformin's mechanism of action. Direct AMPK activation and inhibition studies confirmed that
450 AMPK activation is necessary and sufficient for the cardiac and renal benefits observed with metformin
451 treatment. A number of molecular mechanisms exist downstream of AMPK activation, and we have
452 implicated changes to autophagy and mitochondrial dynamics in our mouse studies, though there were
453 no dramatic changes to mitochondrial morphology or electron transport chain subunit expression.
454 Future studies are warranted to investigate specific pathways downstream of metformin and AMPK that
455 may be useful as an acute rescue therapy to replicate the cardiac and renal protection seen in our model
456 and patient studies.

457

458

459 **Methods**

460 **Pre-clinical Data**

461 **Sudden Cardiac Arrest Model**

462 Eight-week-old male and female C57BL/6J mice (Jackson Labs, Bar Harbor, ME, #000664)
463 underwent cardiac arrest by delivery of potassium chloride (KCl) directly into the LV by percutaneous,
464 ultrasound-guided needle injection as previously described (31). Briefly, mice were anesthetized using
465 vaporized isoflurane (Henry Schein, Melville, NY, #1182097) and endotracheally intubated, then
466 mechanically ventilated (MiniVent, Harvard Apparatus, Holliston, MA, #73-0043) at a rate of 150 bpm
467 and volume of 125 μ L for females and 140 μ L for males. Body temperature was maintained by utilizing a
468 rectal temperature probe and heating pad (Indus Instruments, Webster, TX, #THM150). The chest was
469 cleaned of hair using Nair and sterilized with betadine prior to introduction of a 30-gauge needle into
470 the LV under ultrasound guidance (Visual Sonics Vevo 3100 with Vevo LAB v 5.5.1 software, Toronto,
471 Canada), followed by delivery of 40 μ L of 0.5M KCl to induce asystole. The ventilator was discontinued,
472 and mice remained in asystole for a total of 8 minutes. 7.5 minutes after KCl dosing, 500 μ L of 15 μ g/mL
473 epinephrine in saline (37°C) was injected into the LV over approximately 30 seconds. At 8 minutes, CPR
474 was initiated by finger compression at about 300 bpm for 1-minute intervals. Electrocardiogram (ECG)
475 was evaluated for return of sinus rhythm after each 1-minute interval. Animals not achieving ROSC by 3
476 minutes after CPR initiation were euthanized. Mice remained on the ventilator until breathing frequency
477 was greater than 60 times per minute. Sham mice received no KCl, but rather a single injection of
478 epinephrine. All animals were placed in a recovery cage under a heat lamp after the procedure.

479 **Animal Treatment Groups**

480 Treatment groups in the study were as follows: untreated sham, untreated cardiac arrest,
481 metformin-pretreated sham, metformin-pretreated cardiac arrest, 5-aminoimidazole-4-carboxamide-1-
482 beta-D-ribofuranoside (AICAR) pretreated cardiac arrest, metformin + compound c pretreated cardiac
483 arrest, metformin rescue treatment cardiac arrest, and low-dose metformin pretreatment cardiac
484 arrest. Metformin pretreatment consisted of *ad libitum* access to metformin (Major Pharmaceuticals,
485 Livonia, MI, #48152) in drinking water (1 mg/mL) for 14 days prior to surgery. AICAR pretreated mice
486 were given IP injections of 500 mg/kg AICAR (Toronto Research Chemicals, Toronto, CA, #A611700) in
487 saline every other day for 14 days prior to surgery. Compound C pretreated mice were given 20 mg/kg IP
488 injections of compound C (Cayman Chemical, Ann Arbor, MI, #11967) in saline daily for 14 days prior to
489 arrest. Metformin rescue therapy was given as a single direct LV injection (1250 μ g/kg) dissolved in
490 saline along with 500 μ L of 1mg/mL epinephrine (Par Pharmaceutical, Chestnut Ridge, NY, #10977) at
491 the time of resuscitation. Low-dose metformin pretreatment consisted of *ad libitum* access to
492 metformin in drinking water (0.2 mg/mL) for 14 days prior to surgery.

493 **Echocardiography and Ultrasound**

494 Immediately prior to arrest surgery, mice were evaluated by transthoracic echocardiography
495 using Vevo 3100 imaging systems (Visual Sonics) with a 40MHz probe. Repeat echocardiography was
496 performed one day after arrest under isoflurane anesthesia delivered by nose cone. Heart rate was
497 maintained between 400-500 bpm during imaging by adjusting isoflurane concentration. B-mode images
498 taken from the parasternal long-axis were captured and LV EF calculated using modified Simpson's

499 methods (70). A cohort of all groups were assessed for renal perfusion after resuscitation. The
500 ultrasound probe was oriented transversely across the abdomen at the plane of the right kidney and
501 monitored for renal artery blood flow by doppler imaging. Image analysis was performed by a blinded
502 sonographer (Vevo Lab 5.5.1, Visual Sonics).

503 **Tissue and Serum Collection**

504 After euthanasia with isoflurane and cervical dislocation, mice underwent cardiac puncture for
505 collection of blood by heparinized syringe. Blood was separated by centrifugation at 2,000 x g at 4°C for
506 10 minutes and the serum was flash frozen. These samples were evaluated for blood urea nitrogen
507 (BUN), creatinine, alanine aminotransferase (ALT), and creatine kinase (CK) by the Kansas State
508 Veterinary Diagnostic Laboratories (Manhattan, KS). Hearts from the mice were excised and LVs were
509 isolated and flash frozen.

510 **Western Blot**

511 Frozen LV tissue was homogenized in lysis buffer containing a protease/phosphatase cocktail
512 (Sigma-Aldrich, St. Louis, MO, #11697498001) and normalized for protein content using a BCA assay (Life
513 Technologies, Carlsbad, CA, #23235). Samples were separated on NuPage 4-12% gradient SDS-PAGE gels
514 (ThermoFisher, Waltham, MA, #WG1403BOX) and transferred onto iBlot nitrocellulose membranes
515 (Invitrogen, #IB301001). Membranes were blocked in 5% milk for 1 hour and then incubated overnight
516 at 4°C with primary antibodies, including OxPhos Rodent Antibody Cocktail 1:5000 (ThermoFisher,
517 #458099), p-AMPK (Thr172, 1:1000, Cell Signaling, Danvers, MA, #2535), AMPK (1:1000, Cell Signaling,
518 #2532), p-mTOR (Ser2448, 1:1000, Cell Signaling, #2971), DRP1 (1:1000, Cell Signaling, #5391), pDRP1
519 (Ser616, 1:1000, Thermo, #PA5-64821), mTOR (1:1000, Cell Signaling, #2972), p-S6 (Ser240/244, 1:1000,
520 Cell Signaling, #2215), S6 (1:1000, Cell Signaling, #2217), p-AKT (Ser473, 1:1000, Cell Signaling, #9271),
521 AKT (1:1000, Cell Signaling, #9272), p62 (1:1000, Sigma-Aldrich, #P0067), and GAPDH (1:5000, Millipore,
522 St. Louis, MO, #AB2302). Following incubation, membranes were washed with TBS-tween and then
523 probed for 1 hour at room temperature with anti-mouse or anti-rabbit secondary antibodies (Jackson
524 ImmunoResearch, West Grove, PA, #115-035-003 and #115-035-144). Images were obtained by
525 developing on a ChemiDoc XRS imaging system (BioRad, Hercules, CA) and analyzed using ImageJ
526 software (National Institutes of Health, Bethesda, MD).

527 **Tissue Histology**

528 Kidneys were fixed overnight in 10% formalin (Thermo, #SF100) at 4°C, then washed with PBS
529 and transferred to 70% ethanol at room temperature. After fixation and dehydration, tissues were
530 embedded in paraffin prior to sectioning at 4 µm by the Histology Core at the Children's Hospital of
531 Pittsburgh. Sections were stained with hematoxylin and eosin (H&E). Renal tubular pathology was semi-
532 quantitatively scored (0: no injury to 4: severe injury) in terms of tubular dilatation, formation of
533 proteinaceous casts, and loss of brush border(71). Histological scoring was performed in a blinded
534 fashion at 40x magnification on outer medullary regions of the tissue sections. Eight fields were
535 evaluated per kidney (n=6-8 animals/group). Samples were imaged using a Leica DM 2500 microscope
536 (Leica, Wetzlar, Germany) and analyzed with LAS X software (Leica).

