

1 **Inhibiting succinate release worsens cardiac reperfusion injury by enhancing mitochondrial**
2 **reactive oxygen species generation.**

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21 **ABSTRACT**

22 The metabolite succinate accumulates during cardiac ischemia. Within 5 min. of
23 reperfusion, succinate returns to baseline levels via both its release from cells and oxidation by
24 mitochondrial complex II (Cx-II). The latter drives reactive oxygen species (ROS) generation and
25 subsequent opening of the mitochondrial permeability transition (PT) pore, leading to cell death.
26 Targeting succinate dynamics (accumulation/oxidation/release) may be therapeutically
27 beneficial in cardiac ischemia-reperfusion (IR) injury. It has been proposed that blocking
28 monocarboxylate transporter 1 (MCT-1) may be beneficial in IR, by preventing succinate release
29 and subsequent engagement of downstream inflammatory signaling pathways. In contrast,
30 herein we hypothesized that blocking MCT-1 would retain succinate in cells, exacerbating ROS
31 generation and IR injury. Using the mitochondrial ROS probe mitoSOX, and a custom-built murine
32 heart perfusion rig built into a spectrofluorometer, we measured ROS generation *in-situ* during
33 the first moments of reperfusion, and found that acute MCT-1 inhibition enhanced mitochondrial
34 ROS generation at reperfusion, and worsened IR injury (recovery of function and infarct size).
35 Both these effects were abrogated by tandem inhibition of Cx-II, suggesting that succinate
36 retention worsens IR due to driving more mitochondrial ROS generation. Furthermore, using the
37 PT pore inhibitor cyclosporin A, along with monitoring of PT pore opening via the mitochondrial
38 membrane potential indicator TMRE, we herein provide evidence that ROS generation during
39 early reperfusion is upstream of the PT pore, not downstream as proposed by others. In addition,
40 pore opening was exacerbated by MCT-1 inhibition. Together, these findings highlight the
41 importance of succinate dynamics and mitochondrial ROS generation, as key determinants of PT
42 pore opening and IR injury outcomes.

43

44

45 1. INTRODUCTION

46 The metabolite succinate has a central role in tissue ischemia. Several mechanisms exist
47 for ischemic succinate accumulation ^{1, 2}, and this process is conserved across diverse tissues,
48 species, and physiologic contexts ^{3, 4}. However, rapid oxidation of accumulated succinate at the
49 onset of tissue reperfusion is a key event in ischemia-reperfusion (IR) injury, the underlying
50 pathology of myocardial infarction (heart attack) ^{1, 2}. This has led to intense interest in succinate
51 as a potential therapeutic target ⁵⁻⁹.

52 Approximately $\frac{1}{3}$ of succinate accumulated during ischemia is rapidly oxidized by complex
53 II (Cx-II) of the mitochondrial respiratory chain ^{2, 10}. This leads to the generation of reactive oxygen
54 species (ROS) via mechanisms that are thought to involve either reverse electron transport (RET)
55 at complex-I (Cx-I) ¹¹⁻¹³ or forward electron transport at complex-III (Cx-III) ¹⁴⁻¹⁸. Mitochondrial
56 ROS generation potentiates opening of the mitochondrial permeability transition (PT) pore ¹⁹⁻²²,
57 which triggers necrotic cell death.

58 The remaining $\frac{2}{3}$ of succinate accumulated during ischemia is released upon reperfusion,
59 in a pH-dependent manner via monocarboxylate transporter 1 (MCT-1) ^{10, 23}. The physiologic
60 function of this released succinate is unclear, with suggestions that it may serve as a metabolic
61 signal of hypoxia ²⁴. Succinate is a ligand for the widely expressed succinate receptor (SUCNR1,
62 formerly GPR91), which elicits a range of physiologic responses ²⁵. Most relevant to IR injury,
63 SUCNR1 signaling promotes inflammation via macrophage activation, which may contribute to
64 the pathology of IR injury ^{10, 26-30}. As such, it has been postulated that inhibiting MCT-1 would be
65 beneficial in IR injury, via blunting of extracellular succinate \rightarrow SUCNR1 signaling ^{10, 31}.

66 In contrast, given the key role of intracellular succinate for ROS generation during
67 reperfusion, we hypothesized herein that acute blockade of succinate release via MCT-1 would
68 worsen IR injury, and that simultaneous blockade of Cx-II would abrogate this effect. This
69 hypothesis was tested by measuring mitochondrial ROS generation *in-situ* in perfused mouse
70 hearts using a custom-built perfusion rig within the chamber of a benchtop spectrofluorometer.
71 In addition, since it has been proposed that PT pore opening itself may lie upstream of the burst

72 of ROS seen upon reperfusion ^{23, 32}, we used this apparatus to interrogate the temporal
73 relationship between ROS generation and PT pore opening during early reperfusion.

74

75 **2. MATERIALS AND METHODS**

76 **2.1 Animals and Reagents**

77 Animal and experimental procedures complied with the National Institutes of Health
78 *Guide for Care and Use of Laboratory Animals* (8th edition, 2011) and were approved by the
79 University of Rochester Committee on Animal Resources (protocol #2007-087). Male and female
80 C57BL/6J adult mice (8-20 weeks old) were housed in a pathogen-free vivarium with 12 hr. light-
81 dark cycles and food and water *ad libitum*. Mice were administered terminal anesthesia via intra-
82 peritoneal 2,2,2-tribromoethanol (Avertin) ~250 mg/kg. Avertin was prepared in amber glass
83 vials and stored at 4°C for no more than 1 month. This agent was chosen for anesthesia since it
84 does not impact cardioprotection as reported for volatile anesthetics or opioids ³³⁻³⁵, and does
85 not have mitochondrial depressant effects as reported for barbiturates ³⁶. Euthanasia occurred
86 via cardiac extirpation (see below).

87 MitoSOX red was from Thermo (NJ, USA) and was stored aliquoted under argon prior to
88 use. Fully oxidized mitoSOX was prepared by reaction with Fremy's salt, as described elsewhere
89 ³⁷. AR-C155858 was from MedChemExpress (Monmouth Junction, NJ, USA). Unless otherwise
90 stated, all other reagents were from Sigma (St. Louis MO, USA).

91

92 **2.2 Perfused Mouse Hearts**

93 Following establishment of anesthetic plane (toe-pinch response), beating mouse hearts
94 were rapidly cannulated, excised and retrograde perfused at a constant flow (4 ml/min.) with
95 Krebs-Henseleit buffer (KHB) consisting of (in mM): NaCl (118), KCl (4.7), MgSO₄ (1.2), NaHCO₃
96 (25), KH₂PO₄ (1.2), CaCl₂ (2.5), glucose (5), pyruvate (0.2), lactate (1.2), and palmitate (0.1,
97 conjugated 6:1 to bovine serum albumin). KHB was gassed with 95% O₂ and 5% CO₂ at 37°C. A
98 water-filled balloon connected to a pressure transducer was inserted into the left ventricle and
99 expanded to provide a diastolic pressure of 6-8 mmHg. Cardiac function was recorded digitally at
100 1 kHz (Dataq, Akron OH) for the duration of the protocol. Following equilibration (10-20 min.)

101 ischemia-reperfusion (IR) injury comprised 25 min. global no-flow ischemia plus 60 min.
102 reperfusion. Hearts were then sliced and stained with triphenyltetrazolium chloride (TTC) for
103 infarct quantitation by planimetry (red = live tissue, white = infarct). Infarction analysis was
104 blinded to experimentalists.

