

1 **Modelling cardiac dysfunction following traumatic haemorrhage** 2 **injury: impact on myocardial integrity.**

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10 **Abstract**

11 Cardiac dysfunction (CD) importantly contributes to mortality in trauma patients, who
12 survive their initial injuries following successful haemostatic resuscitation. This poor
13 outcome has been correlated with elevated biomarkers of myocardial injury, but the
14 pathophysiology triggering this CD remains unknown.

15 We investigated the pathophysiology of acute CD after trauma using a mouse model of
16 trauma haemorrhage shock (THS)-induced CD with echocardiographic guidance of fluid
17 resuscitation, to assess the THS impact on myocardial integrity and function. Mice were
18 subjected to trauma (soft tissue and bone fracture) and different degrees of haemorrhage
19 severity (pressure controlled ~MABP <35mmHg or <65mmHg) for one hour, to characterise
20 the acute impact on cardiac function. In a second study, mice were subjected to trauma and
21 haemorrhage (MABP<35mmHg) for one hour, then underwent two echocardiographic-
22 guided resuscitations to baseline stroke volume at 60 min and 120 min, and were monitored
23 up to 180 min to study the longer impact of THS following resuscitation. Naïve and sham
24 animals were used as controls.

25 At 60 min post-THS injury, animals showed a lower cardiac output (CO) and stroke volume
26 (SV) and an early rise of heart fatty acid-binding protein (H-FABP=167±38 ng/ml; 90%
27 increase from shams, 3.54±3.06ng/ml), when subjected to severe haemorrhage and injury.
28 Despite resuscitation, these animals maintained lower CO (6 ml/min vs 23ml/min), lower SV
29 (10 µl vs 46 µl; both ~75% decreased), and higher H-FABP (levels (340±115 ng/ml vs
30 10.3±0.2ng/ml; all THS vs. shams, P<0.001) at 180 min post-TH injury. Histopathological
31 and flow-cytometry analysis of the heart confirmed an influx of circulatory leukocytes,
32 compared to non-injured hearts. Myocardial injury was supported by an increase of troponin I
33 and h-FABP and the widespread ultrastructural disorganisation of the morphology of
34 sarcomeres and mitochondria. DNA fragmentation and chromatin condensation driven by
35 leakage of apoptosis-inducing factor (AIF) may suggest a mitochondria-driven progressive
36 cell death.

37 THS modelling in the mouse results in cardiomyocyte damage and reduced myocardial
38 function, which mimics the cardiac dysfunction seen in trauma patients. This CD model may,
39 therefore, provide further understanding to the mechanisms underlying CD and act as a tool
40 for developing cardioprotective therapeutics to improve survival after injury.

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46 **Key words:** Trauma, Cardiac dysfunction, Myocardial damage

47 **Heading Title:** Trauma associated cardiac dysfunction mechanism

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49

50 **Introduction**

51 Trauma is a large and growing problem worldwide, accounting for 10.1% of the global
52 burden of disease [1], with half of all trauma deaths being due to excessive bleeding and the
53 subsequent severe shock [2, 3]. New paradigms of haemostatic resuscitation to manage
54 coagulopathy have led to large decreases in mortality worldwide [4, 5]. With more patients
55 surviving the initial bleeding episode, cardiac dysfunction is increasingly common and an
56 important determinant of outcome. Over half of all critically injured trauma patients admitted
57 to intensive care develop cardiovascular dysfunction within the first 48 hours, of which 20%
58 will die [6]. Identifying these patients early in their care and rescuing them from this
59 downward trajectory would have a dramatic impact on trauma mortality.