537 **Transmission Electron Microscopy**

538 LV tissue was immersion fixed in 2.5% glutaraldehyde in 0.1M PBS overnight at 4°C. Fixed
539 samples were washed 3x in PBS then post-fixed in aqueous 1% OsO₄, 1% K₃Fe(CN)₆ for 1 hour at room

540 temperature. Following 3 PBS washes, the pellet was dehydrated through a graded series of 30-100%
541 ethanol, and then 100% propylene oxide then infiltrated in 1:1 mixture of propylene oxide:Polybed 812
542 epoxy resin (Polysciences, Warrington, PA) for 1 hour. After several changes of 100% resin over 24
543 hours, the samples were embedded in molds and cured at 37°C overnight, followed by additional
544 hardening at 65°C for two more days. Semi-thin (300 nm) sections were heated onto glass slides, stained
545 with 1% toluidine blue and imaged using light microscopy to assure proper tissue orientation. Ultrathin
546 (60-70 nm) sections were collected on 100 mesh copper grids, and stained with 4% uranyl acetate for 10
547 minutes, followed by 1% lead citrate for 7 min. Sections were imaged using a JEOL JEM 1400 PLUS
548 transmission electron microscope (Peabody, MA) at 80 kV fitted with a side mount AMT digital camera
549 (Advanced Microscopy Techniques, Danvers, MA). Twenty random images were obtained from sections
550 throughout each LV. Individual mitochondria (n=50/sample) were randomly selected and traced for
551 blinded quantification of size, roundness, and density via ImageJ (v1.8.0) software(72).

552 **RNA Isolation, qPCR, and Microarray Analysis**

553 Total genomic and mitochondrial DNA were isolated from frozen, powdered LV tissue. Tissue
554 was homogenized in a buffer containing Proteinase K digestion buffer and proteinase K (Genesee
555 Scientific, Genesee, NY, #42-700) overnight at 55°C, as previously described (73,74). The next day, DNA
556 was purified by centrifuging the homogenized buffer with staged EtOH resuspension, followed by
557 centrifugation and resuspension in TE buffer supplemented with RNase A (Invitrogen, #12091039). DNA
558 concentration was measured using an AccuBlue Broad Range kit (Biotium, Fremont, CA, #31009).
559 Relative mtDNA abundance was measured using a TaqMan primer/probe for mitochondrial ND1 (VIC-
560 labeled; primer:probe 1:1) versus nuclear HDAC1 (FAM-labeled; primer:probe of 3:1; supplemental
561 Table 1). Multiplex assessment of relative abundance of mtDNA was quantified using real-time
562 quantitative PCR (qPCR) with TaqMan Fast Advanced Master Mix (ThermoFisher, #4444965) performed
563 on QuantStudio 5 PCR system (Applied Biosystems). 4.6 ng of DNA was included in each reaction with 5
564 µM of primers/probes in a 10 µL total reaction volume and calculated by the $\Delta\Delta Cq$ method (75). The
565 qPCR amplification profile was: one cycle (95°C for 20 seconds) followed by 40 cycles (95°C for 1 sec
566 then 60°C for 20 seconds). All primers and probes were produced by Integrated DNA Technologies
567 (Coralville, Iowa).

568
569 mtDNA damage was assessed by comparing total PCR product after long-amplicon mtDNA
570 replication with LongAmp Hot Start Taq Polymerase (New England Biolabs, Ipswich PA, #M0533;
571 Supplemental Table 1)(76). 15 ng of DNA was amplified using the following profile: one cycle (94°C for 2
572 minutes) followed by 17 cycles (94°C for 15 seconds then 64°C for 12 seconds) then 1 cycle (72°C for 10
573 min)(77). Final product was measured by Accublu within the linear detection range. Specific DNA
574 products were confirmed by gel electrophoresis.

575
576 Microarray analysis was performed on cDNA through the Affymetrix microarray analysis service
577 (ThermoFisher). Eight untreated sham and eight untreated arrest mice were included in these studies
578 with an even distribution of males and females. Differential gene expression analysis was performed
579 using Transcription Analysis Console (Thermofisher). Gene-level p-values less than 0.05 were considered
580 significant for gene inclusion. Subsequent pathway analysis was performed to compare untreated sham
581 and arrest groups through Ingenuity Pathway Analysis (Qiagen). Complete datasets were deposited in
582 GEO (accession no. GSE176494).

583 584 **Statistical Analysis** 585

586 Data were expressed as mean \pm standard error in all figures. $p \leq 0.05$ was considered significant
587 for all comparisons. One-way ANOVA with either Dunnett's multiple comparisons test or Tukey's
588 multiple comparisons post-hoc analysis was used to compare groups either to a single group or all
589 groups as detailed in figure legends. All statistical analysis was completed using Graphpad Prism 8
590 software (San Diego, CA).

591 **Clinical Data**

592

593 **Clinical Data Collection**

594

595 De-identified adult patients with a documented history of diabetes mellitus treated at a single
596 academic medical center after resuscitation from cardiac arrest from January 2010 to December 2019
597 were included in this study. Patients with a history of known kidney disease prior to arrest, patients who
598 arrived at our facility over 24 hours after collapse, patients for whom home medications were unknown,
599 patients who rearrested and died before blood work could be acquired, and patients resuscitated with
600 extracorporeal support were excluded from analysis. Demographic and arrest characteristics from a
601 prospective registry, including patient age, sex, shockable presenting arrest rhythm, witnessed arrest,
602 layperson CPR, arrest duration, number of epinephrine doses administered, cardiac etiology of arrest,
603 and Charlson Comorbidity Index were extracted. Admission pharmacy documentation in each patient's
604 electronic medical record was evaluated to determine whether patients were prescribed metformin,
605 insulin, or other oral antihyperglycemic medications prior to their arrest, and classified each of these as
606 three independent binary predictors. The primary outcomes of interest were peak serum creatinine and
607 peak serum troponin at 24 hours post-arrest.

608 **Clinical Statistical Analysis**

609 Baseline population characteristics and outcomes were summarized using descriptive statistics.
610 Multiple imputation with chained equations with predictive mean matching was used to impute missing
611 continuous variables. Generalized linear models with a gamma distribution and log link were used to
612 test the association of metformin with peak 24-hour serum creatinine and peak 24-hour serum
613 troponin. For primary adjusted analysis, covariates were included based on biological plausibility. A
614 series of sensitivity analyses, including a backward stepwise model sequentially removing predictors
615 with $p < 0.1$, complete case analysis, and adjusted modeling were performed, excluding patients receiving
616 insulin who may be fundamentally sicker than those receiving no medications or oral antihyperglycemics
617 only.

618 **Study Approvals**

619 All mouse studies were performed at the University of Pittsburgh in compliance with the
620 National Institutes of Health Guidance for Care and Use of Experimental Animals. This protocol was
621 approved by the University of Pittsburgh Animal Care and Use Committee (Protocol #18032212). The
622 University of Pittsburgh Human Research Protection Office approved all aspects of the reported human
623 subject's research with a waiver of informed consent due to its no greater than minimal risk to
624 participants (STUDY19020205).

625 **Author Contributions**

626 CR, CD, and BK were responsible for conceptualization of these studies. CR designed the methodology
627 and with CL and KR performed the investigation. CR and JE were responsible for data curation. Formal
628 analyses were performed by CR, CL, TC, SSL, CD, and JE. Visualization was completed by TC and DS. CR,
629 SSL, CD, JE, and BK wrote the manuscript. SSL and BK supervised the project, and BK provided resources
630 for its completion. All authors reviewed the final manuscript and are responsible for its integrity.