105 The following conditions (Figure 1F) were examined: **(i) Control:** DMSO vehicle infusion 5
106 min. prior to ischemia and 5 min. into reperfusion. **(ii) S1QEL:** 1.6 μ M S1QEL1.1, delivered 5 min.
107 prior to ischemia and 5 min. into reperfusion. **(iii) DMM5:** 5 mM dimethyl malonate at the onset
108 of reperfusion and 5 min. into reperfusion. **(iv) AR:** 10 μ M AR-C155858 infusion, 5 min. prior to
109 ischemia and 5 min. into reperfusion. **(v) AR + DMM5:** 10 μ M AR-C155858 infusion, 5 min. prior
110 to ischemia and 5 min. into reperfusion plus 5 mM dimethyl malonate at the onset of reperfusion
111 and 5 min. into reperfusion. **(vi) AR + DMM10:** 10 μ M AR-C155858 infusion, 5 min. prior to
112 ischemia and 5 min. into reperfusion plus 10 mM dimethyl malonate at the onset of reperfusion
113 and 5 min. into reperfusion. **(vii) AR + AA5:** 10 μ M AR-C155858 infusion, 5 min. prior to ischemia
114 and 5 min. into reperfusion plus 100 nM atpenin A5 at the onset of reperfusion and 5 min. into
115 reperfusion. **(viii) CsA:** 0.8 μ M cyclosporin A infusion³⁸, 5 min. prior to ischemia and 5 min. into
116 reperfusion. Simultaneously with these conditions, hearts were delivered either 1.5 μ M mitoSOX
117 for 5 min. or 500 nM tetramethylrhodamine ethyl ester (TMRE) for 20 min., prior to ischemia.

118

119 **2.3 Perfusion Apparatus Within Spectrofluorometer**

120 A cardiac perfusion apparatus was custom-built with an umbilicum, to position the heart
121 within the light-proof enclosure of a Varian/Cary Eclipse benchtop spectrofluorometer (Agilent,
122 Santa Clara CA, USA), as shown in Figure 1A-1C. Hearts were placed against the wall of a water-
123 jacketed cuvette maintained at 37°C in which the excitation light source strikes the left ventricle
124 of the heart at a 45° angle relative to the photomultiplier tube (PMT) window. The cuvet holder
125 was 3D printed (.stl file in supplementary materials). Data was collected using Cary WinUV
126 Kinetics software, which permitted real-time fluorescence monitoring (1 s. reads / 15 s. cycle,
127 PMT voltage 600, Ex/Em slit width = 5 nm) for the duration of the perfusion protocol. At the heart
128 rates observed (438 ± 54 bpm, mean \pm SD, N=54) a 1 s. fluorescent read averages across ~7 heart
129 beats, so gating for motion artifacts was unnecessary. Initial validation was performed by

130 monitoring endogenous NAD(P)H fluorescence (λ_{EX} 340 nm, λ_{EM} 460 nm) and flavoprotein
131 fluorescence (λ_{EX} 460 nm, λ_{EM} 520 nm), as shown in Figure 1D-1E.

132

133 **2.4 Measuring *in situ* Mitochondrial ROS**

134 Mitochondrial ROS generation was measured using the probe mitoSOX^{39, 40}
135 (tetraphenylphosphonium-conjugated dihydroethidium, TPP⁺-DHE) at a concentration of 1.5
136 μ M, since the TPP⁺ moiety is known to uncouple mitochondria at concentrations $\geq 2.5 \mu$ M⁴¹⁻⁴³.
137 Hearts subject to conditions (i-vii) described above were equilibrated for 10 min., loaded with 1.5
138 μ M mitoSOX for 5 min. before immediately being subjected to 25 min. ischemia. Fluorescence
139 was monitored at λ_{EX} 510 nm, λ_{EM} 580 nm.

140 Since the λ_{EX} and λ_{EM} of mitoSOX overlap with absorbance spectra of endogenous
141 chromophores in cardiomyocytes (e.g., myoglobin and cytochromes), changes in chromophore
142 absorbance during the course of IR could impact the fluorescent signal. Furthermore, the
143 distribution of the mitoSOX probe between cytosolic and mitochondrial matrix compartments is
144 determined by the mitochondrial membrane potential ($\Delta\Psi_m$), which would also be expected to
145 change during IR. To correct for these potential confounding effects, a series of hearts were loaded
146 with fully-oxidized mitoSOX³⁷, which fluoresces in a manner that is independent of ROS
147 generation, but is still subject to the effects of chromophore absorbance and probe distribution.
148 Fluorescent data from these hearts was used to correct data obtained with naïve mitoSOX
149 (conditions i-vii), yielding a net signal that originated only from *in-situ* probe oxidation, without
150 contribution from other factors. The data process for this correction is illustrated in Supplemental
151 Figure 1, which also shows a minimal change in signal during IR when no mitoSOX was present.
152 In addition, absorbance spectra of compounds tested in this study (AR, DMM, S1QEL, etc.) were
153 also measured, to ensure these chemicals did not absorb significant amounts of light at the λ_{EX}
154 and λ_{EM} of mitoSOX (Supplemental Figure 2).

155

156 **2.5 Measuring *in situ* Mitochondrial Membrane Potential ($\Delta\Psi_m$)**

157 In order to assess the timing of mitochondrial PT pore opening, mitochondrial membrane
158 potential was measured using TMRE at 500 nM delivered for 20 min. prior to ischemia. The PT

159 pore inhibitor CsA (0.8 μ M) was optionally delivered for 5 min. prior to ischemia. To determine
160 changes upon reperfusion, TMRE data were normalized to the fluorescent signal during the last
161 5 min. of ischemia. Two conditions were examined: control IR, and IR with MCT-1 inhibition (same
162 as condition (vi) in section 2.2 above). Attempts to determine the effect of CsA on PT pore
163 opening in the presence of AR-C155858 were confounded by interactions between these
164 molecules and TMRE, resulting in precipitation of components in Krebs-Henseleit buffer.

165

166 **2.6 Effluent Analysis by High-Performance Liquid Chromatography**

167 Effluents from control and AR treated hearts were collected in 1 min. intervals for the first
168 3 min. of reperfusion, and immediately treated with 10 % perchloric acid. Following addition of
169 100 nmols butyrate as an internal standard, samples were frozen in liquid N₂ and stored at -80°C
170 until analysis. Samples were centrifuged at 20,000 x g to remove insoluble materials. Metabolites
171 were resolved on HPLC (Shimadzu Prominence 20 system) using two 300 x 7.8 mm Aminex HPX-
172 87H columns (BioRad, Carlsbad CA, USA) in series with 10 mM H₂SO₄ mobile phase (flow rate: 0.7
173 ml/min) and 100 μ l sample injected on column. Succinate and lactate were detected using a
174 photodiode array measuring absorbance at 210 nm as previously described ². A standard curve
175 was constructed for calibration. Lactate data were corrected for 1.2 mM lactate contained in the
176 Krebs-Henseleit buffer.

177

178 **2.7 Quantitation and Statistical Analysis**

179 Comparisons between groups were made using ANOVA, followed by unpaired Student's
180 t-tests. Data are shown as means \pm SEM. Numbers of biological replicates (N) are noted in the
181 figures. Significance was set at $\alpha = 0.05$.

182

183 **3. RESULTS**

184 **3.1 Cardiac Fluorescence**

185 Although similar spectrofluorometric cardiac perfusion apparatus has previously been
186 constructed ^{32, 44-48}, prior efforts have used larger animal hearts (rats, guinea pigs) or have not

187 simultaneously measured cardiac function. To the best of our knowledge, this is the first study of
188 mouse hearts with simultaneous fluorescence and functional assessment.

189 The spectrofluorimetric perfusion system was first validated by monitoring NAD(P)H and
190 flavoprotein autofluorescence during ischemia and reperfusion (Fig. 1D and 1E). As expected,
191 NAD(P)H (λ_{EX} 340 nm, λ_{EM} 460 nm) autofluorescence immediately rose upon ischemia, in
192 agreement with previous reports^{32, 44, 49, 50}. Likewise, flavoprotein fluorescence (λ_{EX} 460 nm, λ_{EM}
193 520 nm) decreased upon ischemia^{32, 50, 51}. Both parameters returned to baseline levels
194 immediately upon reperfusion.