60 Cardiac dysfunction in trauma patients is initially difficult to recognise, as it develops within
61 the context of hypovolaemia and a widespread inflammatory response [7]. Patients often
62 initially show a normal cardiac response with a high cardiac output, but over a relatively short
63 period they experience a dramatic fall in SV and CO, despite inotrope and vasopressor
64 support. We have previously shown that trauma patients have elevated levels of biomarkers
65 of myocardial damage within the first 2 hours of injury, and this is associated with increased
66 risks of adverse cardiac events and mortality [8, 9]. Cardiac histopathology in non-survivors
67 has shown multiple pathological identifiers of indirect or secondary cardiac injury [10, 11].
68 The pathophysiology and mechanisms of secondary cardiac dysfunction are unknown, with
69 most critical care studies limited to sepsis [12]. There is some pre-clinical evidence for the
70 development of cardiac injury and dysfunction arising as an indirect consequence of trauma
71 and haemorrhage in pigs [13] and in rodents [14-16], suggesting a local cardiac inflammatory
72 response as the main driver of cardiomyocyte structural and functional damage.

73 Our overall **aim** for this study was to investigate the pathophysiology of cardiac dysfunction
74 after trauma by implementing a clinically relevant murine model of post trauma haemorrhage
75 cardiac dysfunction. Specifically, **our objectives were** to: 1) determine the nature and extent
76 of myocardial damage and cardiac dysfunction in an un-resuscitated model of trauma
77 haemorrhage; 2) determine the progression of cardiac dysfunction over time in a resuscitated
78 model of trauma haemorrhage; and 3) explore the inflammatory myocardial response and
79 ultrastructural integrity, to elucidate possible mechanistic pathways for cardiomyocyte cell
80 damage.

81

82 **Materials and methods**

83

84 **Ethical statement**

85 All animal procedures were carried out under a Project Licence (PC5F29685) approved by
86 the Animal Welfare and Ethical Review Body at Queen Mary University of London and the
87 UK Home Office, in accordance with the EU Directive 2010/63/EU. All animal facilities and
88 suppliers have been approved by the UK Home Office Licensing Authority and meet all
89 current regulations and standards for the UK. A total of 98 mice were used for the work
90 described in this study (details in supplemental Table 1).

91 For this study we used n=6-10 animals per group, to provide a valuable discriminatory power
92 of 90% with a significant level $\alpha = 0.05$ to detect up to 15-20% relative differences in primary
93 outcomes (lactate levels, cardiac function, cardiac injury biomarkers). Experimental planning
94 for data randomization and blinded data acquisition and analysis was carried out following
95 the ARRIVE guidelines [17].

96 **Animal housing and husbandry**

97 Adult male C57BL/6 mice (weight range 25-30g; 9-11 weeks old) were obtained from
98 Charles River Laboratories (Margate, UK). Health screens provided by the official vendor
99 indicated that animals were free of known pathogens in accordance with FELASA guidelines
100 for health monitoring of rodent colonies [18]. Animals were housed in groups of 4-6 per
101 individually ventilated cage (IVC; Allentown Europe, UK), in a 12 h light dark cycle (06:30 -
102 18:30 light; 18:30 - 06:30 dark), with controlled room temperature ($21 \pm 1^\circ\text{C}$) and relative
103 humidity (40-60 %). Animals were allocated to cages on arrival and remained in the same
104 social group throughout the study, including a 7-day acclimatization phase prior to any study,
105 with ad libitum access to standard diet and water.

106 **Induction of haemorrhagic traumatic injury**

107 Animals were anaesthetised (Isoflurane: 4-5% induction, 0.5-1.5% for maintenance in 0.8-1lt
108 /min 100% medical O₂). Anaesthesia depth was controlled clinically and by haemodynamic
109 monitoring (Mean Arterial Blood Pressure; MABP). Core temperature was maintained at
110 $36 \pm 1^\circ\text{C}$ throughout the study with a homoeothermic blanket (Harvard Apparatus Ltd., UK)
111 and heat lamps. All experiments were carried out under terminal anaesthesia with no
112 recovery, and all animals were humanely culled at the end of the experiment.

113 A 1cm incision was made in the middle of the cervical skin and the left jugular vein was
114 cannulated [polyethylene tubing pre-flushed with heparinised saline (25 IU/mL); Portex.
115 Smith's Medical Int. Ltd. Kent, UK] The right carotid artery was then cannulated in the same
116 fashion and connected to a pressure transducer (Capto SP 844, AD Instruments, UK) attached
117 to a PowerLab 8/30 (ML870, AD Instruments Ltd, Oxford, UK) to monitor MABP using the
118 LabChart software (ADInstruments Ltd, UK). The neck incision was covered and regularly
119 checked for evidence of line displacement and/or bleeding. If either of these developed, the
120 animal was euthanised and removed from the study.