631

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638

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- 843
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845 **Figures and figure legends**

846

847 **Figure 1. Mouse model of sudden cardiac arrest (SCA) and Microarray Pathway Analysis.** A) Cartoon
848 representation of direct left ventricular injection of potassium chloride (KCl) to cause asystole with
849 representative ultrasound image of needle guidance. B) Time course of SCA protocol. C) Pathway
850 analysis of microarray data from left ventricles collected one day after SCA (n=8) versus sham (n=8)
851 surgeries, demonstrating the ten most significantly changed canonical signaling pathways by Ingenuity
852 Pathway Analysis (IPA). LV, left ventricle; ROSC, return of spontaneous circulation.

853

854 **Figure 2. Metformin treated mice have preserved ejection fraction (EF) and lower kidney damage than**
855 **untreated mice one day after sudden cardiac arrest (SCA).** A) At baseline, there is no difference in EF
856 between treatment groups. One day after SCA, EF in arrest mice (n=20) is significantly lower than sham
857 mice (n=15). Metformin pretreatment did not change EF in mice receiving sham surgery (n=8), but
858 metformin pretreatment did lead to higher EF at 24 hours post-SCA (n=20) when compared to untreated
859 arrest mice. B) Representative histologic sections from untreated sham and untreated arrest mice
860 demonstrating proteinaceous casts in renal tubules (black stars) and infiltrates (white arrowheads) with
861 glomeruli marked (white X's). Scale bar = 50 μ M. C) Markers of kidney damage, including serum
862 creatinine, blood urea nitrogen (BUN), and histologic tubular injury score demonstrate significant injury
863 in untreated arrest mice. Untreated sham (n=8) and metformin treated sham (n=6) mice have no
864 evidence of damage. Metformin treated arrest mice (n=7) have significantly lower creatinine and tubular
865 injury score than untreated arrest mice (n=8). D) There is no change in renal ischemic duration between
866 untreated arrest and metformin treated arrest mice. E) Western blot analysis of pAMPK/AMPK in arrest
867 mice pretreated with metformin when compared to sham, untreated arrest, and metformin-pretreated
868 sham groups demonstrating increased p-AMPK/AMPK and p-AMPK/GAPDH (n=6 for all groups). Data are
869 expressed as mean \pm SEM. P-values: * < 0.05, ** < 0.01, *** < 0.001 by one-way ANOVA with Tukey post-
870 hoc analysis. EF, ejection fraction; BUN, blood urea nitrogen.

871

872 **Figure 3. AMPK activation alone improves cardiac outcomes after SCA, and AMPK activation is**
873 **necessary to exert metformin's cardioprotection.** A) SCA mice pretreated with the AMPK-activator
874 AICAR (Arrest AICAR; n=9) have improved EF when compared to untreated arrest mice (n=15; data
875 presented in Figure 2). Mice pretreated with both metformin and the AMPK-inhibitor Compound C
876 (Arrest Met + Comp C; n=9) have decreased EF when compared to untreated arrest mice. B) AICAR
877 treatment causes significant p-AMPK/AMPK elevation when compared to untreated arrest mice (n=6 for
878 all groups). Data are expressed as mean \pm SEM. P-values: * < 0.05, ** < 0.01, *** < 0.001 by one-way
879 ANOVA with Dunnett's post-hoc analysis (A) or Tukey post-hoc analysis (B). AICAR, 5-aminoimidazole-4-
880 carboxamide-1- β -D-ribofuranoside; Comp C, Compound C; EF, ejection fraction.

881

882 **Figure 4. Metformin affects mitochondrial characteristics after SCA.** A) Representative electron
883 microscope images of intrafibrillar mitochondria in untreated sham, untreated arrest, and metformin
884 pretreated arrest mice one day after surgery (40,000x magnification). B) Mitochondria are smaller and

885 more circular in untreated arrest and metformin treated arrest mice when compared to sham, and
886 metformin treated arrest mice have larger mitochondria than untreated arrest mice (n=50 per group). C)
887 Electron transport chain expression is largely unchanged between treatment groups, with the exception
888 of complex II (CII) expression in metformin-pretreated arrest mice being significantly higher than
889 untreated sham mice one day after arrest (n=6/group, normalized to ponceau stain). D) mtDNA copy
890 number is increased and has less damage in metformin treated arrest mice (n=7) when compared to
891 untreated arrest mice (n=5). Untreated arrest mice had no changes to mtDNA copy number or damage
892 when compared to sham. Data are expressed as mean \pm SEM. P-values: * < 0.05, ** < 0.01, *** < 0.001 by
893 one-way ANOVA with Tukey post-hoc analysis. ND1, NADH dehydrogenase 1; HDAC1, Histone
894 deacetylase 1; mtDNA, mitochondrial DNA; WT, wild-type.

895

896 **Figure 5. Metformin pretreatment affects autophagy and mitochondrial dynamics after sudden**
897 **cardiac arrest (SCA).** A) Representative western blot images of markers of mitochondrial fission and
898 fusion. Mitofusin 2 (MFN2) is significantly depressed in metformin treated arrest mice (n=6) compared
899 to untreated sham (n=6), untreated arrest (n=6), and sham metformin (n=5) groups. OPA1 is significantly
900 depressed in metformin treated arrest compared to untreated arrest mice. There is no significant
901 change to p-DRP expression between groups. B) Representative western blot images of autophagy
902 related proteins downstream of AMPK. p-mTOR expression is reduced in untreated arrest, metformin
903 treated sham, and metformin treated arrest mice when compared to untreated sham, and metformin
904 treated mice have lower total mTOR than untreated sham and arrest groups. p-S6, a marker of mTOR
905 activity, is also reduced in untreated arrest and metformin treated arrest mice when compared to sham.
906 S6 expression is increased in sham metformin mice, but depressed in metformin arrest mice compared
907 to sham, while the p-S6/S6 ratio is only significantly changed in the untreated arrest mice when
908 compared to sham. P62, a marker of autophagosome formation, is lower in metformin treated arrest
909 mice compared to untreated arrest mice. Untreated arrest mice have significantly elevated LC3II/LC3I, a
910 marker of autophagy, compared to sham mice and metformin treated arrest mice (n=6/group). Data are
911 expressed as mean \pm SEM. P-values: * < 0.05, ** < 0.01, *** < 0.001 by one-way ANOVA with Tukey post-
912 hoc analysis for all groups, except for the p-S6 analyses, which used Dunnett's post-hoc analysis. DRP,
913 dynamin-related protein1; LC3, microtubule-associated protein light chain; MFN2, mitofusin 2; mTOR,
914 mechanistic target of rapamycin; OPA1, dynamin-like 120 kDa protein, mitochondrial.

915

916 **Figure 6. Metformin does not work as a rescue therapy following arrest.** A) When given concomitantly
917 with epinephrine, intravenous metformin (n=6) did not demonstrate any change to ejection fraction (EF)
918 when compared to untreated arrest mice (n=15; data presented in Figure 2) or metformin treated arrest
919 mice (n=20, data presented in Figure 2). P-values: * < 0.05 by one-way ANOVA with Tukey post-hoc
920 analysis.

921

922 **Figure 7. Summary of clinical inclusion and exclusion data for retrospective analysis of the Pittsburgh**
923 **Post-Cardiac Arrest Service patient database.** Primary outcomes evaluated included peak serum
924 troponin and peak serum creatinine in diabetic patients with and without a history of metformin
925 therapy.

926

927 **Figure 8. Distributions of peak 24-hour serum troponin and peak 24-hour serum creatinine for diabetic**
928 **patients with and without metformin therapy prior to arrest.** Dashed lines represent median values
929 and dotted lines represent upper and lower quartiles.

930

931

932

933 **Table Legends**

934

935 **Table 1. Surgical data for arrest mice.** There are no significant changes to body weight or body
936 temperature at time of extubation between treatment groups. There is no change in time to return of
937 spontaneous circulation (ROSC) or time to extubation between untreated arrest and treated arrest
938 groups. There is no change to random blood glucose 24-hours after arrest between sham, untreated
939 arrest, and metformin pretreated arrest mice. Data are presented as mean \pm SEM unless otherwise
940 noted. Analysis by one-way ANOVA with Tukey post-hoc analysis.

941

942 **Table 2. Baseline demographics and clinical characteristics.** Data are presented as mean \pm standard
943 deviation, median [interquartile range], or sample number (corresponding percentage).