195 NAD(P)H autofluorescence in the heart is thought to mainly represent mitochondrial
196 NADH, the substrate for Cx-I which accumulates during ischemia due to a highly reduced
197 respiratory chain⁵⁰. The mitochondrial ATP synthase is thought to operate in reverse during
198 ischemia, to maintain mitochondrial membrane potential ($\Delta\Psi_m$)^{52, 53}, and this also contributes
199 to feedback inhibition of Cx-I resulting in high [NADH]¹¹. Notably, a decrease in NAD(P)H signal
200 was observed ~12 min. into ischemia, concurrent with the onset of ischemic hyper-contracture
201 (not shown). The latter is thought to indicate the onset of an energetic crisis (no ATP to relax
202 contractile machinery)⁵⁴.

203

204 **3.2 MitoSOX Fluorescence**

205 Having validated the cardiac fluorescence system, we next sought to use it for
206 measurement of ROS generation during IR. Figure 2A shows cardiac functional measurements
207 throughout IR (heart rate x pressure product, RPP), while Figure 2B shows the corrected mitoSOX
208 fluorescent readout over the same period.

209 To account for changes in the absorbance of endogenous chromophores during IR, or
210 probe distribution, additional hearts were perfused with oxidized mitoSOX or with no probe at
211 all. No significant change in the fluorescent signal was observed in hearts without mitoSOX
212 (Supplemental Figure 1A). However, delivery of oxidized mitoSOX resulted in a rapid increase in
213 the fluorescent signal during dye loading (Supplemental Figure 1B). At the onset of ischemia, an
214 additional signal increase was seen. This is expected, since it has been shown that cardiac tissue
215 absorbance at 510 nm and 580 nm decreases during ischemia⁵⁵. Upon reperfusion, after a small

216 increase, the oxidized mitoSOX signal steadily declined for the remainder of the experiment. This
217 is possibly due to loss of the dye from mitochondria as a result of changes in ($\Delta\Psi_m$) as
218 mitochondrial integrity becomes compromised.

219 Raw mitoSOX fluorescence traces (Supplemental Figure 1C) also increased slightly during
220 loading, but not to the same extent as the fully oxidized probe. A sharp signal increase at the
221 onset of ischemia may represent a burst of ROS generation as the terminal respiratory chain
222 becomes inhibited. Upon reperfusion, a sustained signal increase was observed for ~10 min.,
223 followed by a decline. Using the oxidized mitoSOX data to correct the raw mitoSOX data, Figure
224 2B (Supplemental Figure 1D) shows the signal resulting from oxidation of the probe during IR.
225 Upon reperfusion, a sustained increase in the redox-dependent mitoSOX signal was seen, and
226 this is consistent with the concept that a burst of mitochondrial ROS generation occurs during
227 the first minutes of reperfusion ^{1, 56-58}.

228 It has been posited that reverse electron transport (RET) at Cx-I is the primary source of
229 ROS during reperfusion, and recently a novel series of inhibitors that target ROS generation at
230 the ubiquinone (Q) binding site of Cx-I (termed S1QELs) were shown to elicit cardioprotection
231 against IR injury ⁵⁹. Succinate levels return to baseline within the first 5 min. of reperfusion ², and
232 accordingly, examining mitoSOX fluorescence during this time period revealed that the
233 immediate signal increase at the onset of reperfusion in control hearts was suppressed in hearts
234 treated with a S1QEL (Figure 3A-3C). These data suggest that the ROS signal detected by mitoSOX
235 during the first minutes of reperfusion originates from Cx-I RET. However, by 2 minutes the rate
236 of signal increase in S1QEL hearts had returned to that seen in control hearts, potentially
237 indicating a role for other sources of ROS ¹⁷. While no effect of S1QEL on flavoprotein
238 fluorescence was observed (Figure 3E), S1QEL did cause a slight detriment in the elevation of
239 NAD(P)H fluorescence at the start of ischemia (37±4% with S1QEL vs. 53±5% in controls,
240 p=0.042). However, it also blunted the NAD(P)H response to ischemic hyper-contracture (Figure
241 3D). The combination of these effects was such that the drop in NAD(P)H signal at the onset of
242 reperfusion was not significantly different between S1QEL vs. control (30±4% vs. 36±4%
243 respectively), suggesting that consumption of NADH by Cx-I during early reperfusion was not
244 impacted by the compound.

245

246 **3.3 Preventing Succinate Efflux Exacerbates Mitochondrial ROS at Reperfusion**

247 Upon reperfusion of ischemic heart, $\frac{1}{3}$ of accumulated succinate is oxidized by Cx-II,
248 driving ROS generation ^{2, 10}. Consistent with this, the Cx-II competitive inhibitor malonate is
249 reported to be cardioprotective when delivered at reperfusion ^{6, 7, 60}. The remaining $\frac{2}{3}$ of
250 succinate accumulated during ischemia is released from tissue upon reperfusion, and the
251 pathway for this release was recently elucidated as monocarboxylate transporter 1 (MCT-1) ^{2, 10}.
252 To test the impact of MCT-1 inhibition on ROS generation at reperfusion, hearts were infused
253 peri-ischemically with the MCT-1 inhibitor AR-C155858 (AR) ⁶¹, which resulted in a significant
254 decrease in succinate release into the post-cardiac effluent during the first 3 minutes of
255 reperfusion (Supplemental Figure 3). As expected, lactate efflux was also significantly diminished,
256 confirming MCT-1 inhibition.

257 Figures 4C & 4F show that AR resulted in a significantly greater rate of ROS generation
258 during the first minute of reperfusion, compared to control. To investigate the requirement for
259 Cx-II in the additional AR-induced mitoSOX signal, the competitive Cx-II inhibitor dimethyl
260 malonate (DMM, 5mM) was used. Surprisingly, DMM alone at this concentration did not
261 significantly impact the mitoSOX signal, and it also did not significantly blunt the additional signal
262 induced by AR (Figure 4B, 4D & 4F). We hypothesized that because malonate is a competitive Cx-
263 II inhibitor, it may not be able to out-compete the additional succinate present in cells caused by
264 MCT-1 inhibition. Supporting this hypothesis, tandem administration of a higher dose of DMM
265 (10 mM) was capable of blocking the elevated mitoSOX signal elicited by AR, returning it to
266 control levels (Figure 4E & 4F). Furthermore, the potent Cx-II inhibitor atpenin A5 (AA5, 100 nM)
267 was also effective in blocking the additional mitoSOX signal induced by AR (Supplemental Figure
268 4A & 4B). Overall, these data suggest that MCT-1 inhibition enhances mitochondrial ROS
269 generation in the first minute of reperfusion, in a manner that can be blocked by inhibitors of Cx-
270 II.

271

272 **3.4 Blocking Succinate Release via MCT-1 Worsens IR Injury**

273 In agreement with the role of succinate-derived ROS as a driver of post-IR pathology,
274 DMM alone improved and AR worsened, IR injury (Figure 5A-C, cardiac functional recovery and
275 infarct size). In addition, tandem administration of low dose DMM (5 mM) failed to reverse the
276 impact of AR, whereas high dose DMM (10 mM) was protective. Furthermore, administration of
277 the potent Cx-II inhibitor AA5 also blocked the impact of AR on functional recovery and infarct
278 (Supplemental Figure 4C-E), and in-fact was more protective than AA5 alone ². These effects of
279 MCT-1 inhibition and Cx-II inhibition were more significant for myocardial infarction (Figure 5C)
280 than for functional recovery (Figure 5B), but all trended in the same direction similar to mitoSOX
281 data (Figure 4), i.e., interventions that increased mitoSOX signal at reperfusion worsened IR
282 injury, and those that decreased it improved IR injury. Overall, a correlation was observed
283 between the effect of interventions on mitoSOX at reperfusion and on infarct size and functional
284 recovery, as illustrated in Figure 6.