944

945 **Table 3. Association between metformin use and peak serum creatine and troponin by log link model.**
946 Model was adjusted for age, sex, arrest location (in- vs out-of-hospital), witnessed collapse, layperson
947 cardiopulmonary resuscitation, presenting rhythm, arrest duration, number of epinephrine
948 administered, cardiac etiology of arrest, Charlson Comorbidity index, insulin, and other oral diabetic
949 medications. Both peak serum creatinine level and peak serum troponin level 24 hours post-arrest are
950 significantly associated with history of metformin use.

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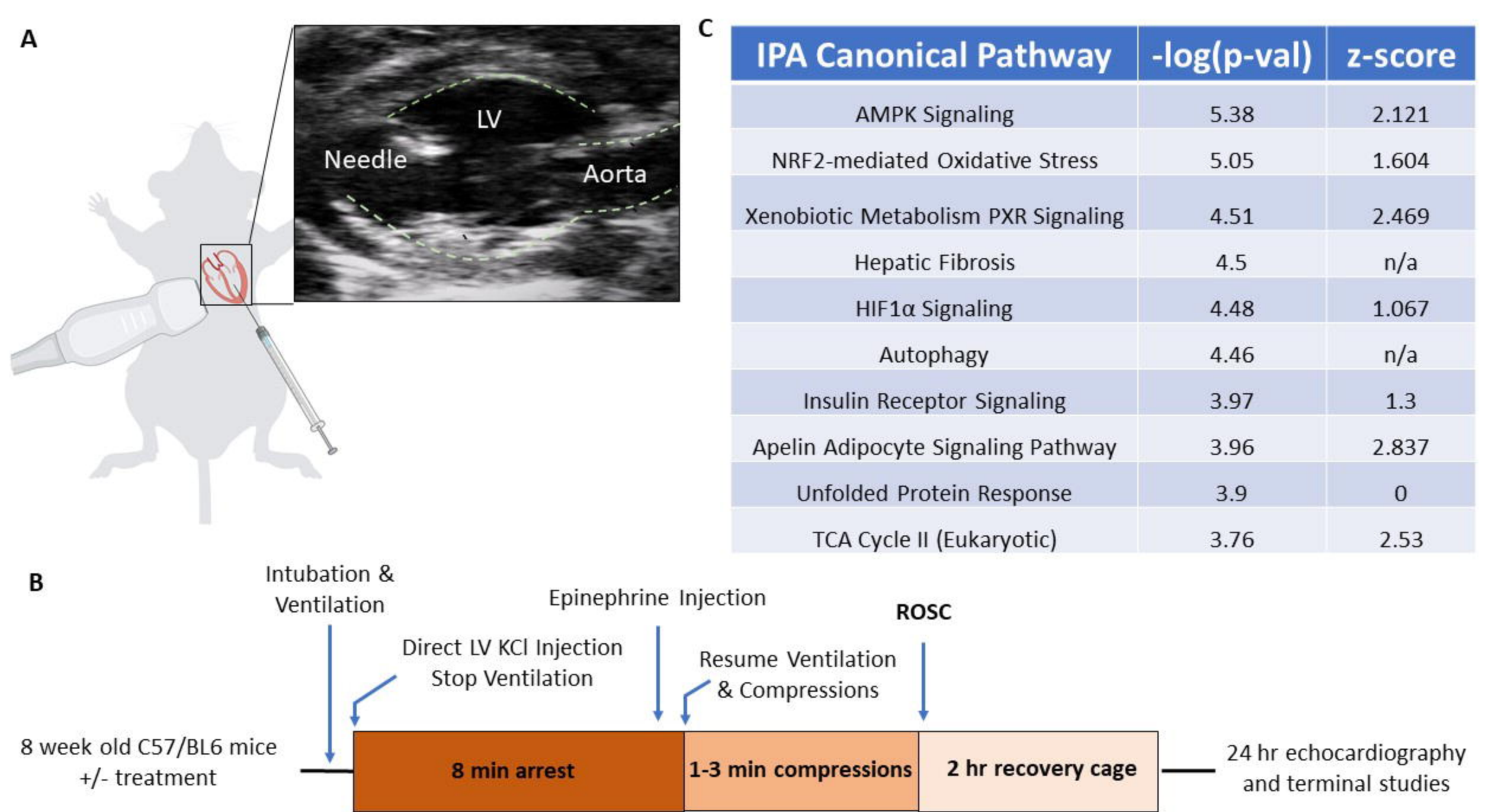


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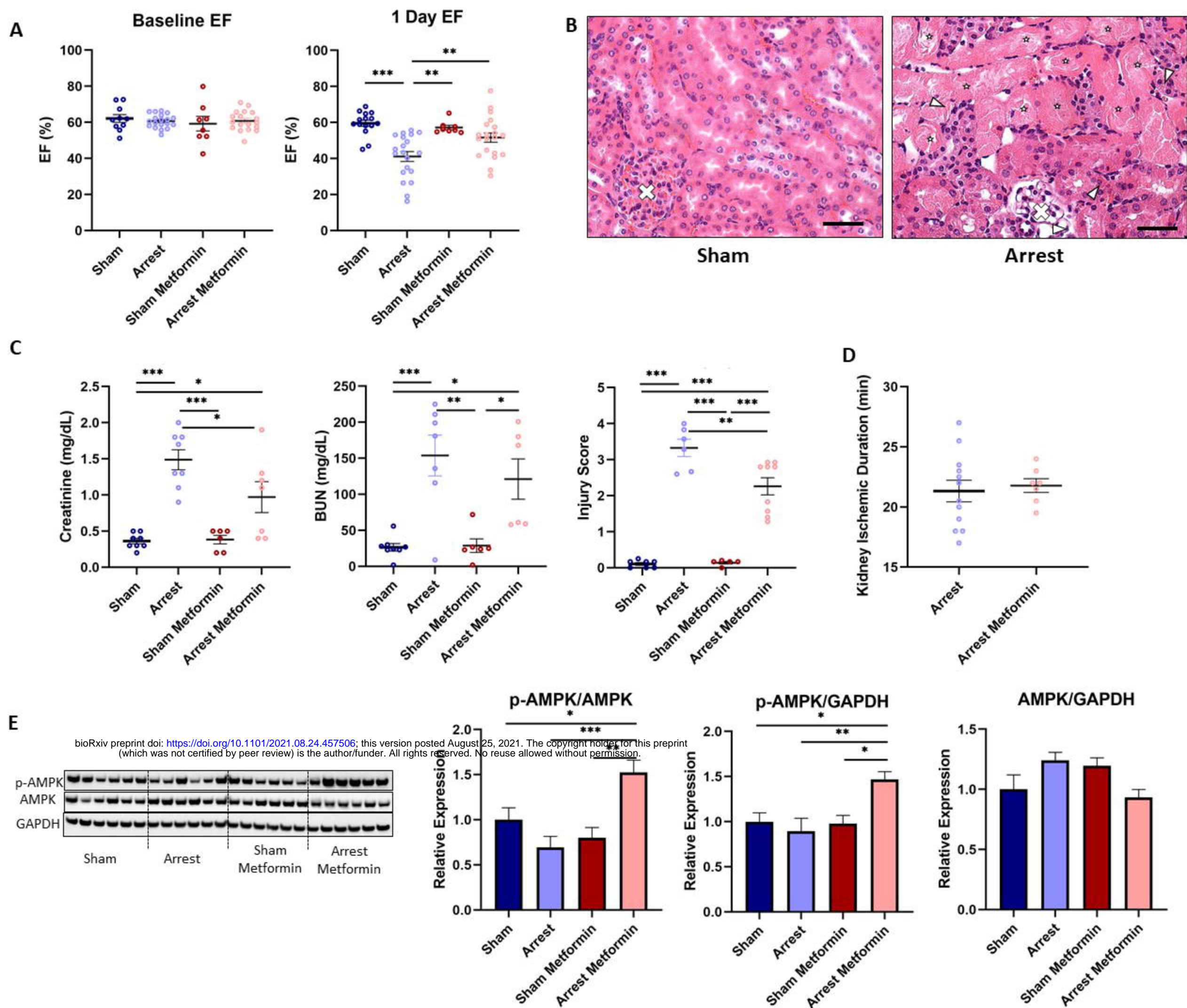
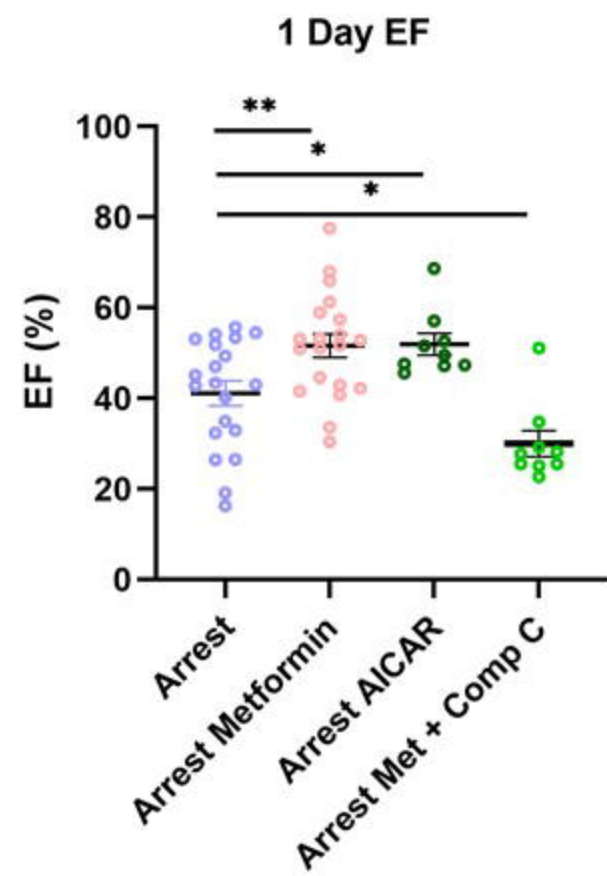


Figure 2. Metformin treated mice have preserved ejection fraction (EF) and lower kidney damage than untreated mice one day after sudden cardiac arrest (SCA). A) At baseline, there is no difference in EF between treatment groups. One day after SCA, EF in arrest mice (n=20) is significantly lower than sham mice (n=15). Metformin pretreatment did not change EF in mice receiving sham surgery (n=8), but metformin pretreatment did lead to higher EF at 24 hours post-SCA (n=20) when compared to untreated arrest mice. B) Representative histologic sections from untreated sham and untreated arrest mice demonstrating proteinaceous casts in renal tubules (black stars) and infiltrates (white arrowheads) with glomeruli marked (white X's). Scale bar = 50 μ M. C) Markers of kidney damage, including serum creatinine, blood urea nitrogen (BUN), and histologic tubular injury score demonstrate significant injury in untreated arrest mice. Untreated sham (n=8) and metformin treated sham (n=6) mice have no evidence of damage. Metformin treated arrest mice (n=7) have significantly lower creatinine and tubular injury score than untreated arrest mice (n=8). D) There is no change in renal ischemic duration between untreated arrest and metformin treated arrest mice. E) Western blot analysis of pAMPK/AMPK in arrest mice pretreated with metformin when compared to sham, untreated arrest, and metformin-pretreated sham groups demonstrating increased p-AMPK/AMPK and p-AMPK/GAPDH (n=6 for all groups). Data are expressed as mean \pm SEM. P-values: * < 0.05, ** < 0.01, *** < 0.001 by one-way ANOVA with Tukey post-hoc analysis. EF, ejection fraction; BUN, blood urea nitrogen.

A



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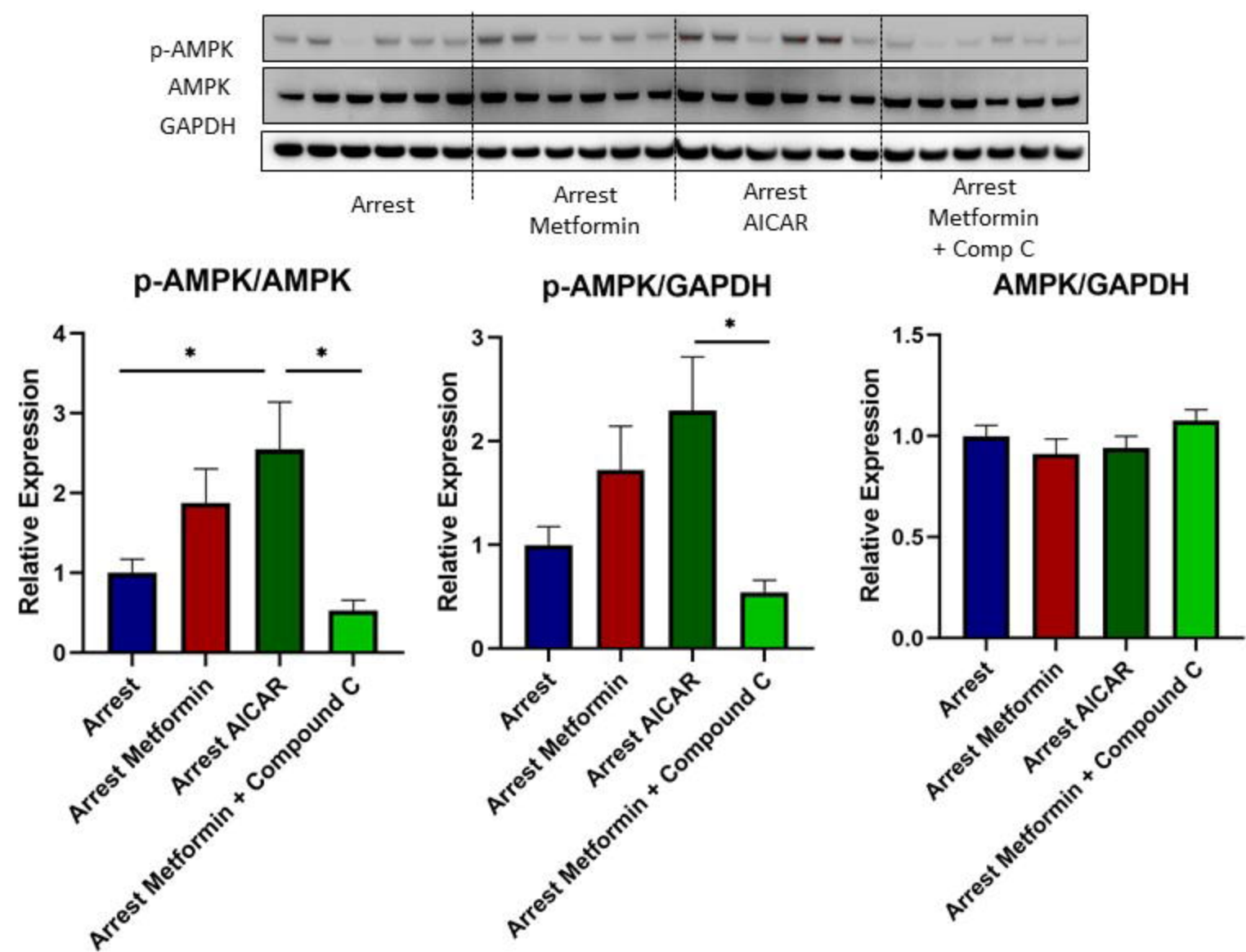