285

286 **3.5 Timing of Mitochondrial ROS at Reperfusion vs. PT Pore Opening.**

287 While it is largely accepted that ROS can trigger PT pore opening, it has also been
288 proposed that during reperfusion injury, the PT pore itself may drive ROS generation ^{23,32}. To test
289 this, we measured the effect of the PT pore inhibitor CsA on mitoSOX fluorescence during
290 reperfusion. 0.8 μ M CsA was chosen, as this was previously shown to elicit protection in mouse
291 hearts ³⁸. As shown in Figure 7B & 7C, CsA had no impact on mitoSOX fluorescence during
292 reperfusion. To confirm that PT pore opening did occur, hearts were loaded with the $\Delta\Psi_m$
293 indicator TMRE. Upon reperfusion, an immediate rise in TMRE fluorescence was observed.
294 Subsequently, in control hearts the TMRE signal declined from ~5 min. into reperfusion, whereas
295 in CsA treated hearts the signal was sustained (Figure 7D & 7E). We thus infer that PT pore
296 opening occurs no sooner than 5 min. into reperfusion. Despite a small blip in the mitoSOX signal
297 in control hearts at ~6.5 min., no substantial difference was seen between control & CsA hearts
298 at this time point, concurrent with divergence of the TMRE traces, thus suggesting no secondary
299 ROS burst due to PT pore opening.

300 Finally, as expected from its impact on mitoSOX fluorescence during early reperfusion
301 (Figure 4), inhibition of MCT-1 led to an accelerated loss of the TMRE signal during reperfusion,
302 indicating faster opening of the PT pore (Supplemental Figure 5).

303

304 4. DISCUSSION

305 Cardiac IR is a complex pathology, with numerous links between early and late events in
306 the development of myocardial infarction. Shortly after reperfusion, a burst of ROS generation
307 and mitochondrial Ca^{2+} overload both trigger opening of the mitochondrial PT pore, a key event
308 in necrotic cell death of cardiomyocytes. Following this, inflammatory cells (macrophages,
309 neutrophils) are recruited to the heart, where they mediate responses that lead to cardiac
310 remodeling, fibrosis, and eventual development of hypertrophy and heart failure⁶²⁻⁶⁴.

311 Accumulation of succinate in hypoxia/ischemia is highly conserved^{3,4}, but this succinate
312 drives ROS generation upon tissue reperfusion, and this has led to a consensus that intracellular
313 succinate plays a detrimental role in reperfusion injury^{1,2,16,65,66}. In contrast, succinate release
314 from cells via MCT-1^{2,10,23}, along with the recent identification of a succinate receptor
315 (GPR91/SUCNR1²⁵), have led to the notion that extracellular succinate may also play a role in IR
316^{6,10}. In this regard, findings from an *in-vivo* model of cardiac IR injury demonstrated that blocking
317 succinate release via MCT-1 was cardioprotective, likely due to inhibiting immune system
318 activation^{6,10}. However, herein our data reveal that acutely blocking MCT-1 in a perfused heart
319 system (where there are no inflammatory cells) leads to greater ROS generation and the
320 worsening of IR injury. Reconciling these findings, it is possible that MCT-1 inhibition is indeed
321 detrimental in the acute setting at the level of cardiomyocytes, but this is balanced *in-vivo* by a
322 longer-term effect of MCT-1 inhibition, possibly involving receptor-mediated succinate effects on
323 other cell types, including inflammatory cells. Together, these studies highlight that the
324 therapeutic targeting of MCT-1 may require careful timing and titration, to balance detrimental
325 vs. beneficial effects of blocking succinate release.

326 Inhibiting MCT-1 in the heart lowered succinate release into the effluent by ~30%.
327 Assuming this succinate was retained in cells and available for oxidation by Cx-II, it is therefore
328 not surprising that MCT-1 inhibition also led to enhanced ROS generation. Furthermore, while 5

329 mM of the competitive Cx-II inhibitor malonate was ineffective at blocking this additional ROS,
330 doubling its concentration to 10 mM effectively overrode the impact of AR. Unfortunately, due
331 to the physical nature of dimethylmalonate (liquid) and our drug infusion system, it was not
332 possible to test additional even higher doses of DMM. Instead, we tested the potent Cx-II
333 inhibitor atpenin A5 (AA5, $IC_{50} \sim 10 \text{ nM}^{67}$), and found that it was able to completely abrogate the
334 additional ROS induced by AR, and this resulted in cardioprotection to a level greater than
335 baseline control IR injury (Supplementary Figure 4). In-fact, the combination of AR and AA5 may
336 be considered optimal from a therapeutic perspective – keeping succinate inside cells to prevent
337 its signaling effects, and preventing its oxidation to generate ROS.

338 Although there is a general consensus that mitochondrial ROS generation during early
339 reperfusion is an upstream event that triggers PT pore opening, it has also been suggested that
340 pore opening itself is the main driver of ROS generation in early reperfusion ^{23, 32}. Our results
341 (Figure 7) suggest that the burst of ROS at reperfusion is independent of the PT pore, since
342 evidence for pore opening (i.e., a CsA-sensitive loss of $\Delta\Psi_m$) was not observed until at least 5 min.
343 into reperfusion, by which time a large amount of ROS generation had already occurred.

344 It should not go unmentioned that Ca^{2+} is also considered to be a major trigger for PT pore
345 opening ^{68, 69}, with both Ca^{2+} and ROS thought to potentiate each other's effects at promoting
346 pore formation ⁷⁰. Thus, future experiments should be directed at using fluorescent Ca^{2+} probes
347 in this or similar perfusion systems, to understand *in-situ* Ca^{2+} kinetics and how they relate to
348 ROS and pore opening kinetics in early reperfusion.

349 A number of caveats regarding the use of mitoSOX as a probe for mitochondrial ROS
350 should be addressed. Firstly, it is known the probe can be oxidized by reactants other than ROS,
351 although notably such one-electron oxidations result in non-fluorescent products that would be
352 undetectable in our measurement system ⁷¹. Furthermore, the mitoSOX signal increase in early
353 reperfusion was inhibited by S1QEL, a specific inhibitor of ROS generation at Cx-I ⁵⁹, so we
354 consider any contribution from other poorly-characterized oxidation sources to be minimal.
355 Secondly, the mitoSOX signal is impacted both by its distribution between extracellular, cytosolic
356 and mitochondrial compartments ³⁷, and by the primary and secondary filter effects from
357 endogenous chromophores ^{47, 55} (see methods section 2.4). However, our use of fully-oxidized

358 mitoSOX as a control (Supplementary Figure 1) ensures that any changes in the mitoSOX signal
359 we observed originated only from *in-situ* probe oxidation, and not from redistribution of the
360 probe or changes in the absorbance of myoglobin, cytochromes, etc.

361 While mitoSOX does confer a number of limitations regarding the assignment of the signal
362 to a particular reactive oxygen species, similar issues of probe specificity are also broadly
363 applicable to genetically encoded biosensors^{72, 73}. In addition, more precise analytical methods
364 such as LC-MS separation of mitoSOX oxidation products^{37, 74} require time-consuming isolation
365 steps that may release components that further oxidize the probe during isolation, and so may
366 not be readily compatible with the rapid kinetics of the events observed herein. The development
367 of more precise genetically-encoded and non $\Delta\Psi_m$ dependent mitochondrial ROS probes may
368 therefore be useful in future studies.

369 Overall, the findings herein demonstrate that blocking MCT-1 in cardiac IR leads to
370 inhibition of succinate release, which worsens IR injury due to enhanced mitochondrial ROS
371 generation. Concurrent inhibition of Cx-II abrogates these effects, highlighting the importance of
372 succinate oxidation at Cx-II in the pathology of IR injury. Furthermore, these events appear to lie
373 temporally upstream of PT pore opening. Future experiments could explore the role of local
374 succinate signaling in the heart, to determine if SUCNR1 may modulate responses to IR in a
375 manner independent of inflammatory cells, providing further insight on the delicate balance of
376 succinate dynamics in IR injury.

377

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382

383 **Author Contribution Statement**

384 ASM and PSB designed the research. ASM and SMN performed experiments. ASM and
385 PSB analyzed the data and wrote the manuscript. All authors approved the final version of the
386 manuscript.

387

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391

392 **Conflict of Interest:** The authors declare that they have no conflicts of interest.

393

394 **Data Availability Statement**

395 All original data used to prepare the figures is available in spreadsheet form, on the data
396 sharing website FigShare (<https://doi.org/10.6084/m9.figshare.19319627>, to be unembargoed
397 upon publication).

398

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- 624

625 **FIGURE LEGENDS**

626 **Figure 1. Fluorescence Cardiac Perfusion System. (A):** Schematic showing arrangement of the
627 perfused mouse heart relative to light source and photomultiplier tube (PMT). **(B):** Photograph
628 of heart in-situ prior to mounting inside spectrofluorometer. **(C):** Photograph of cardiac perfusion
629 apparatus from above mounted inside spectrofluorometer. Components of the system are
630 labeled, including a custom 3D-printed water-jacketed cuvet holder, heated umbilical lines to
631 deliver KHB at 37°C to the heart, and vacuum line to remove perfusate/effluent. **(D):** NAD(P)H
632 autofluorescence (340/460 nm) during ischemia and reperfusion. Traces in gray are individual
633 hearts, with the average (N=7) shown in black. **(E):** Flavoprotein autofluorescence (462/520 nm)
634 during ischemia and reperfusion. Traces in gray are individual hearts, with the average (N=4)
635 shown in black. Fluorescence data are normalized to pre-ischemic levels. **(F):** Schematic showing
636 design of the 7 experimental perfusion conditions examined herein. Where indicated, S1QEL1.1
637 (Cx-I ROS inhibitor), dimethylmalonate (Cx-II inhibitor), AR-C155858 (MCT-1 inhibitor) or atpenin
638 A5 (Cx-II inhibitor) were administered at the listed concentrations. Colors and terms used to
639 denote each condition are used throughout the remainder of the Figures.

640
641 **Figure 2. Cardiac Function vs. mitoSOX Fluorescence During IR. (A):** Cardiac functional
642 measurements. Graph shows heart rate x left ventricular developed pressure (i.e., rate pressure
643 product, RPP). The period of 25 min. ischemia is indicated. Data are means \pm SEM, N= 16. **(B):**
644 Corrected mitoSOX fluorescence (510/580 nm) during IR. Fluorescent data were processed as
645 described in the methods and as shown in Supplemental Figure 1, to correct for changes in
646 absorbance of endogenous cytochromes at the fluorescence wavelengths. Data are means \pm
647 SEM, N= 6. Note: mitoSOX fluorescent measurements were not performed for the entire
648 perfusion time for all hearts in panel A (e.g., some hearts were used for controls or other
649 measurements), hence different N between panels.

650
651 **Figure 3. Impact of S1QEL on IR injury dynamics.** Hearts were treated with the Cx-I Q-site ROS
652 generation inhibitor S1QEL1.1 for 5 min. pre- and post-ischemia (Figure 1 schematic). **(A):** Control
653 mitoSOX fluorescent data during the first 5 min. of reperfusion (from Figure 2B) for comparative

654 purposes. **(B)**: mitoSOX fluorescent data from S1QEL treated hearts. Gray traces show individual
655 data, with averages shown as bold line. **(C)**: Calculated slopes from 1st minute of reperfusion, for
656 the data in panels A/B. For each condition, individual data points are shown on the left, with
657 mean \pm SEM on the right (N for each condition can be seen from number of data points). p values
658 (ANOVA followed by unpaired Student's t-test) for differences between groups are denoted. **(D)**:
659 NAD(P)H autofluorescence (340/460 nm) during ischemia and reperfusion. Traces in gray are
660 individual hearts, with the average (N=5) shown in green. **(E)**: Flavoprotein autofluorescence
661 (462/520 nm) during ischemia and reperfusion. Traces in gray are individual hearts, with the
662 average (N=5) shown in green. For panels D and E, the average data from control hearts (from
663 Figure 1) is shown in black for comparative purposes. Fluorescence data are normalized to pre-
664 ischemic levels.

665
666 **Figure 4. Impact of AR and Cx-II Inhibitors on mitoSOX Fluorescence at Reperfusion. (A-E)**:
667 Corrected mitoSOX traces 5 min. pre- and post-reperfusion, for the labeled conditions. Gray
668 traces are individual data, with averages shown as bold colored lines (see Figure 1 for color
669 scheme). **(F)**: Calculated slopes from 1st minute of reperfusion, for the data in panels A-E. For
670 each condition, individual data points are shown on the left, with mean \pm SEM on the right (N for
671 each condition can be seen from number of data points). p values (ANOVA followed by unpaired
672 Student's t-test) for differences between groups are denoted.

673
674 **Figure 5. Impact of AR and Cx-II Inhibitors on Outcomes of IR Injury. (A)**: Cardiac functional
675 measurements during IR. Graphs show heart rate \times left ventricular developed pressure (i.e., rate
676 pressure product, RPP) for each of the conditions examined. The period of 25 min. ischemia is
677 indicated. Data are means \pm SEM. **(B)**: Quantitation of percent functional recovery, i.e., cardiac
678 function at 60 min. of reperfusion as a percentage of that immediately before ischemia (-25 min.
679 time point). For each condition, individual data points are shown on the left, with mean \pm SEM
680 shown on the right (N for each condition can be seen from number of data points). p values
681 (ANOVA followed by unpaired Student's t-test) for differences between groups are denoted
682 above the data. **(C)**: Myocardial infarct size for each condition. Images above the graph show

683 representative TTC stained heart cross-sections, with pseudo-colored mask used for planimetry
684 below. Graph shows data for each condition, with individual data points on the left, mean \pm SEM
685 on the right (N for each condition can be seen from number of data points). p values (ANOVA
686 followed by unpaired Student's t-test) for differences between groups are denoted above the
687 data.