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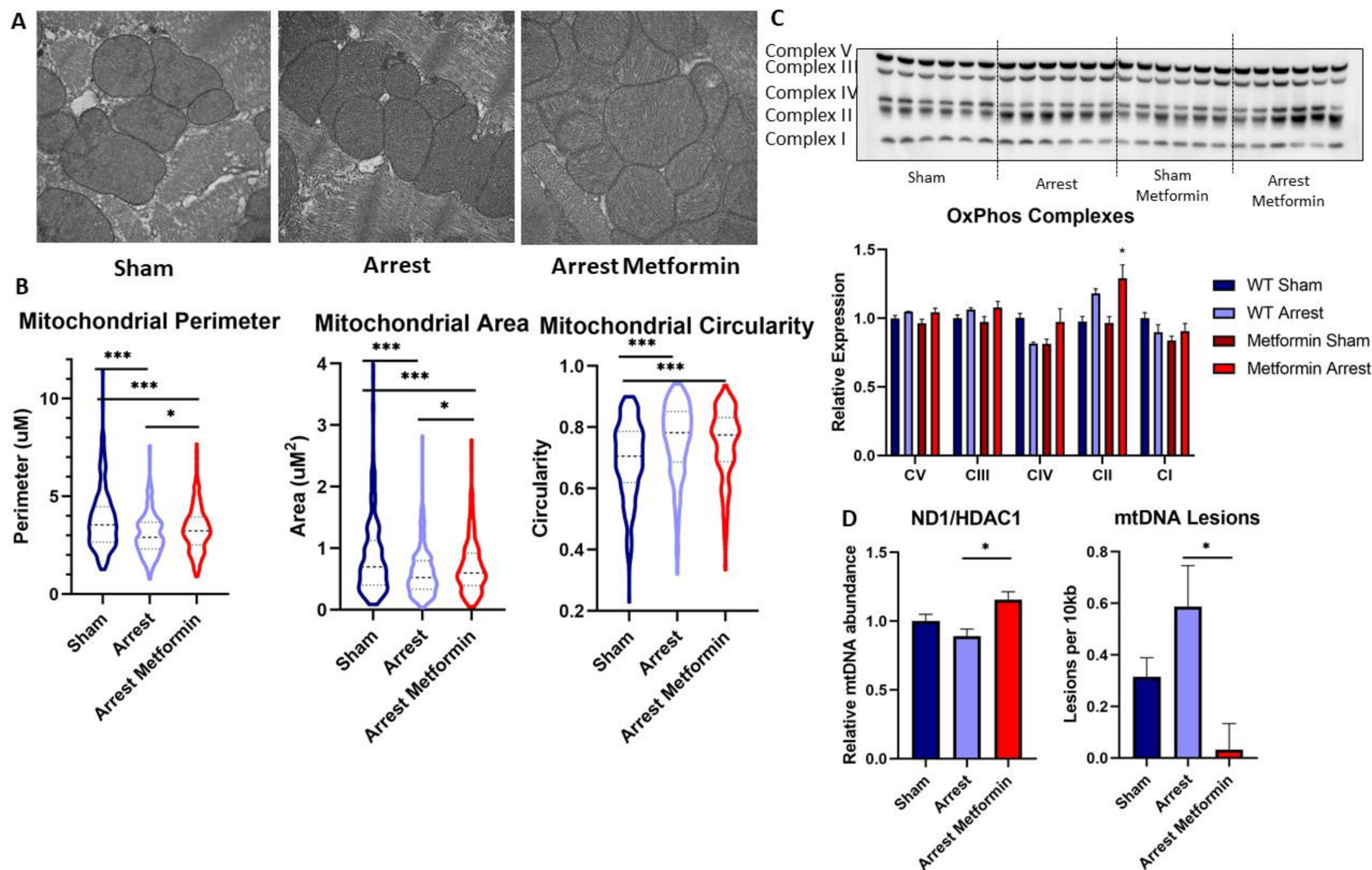


Figure 4. Metformin affects mitochondrial characteristics after SCA. A) Representative electron microscope images of intrafibrillar mitochondria in untreated sham, untreated arrest, and metformin pretreated arrest mice one day after surgery (40,000x magnification). B) Mitochondria are smaller and more circular in untreated arrest and metformin treated arrest mice when compared to sham, and metformin treated arrest mice have larger mitochondria than untreated arrest mice (n=50 per group). C) Electron transport chain expression is largely unchanged between treatment groups, with the exception of complex II (CII) expression in metformin-pretreated arrest mice being significantly higher than untreated sham mice one day after arrest (n=6/group, normalized to ponceau stain). D) mtDNA copy number is increased and has less damage in metformin treated arrest mice (n=7) when compared to untreated arrest mice (n=5). Untreated arrest mice had no changes to mtDNA copy number or damage when compared to sham. Data are expressed as mean \pm SEM. P-values: * < 0.05, ** < 0.01, *** < 0.001 by one-way ANOVA with Tukey post-hoc analysis. ND1, NADH dehydrogenase 1; HDAC1, Histone deacetylase 1; mtDNA, mitochondrial DNA; WT, wild-type.