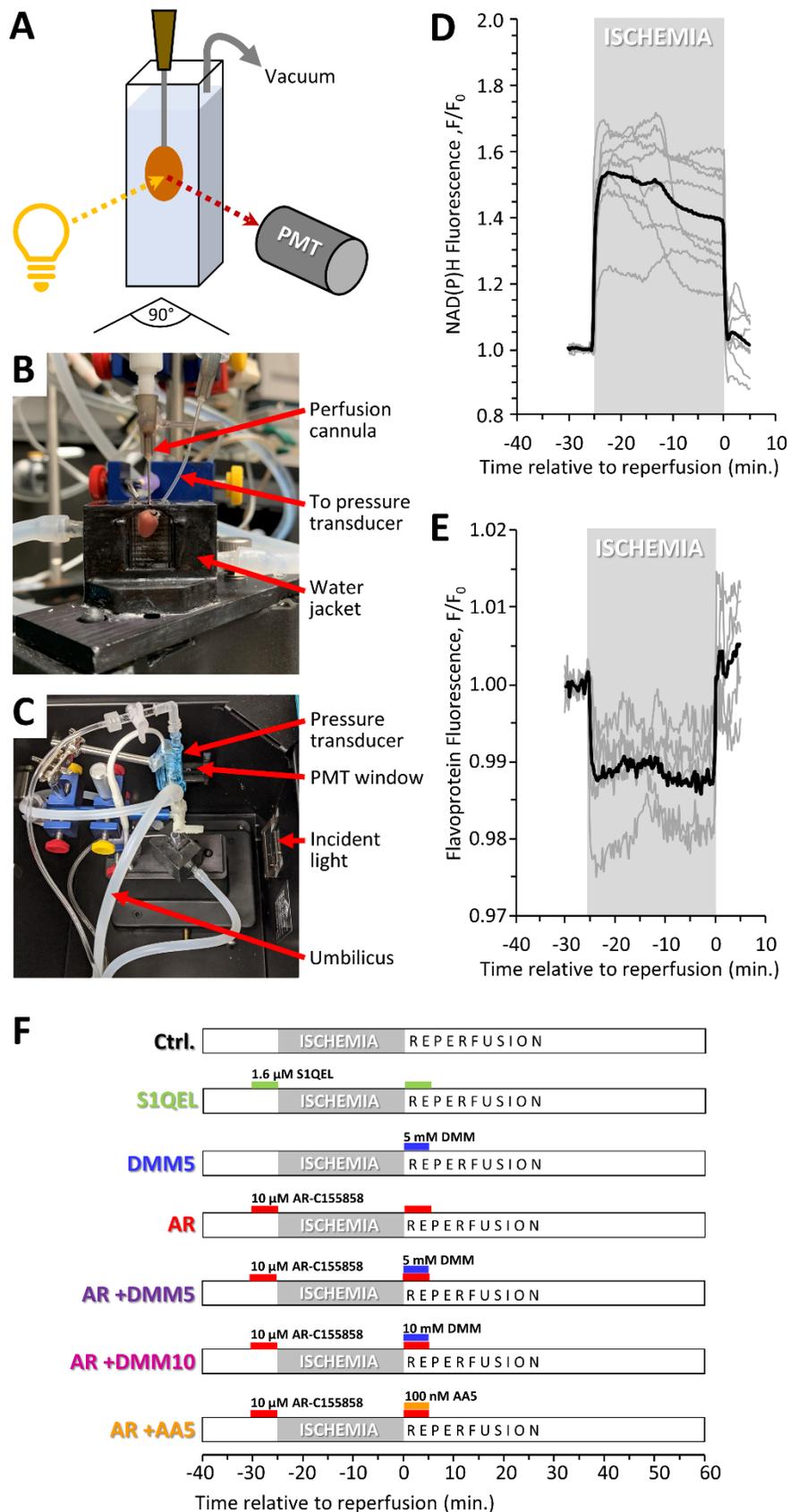
688
689 **Figure 6. Correlations Between mitoSOX Fluorescence During Reperfusion and IR Outcomes.**

690 Graphs show correlation between mitoSOX slope data and **(A)**: infarct size or **(B)**: functional
691 recovery. Each data point is the mean \pm SEM for a condition (see color scheme in Figure 1).
692 mitoSOX data are from Figure 3F and Supplemental Figure 5B. Infarct data are from Figure 4C
693 and Supplemental Figure 5E. Functional data are from Figure 4B and Supplemental Figure 5D.
694 Linear curve fits are shown, with correlation coefficient (r^2) listed alongside.

695
696 **Figure 7. Temporal Relationship Between mitoSOX and PT Pore Opening During IR. (A):**

697 Schematic showing perfusion conditions. Where indicated, mitoSOX (ROS indicator), TMRE ($\Delta\Psi_m$
698 indicator) and/or CsA (PT pore inhibitor) were administered at the listed concentrations. **(B/C)**:
699 mitoSOX signal during early reperfusion under control or CsA condition. The last 5 min. of the
700 ischemic period is indicated. Gray traces show individual data, with averages shown as bold lines.
701 Inset to panel C shows averages for control and CsA superimposed. **(D/E)**: Normalized TMRE
702 fluorescence during early reperfusion under control or CsA condition. The last 5 min. of the
703 ischemic period is indicated. Gray traces show individual data, with averages shown as bold lines.
704 Inset to panel C shows averages for control and CsA superimposed. Red arrow indicates the point
705 at which traces diverge, 5 min. into reperfusion.

706



707

FIGURE 1

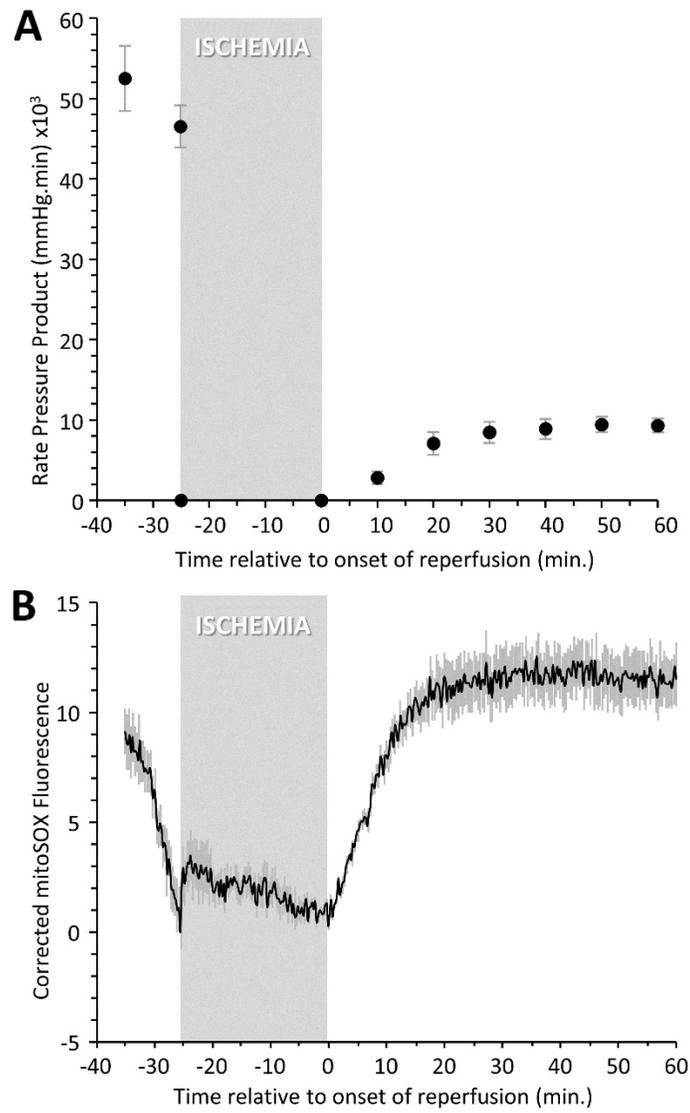
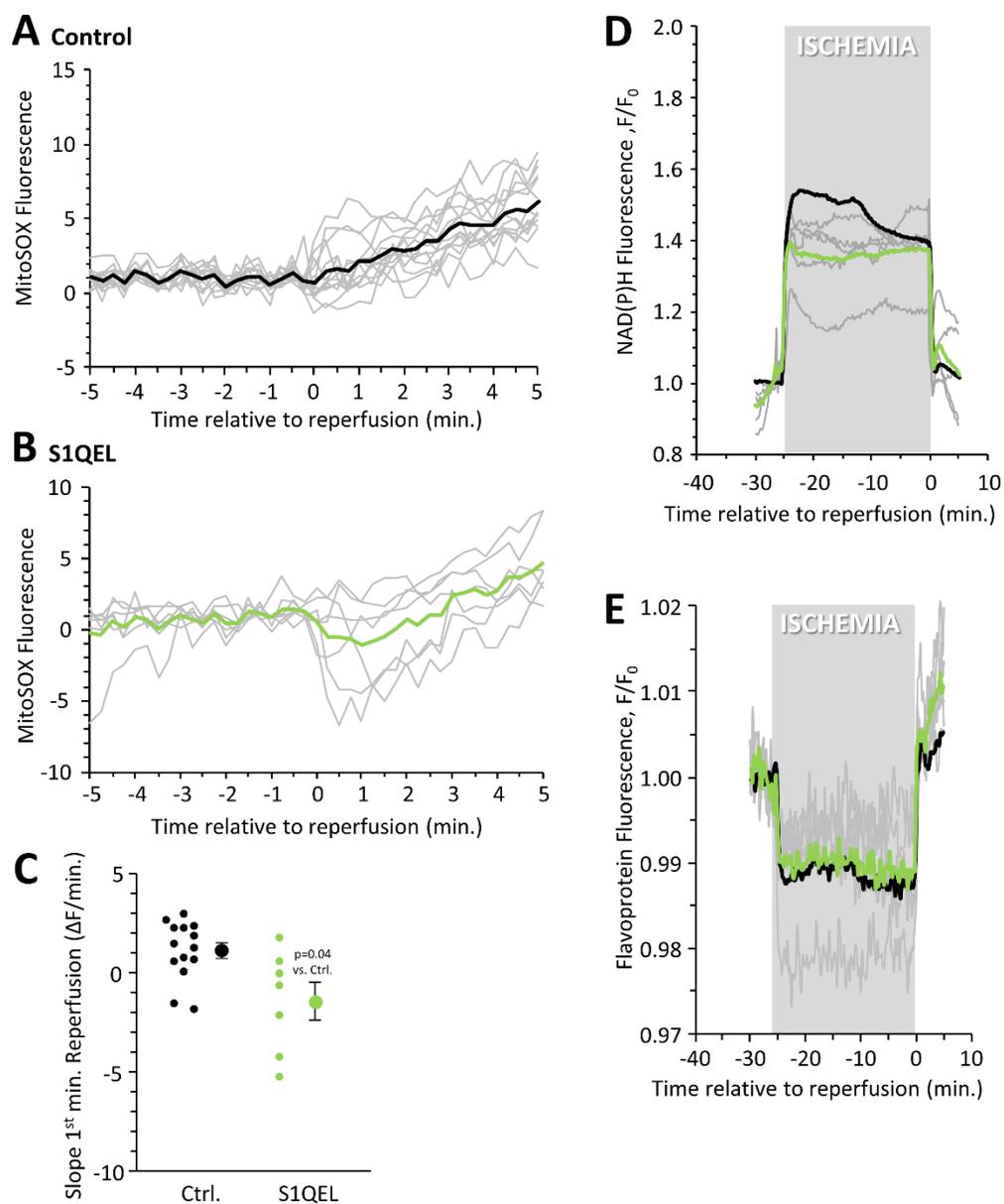


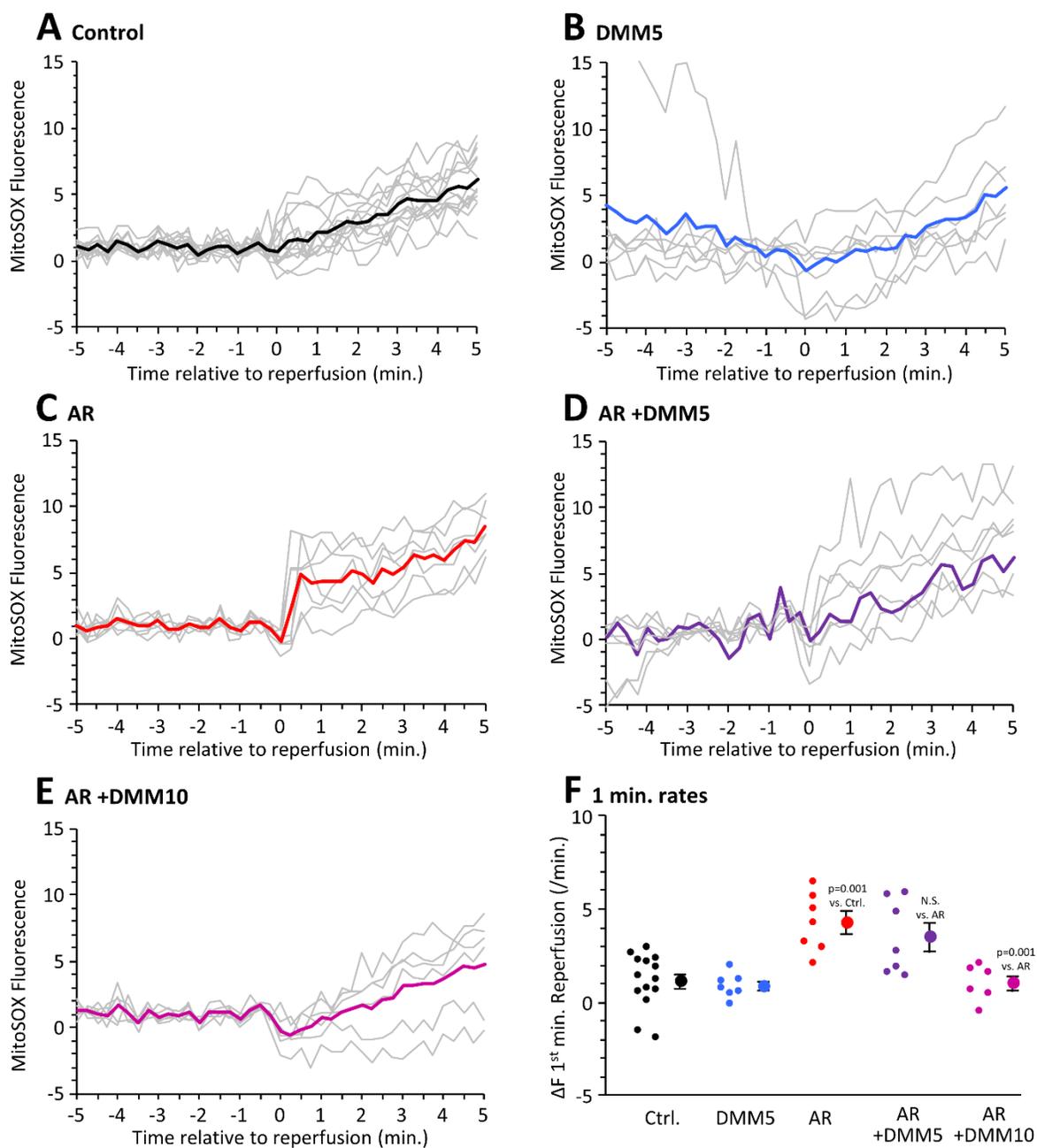
FIGURE 2

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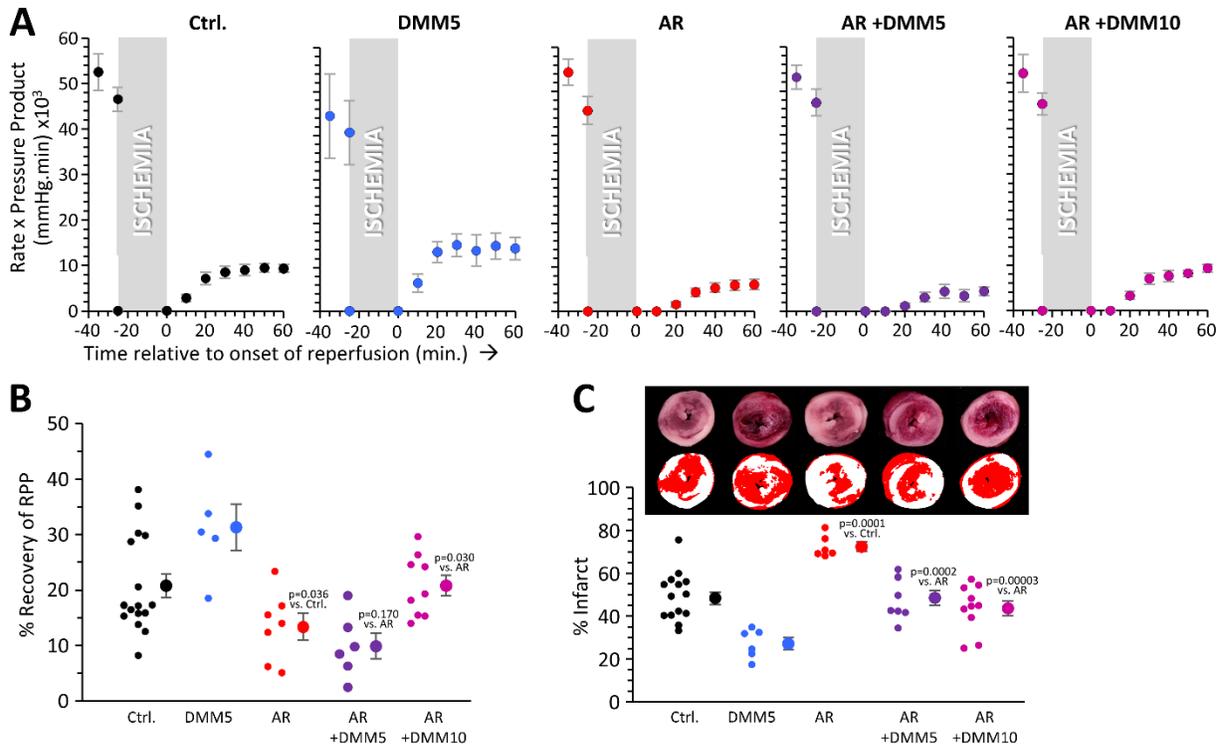
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FIGURE 3