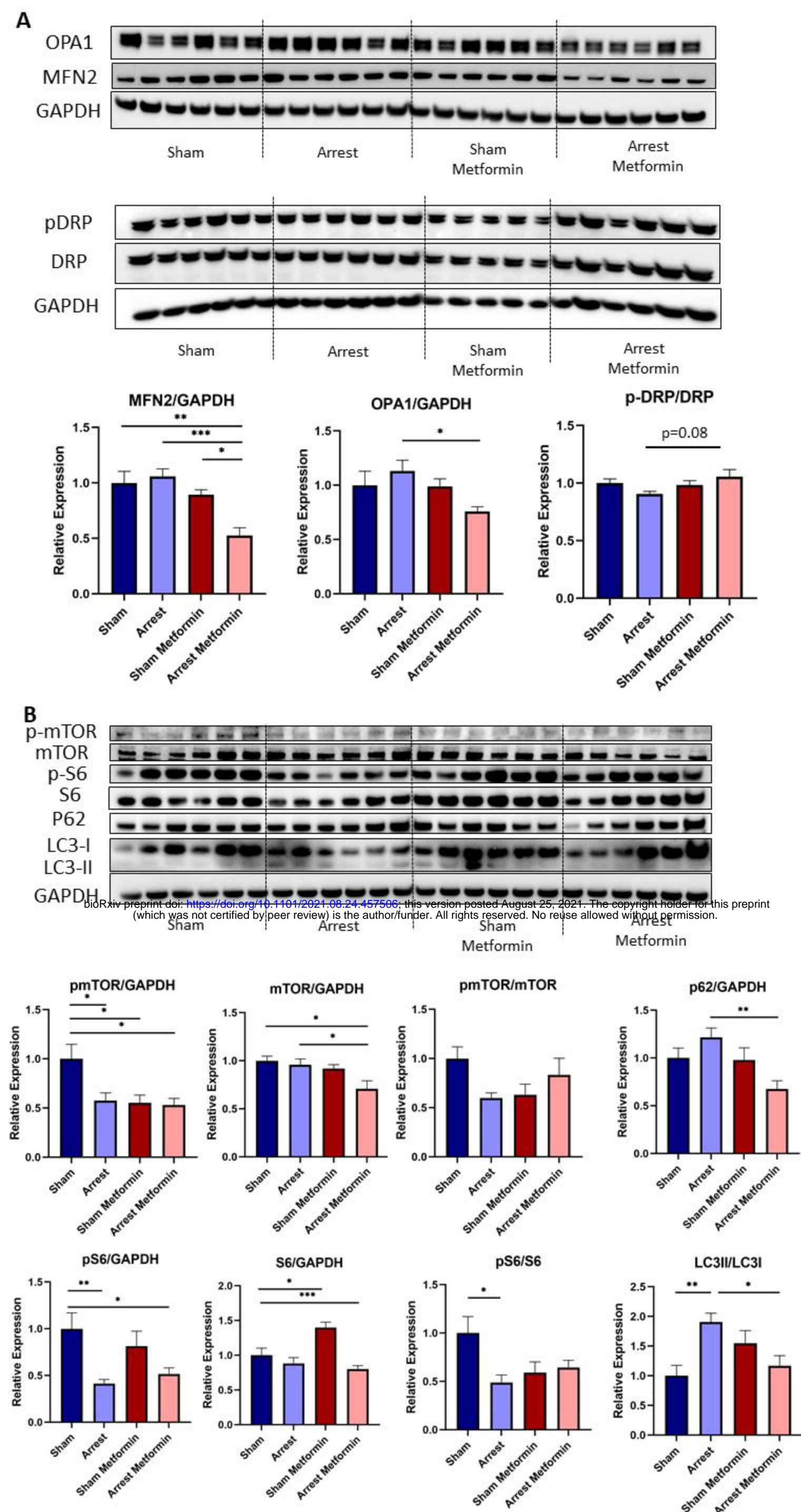


Figure 5. Metformin pretreatment affects autophagy and mitochondrial dynamics after sudden cardiac arrest (SCA). A) Representative western blot images of markers of mitochondrial fission and fusion. Mitofusin 2 (MFN2) is significantly depressed in metformin treated arrest mice (n=6) compared to untreated sham (n=6), untreated arrest (n=6), and sham metformin (n=5) groups. OPA1 is significantly depressed in metformin treated arrest compared to untreated arrest mice. There is no significant change to p-DRP expression between groups. B) Representative western blot images of autophagy related proteins downstream of AMPK. p-mTOR expression is reduced in untreated arrest, metformin treated sham, and metformin treated arrest mice when compared to untreated sham, and metformin treated mice have lower total mTOR than untreated sham and arrest groups. p-S6, a marker of mTOR activity, is also reduced in untreated arrest and metformin treated arrest mice when compared to sham. S6 expression is increased in sham metformin mice, but depressed in metformin arrest mice compared to sham, while the p-S6/S6 ratio is only significantly changed in the untreated arrest mice when compared to sham. P62, a marker of autophagosome formation, is lower in metformin treated arrest mice compared to untreated arrest mice. Untreated arrest mice have significantly elevated LC3II/LC3I, a marker of autophagy, compared to sham mice and metformin treated arrest mice (n=6/group). Data are expressed as mean \pm SEM. P-values: * < 0.05, ** < 0.01, *** < 0.001 by one-way ANOVA with Tukey post-hoc analysis for all groups, except for the p-S6 analyses, which used Dunnett's post-hoc analysis. DRP, dynamin-related protein1; LC3, microtubule-associated protein light chain; MFN2, mitofusin 2; mTOR, mechanistic target of rapamycin; OPA1, dynamin-like 120 kDa protein, mitochondrial.

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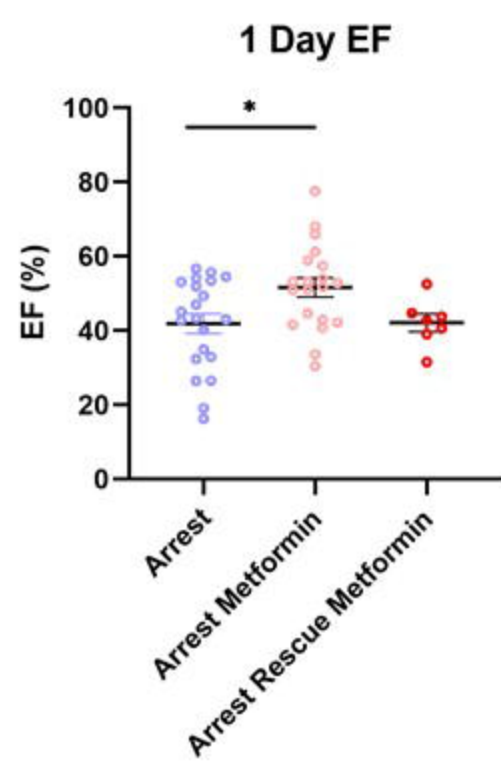


Figure 6. Metformin does not work as a rescue therapy following arrest. A) When given concomitantly with epinephrine, intravenous metformin (n=6) did not demonstrate any change to ejection fraction (EF) when compared to untreated arrest mice (n=15; data presented in Figure 2) or metformin treated arrest mice (n=20, data presented in Figure 2). P-values: * < 0.05 by one-way ANOVA with Tukey post-hoc analysis.

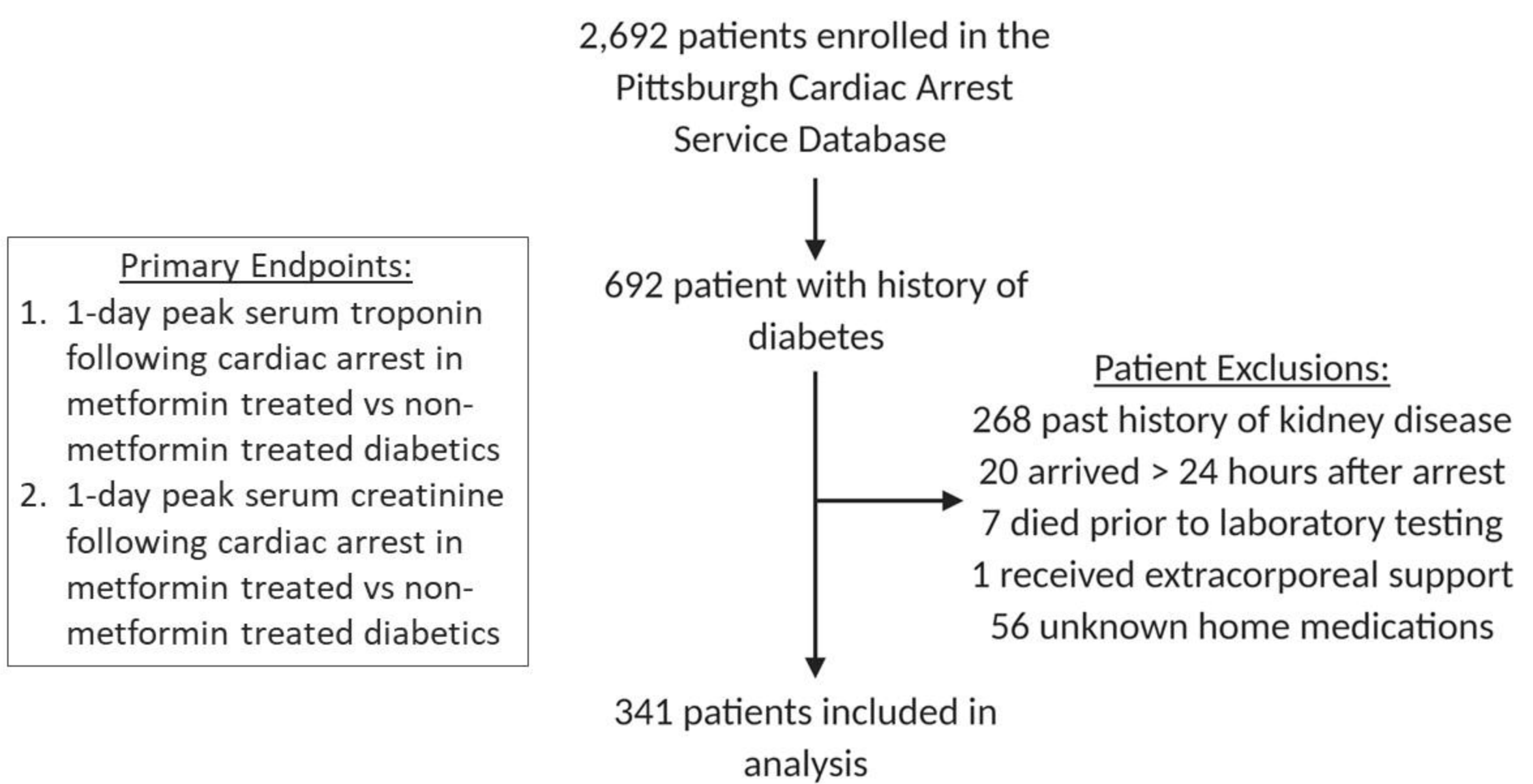


Figure 7. Summary of clinical inclusion and exclusion data for retrospective analysis of the Pittsburgh Post-Cardiac Arrest Service patient database. Primary outcomes evaluated included peak serum troponin and peak serum creatinine in diabetic patients with (n=122) and without (n=174) a history of metformin therapy.