715
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FIGURE 4



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FIGURE 5

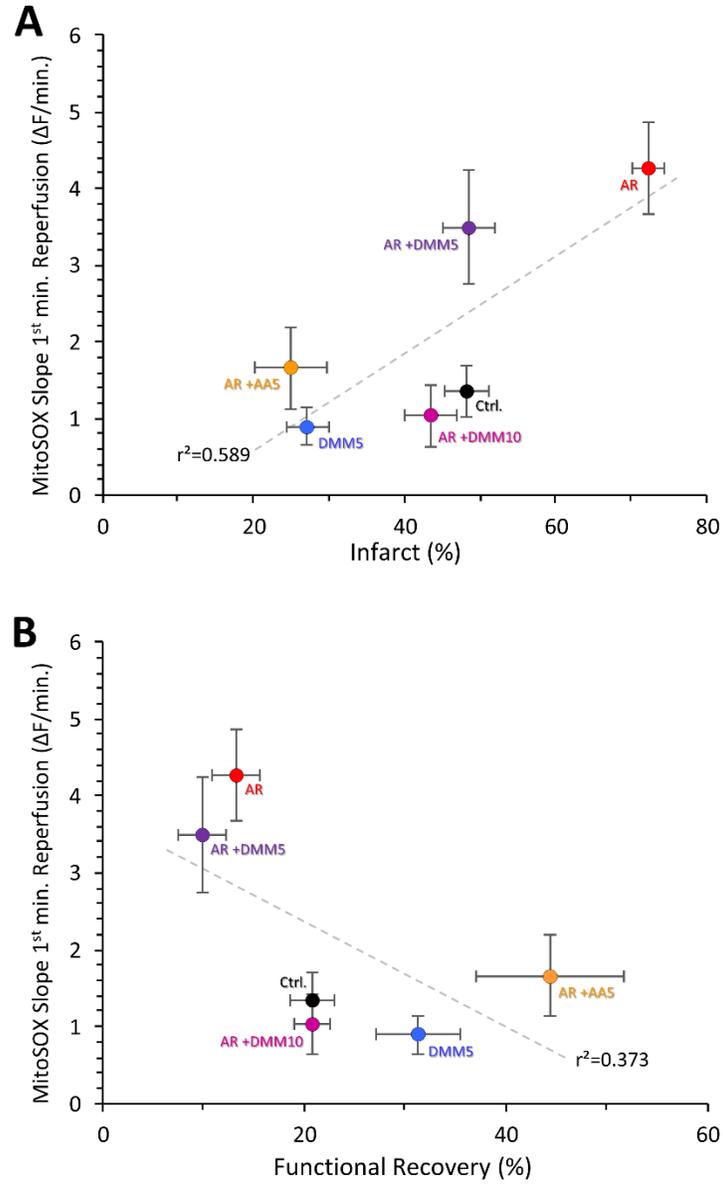
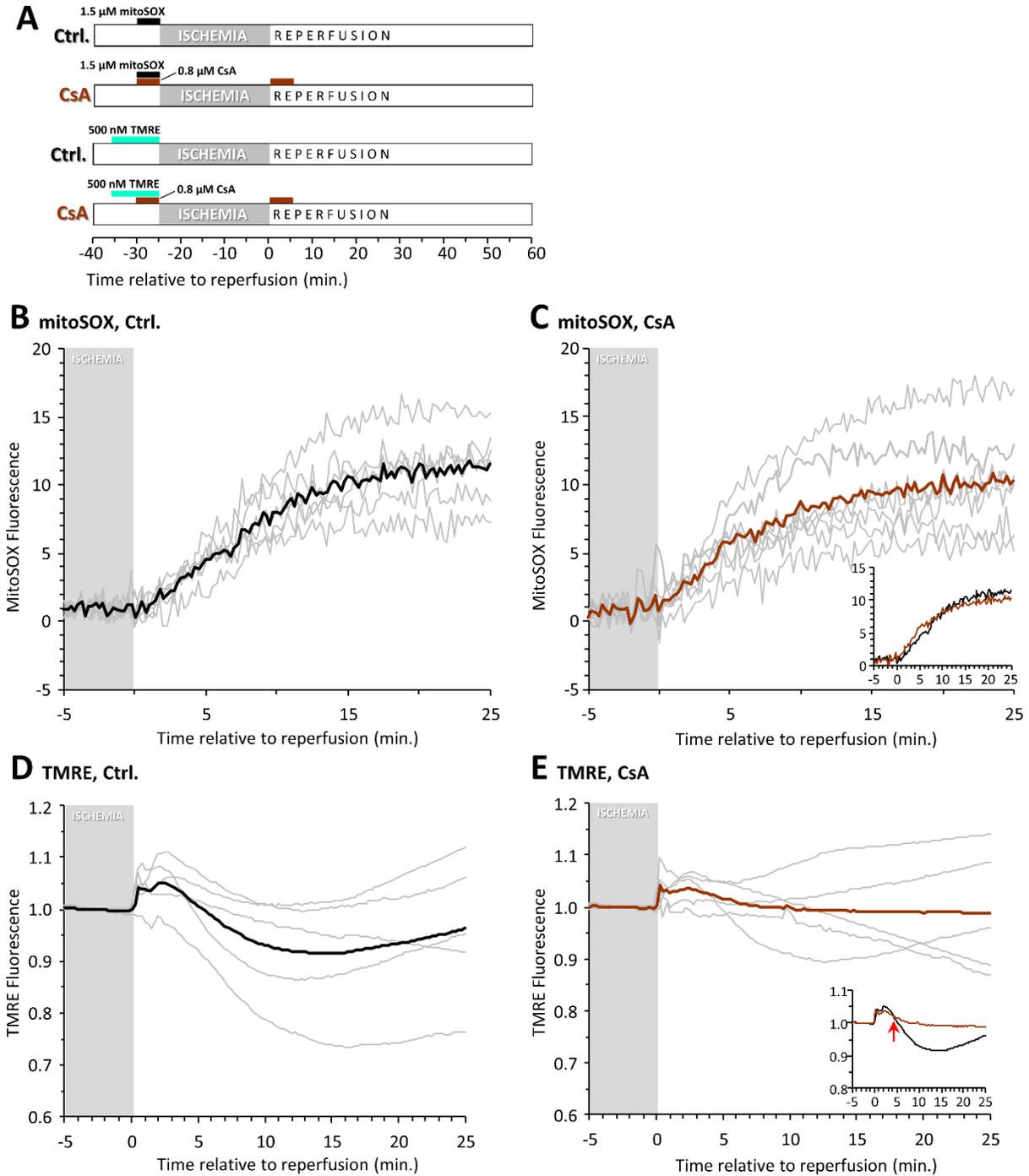


FIGURE 6

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720



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722
723

FIGURE 7