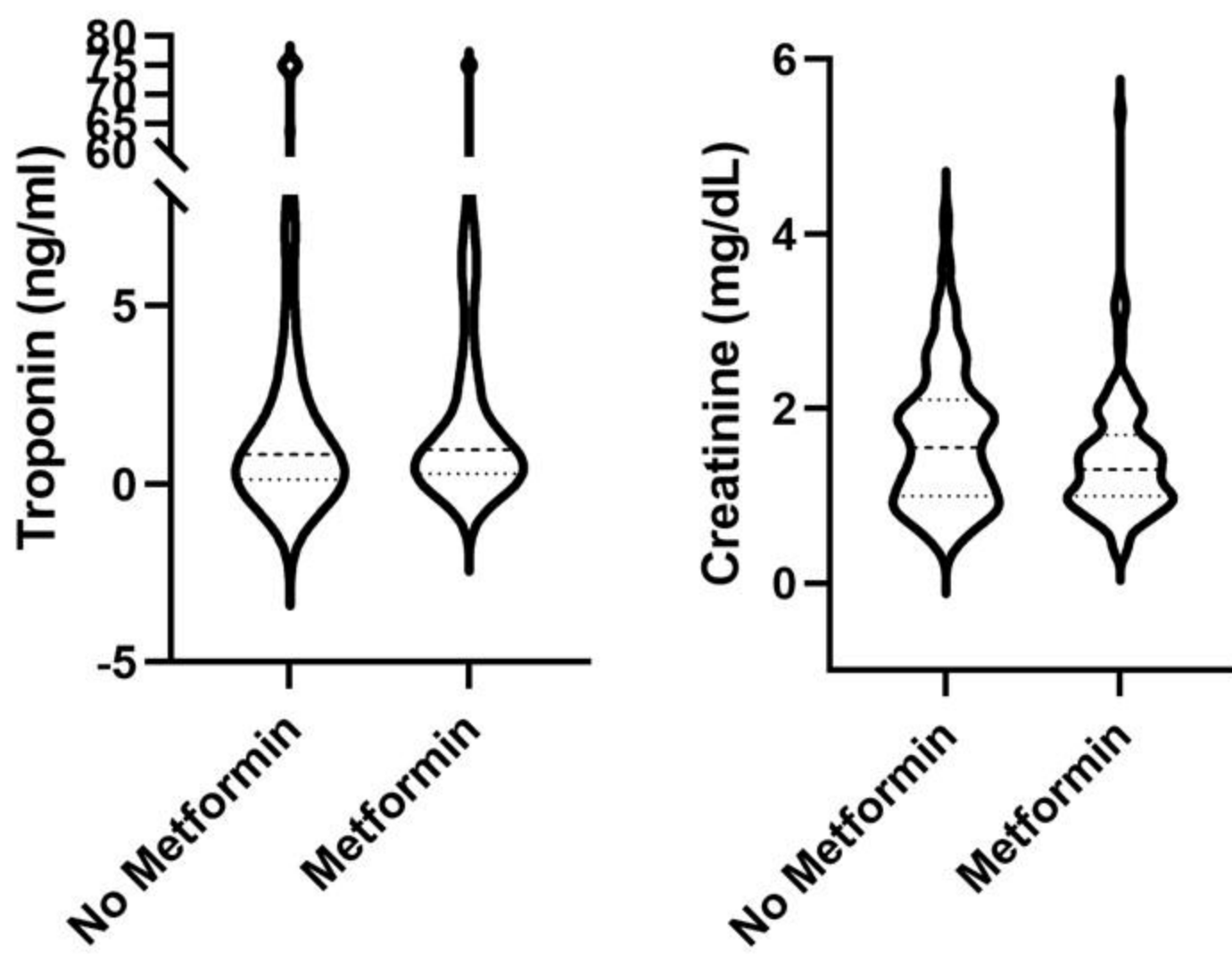


Figure 8. Distributions of peak 24-hour serum troponin and peak 24-hour serum creatinine for diabetic patients with and without metformin therapy prior to arrest. Dashed lines represent median values and dotted lines represent upper and lower quartiles.

Table 1. Surgical data for arrest mice. There are no significant changes to body weight or body temperature at time of extubation between treatment groups. There is no change in time to return of spontaneous circulation (ROSC) or time to extubation between untreated arrest and treated arrest groups. There is no change to random blood glucose 24-hours after arrest between sham, untreated arrest, and metformin pretreated arrest mice. Data are presented as mean \pm SEM unless otherwise noted. Analysis by one-way ANOVA with Tukey post-hoc analysis.

	Sham (n=20)	Arrest (n=26)	Sham Metformin (n=8)	Arrest Metformin (n=21)	Arrest Low Dose Metformin (n=9)	Arrest AICAR (n=9)	Arrest Metformin+ Compound C (n=8)	Arrest Rescue Metformin (n=7)
Age (d)	57.55 \pm 0.68	57.60 \pm 0.59	58.75 \pm 0.40	58.80 \pm 0.48	60.33 \pm 0.53	58.44 \pm 0.50	58.75 \pm 0.37	60.71 \pm 0.61
# Female / Total Mice (percentage)	11/20 (55%)	12/26 (46%)	5/8 (63%)	12/21 (57%)	4/9 (44%)	4/9 (44%)	4/8 (50%)	3/7 (43%)
Body Wt (g)	23.19 \pm 0.83	22.97 \pm 0.62	21.05 \pm 0.88	21.77 \pm 0.78	20.86 \pm 0.84	22.59 \pm 1.1	22.47 \pm 1.23	22.46 \pm 1.34
Temp at Extubation ($^{\circ}$ C)	35.86 \pm 0.15	35.53 \pm 0.20	35.96 \pm 0.18	35.95 \pm 0.12	35.53 \pm 0.21	35.79 \pm 0.19	36.21 \pm 0.11	35.86 \pm 0.23
ROSC (min)	-	1.31 \pm 0.09	-	1.43 \pm 0.15	1.11 \pm 0.11	1.44 \pm 0.24	1.13 \pm 0.12	1.29 \pm 0.18
Time to Extubation (min)	-	22.31 \pm 0.66	-	23.64 \pm 0.32	22.72 \pm 0.49	22.06 \pm 0.90	24.06 \pm 0.52	22.71 \pm 0.84
Time to Kidney Reperfusion (min)	-	21.3 \pm 0.90	-	21.79 \pm 0.57	22.24 \pm 0.49	20.43 \pm 1.01	24.70 \pm 0.64	22.50 \pm 0.97
Random glucose 24 h after arrest (mmol/L)	184.3 \pm 25.0	205.8 \pm 10.0	-	218.8 \pm 28.0	-	-	-	-

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Table 2. Baseline demographics and clinical characteristics. Data are presented as mean \pm standard deviation, median [interquartile range], or sample number (corresponding percentage).

Characteristic	Overall cohort (n = 341)	Metformin (n = 140)	No metformin (n = 201)
Age, years	65 \pm 13	65 \pm 12	64 \pm 14
Female sex	148 (43)	58 (41)	90 (45)
Arrest out-of-hospital	256 (75)	108 (77)	148 (74)
Shockable rhythm	109 (32)	39 (28)	70 (35)
Witnessed collapse	160 (47)	70 (50)	90 (45)
Layperson CPR	156 (46)	72 (51)	84 (42)
Epinephrine doses	2 [1 – 4]	3 [1 – 4]	2 [1 – 4]
Arrest duration, min	16 [8 – 27]	16 [8 – 32]	16 [8 – 23]
Cardiac etiology	96 (28)	36 (26)	60 (30)
Charlson Comorbidity index	2 [2 – 3]	2 [1 – 3]	3 [2 – 3]
Insulin	152 (45)	34 (24)	118 (59)
Other oral diabetic medication	92 (27)	47 (34)	45 (22)
Peak 24h troponin	0.88 [0.19 – 5.7]	0.97 [0.29 – 4.71]	0.84 [0.14 – 7.0]
Peak 24h creatinine	1.4 [1.0 – 2.0]	1.3 [1.0 – 1.7]	1.6 [1.0 – 2.1]

Table 3. Association between metformin use and peak serum creatine and troponin by log link model. Model was adjusted for age, sex, arrest location (in- vs out-of-hospital), witnessed collapse, layperson cardiopulmonary resuscitation, presenting rhythm, arrest duration, number of epinephrine administered, cardiac etiology of arrest, Charlson Comorbidity index, insulin, and other oral diabetic medications. Both peak serum creatinine level and peak serum troponin level 24 hours post-arrest are significantly associated with history of metformin use.

Endpoint	Coefficient (95% CI)	P value
Creatinine	-0.19 (-0.30 – -0.08)	0.001
Troponin	-1.29 (-2.11 – -0.46)	0.002