121 A 2cm midline laparotomy was then performed, and the rectus muscle was crushed using
122 forceps in a systematic fashion in each animal. The abdominal area was examined to exclude
123 inadvertent iatrogenic injury and/or bleeding, and then the incision was closed using 5.0-
124 prolene suture material (Ethicon, UK). Immediately thereafter, animals were subjected to a
125 bilateral hind limb fracture. Fractures were performed using a closed, manual 3-point bending
126 technique. Following 5 minutes' stabilisation after traumatic injury, a 'baseline' MABP was
127 recorded. Then, a pressure-controlled haemorrhage via the carotid cannula was induced to

128 achieve a target MABP of 30 – 40mmHg to reach a Traumatic haemorrhagic shock state
129 (THS). Animals underwent a 60-minute observation period, during which the target blood
130 pressure (30-40mmHg) was maintained with removal of blood as required via the carotid
131 cannula. Shed blood was kept warm in a heparinised 1mL syringe (25 IU/mL), which was
132 occasionally agitated to prevent thrombus formation. In-dwelling vascular catheters were
133 intermittently flushed with small volumes of heparinised saline and wound sites checked for
134 signs of haemorrhage. All volumes of heparinised saline were taken into account when
135 recording volumes of shed and reinfused blood.

136 Sham controls underwent cannulation of the carotid artery for invasive MABP monitoring
137 only, without trauma, haemorrhage or fluid resuscitation.

138 Echocardiography of the left ventricle (LV) was performed at baseline (prior to any traumatic
139 haemorrhagic intervention), and at defined time-points after injury to assess the impact of
140 THS injury at 60 (T60), 120 (T120) and 180 (T180) min post THS, using the Vevo 770 high-
141 resolution system (Visualsonics Inc. Toronto, Canada). M-mode short axis measurements
142 were used to calculate stroke volume (SV), left ventricular end-diastolic volume (LVEDV)
143 and cardiac output (CO). Measurements were carried out in triplicate.

144 The acute impact of THS in cardiac function was studied in animals subjected to sham,
145 trauma only, THS to 60-70 mmHg MABP or THS to 30-40 mmHg MABP procedures (n=6
146 /group) following 1h post-intervention, without any resuscitation. At 60 min post THS (T60),
147 cardiac function and shock status were assessed and animals were culled via terminal
148 exsanguination and blood samples processed for further analysis (Fig.1A).

149 To investigate the CD associated with THS, the THS model with a 30-40 mmHg MABP was
150 extended up to 3h through 2 resuscitation phases at T60 and T120 (THS and sham,
151 n=10/group) and terminated at T180 . Echocardiography was carried out at T60, T120 and
152 T180 to assess cardiac function and also to guide the resuscitation volume requirements to
153 restore the SV to baseline levels during each resuscitation at T60 and T120 (Fig. 2A).

154 For the first resuscitation (RESUS 1 at T60), the whole shed blood was transfused as a bolus
155 over 5 minutes *via* the jugular catheter, and boluses of warmed Hartmann's solution
156 (Vetivex11. Dechra Veterinary Products, Shrewsbury, UK) were then administered to reach
157 SV baseline. In the second resuscitation phase (RESUS 2 at T120), a bolus of Hartmann's
158 alone was administered over 5-10 minutes. At each resuscitation, animals received volume
159 resuscitation to restore left ventricular stroke volume to baseline. During the second
160 resuscitation, if the target SV was not reached and yet there was no incremental response to
161 further fluid boluses (and in the confirmed absence of bleeding from wound sites), this was
162 deemed to represent the completion of the resuscitation phase. All studies were terminated at
163 T180, through controlled exsanguination via carotid line and confirmation of death via HR
164 and MABP.

165 At the end of the experiment, sham animals were culled *via* terminal exsanguination and
166 blood samples and tissue taken.

167 **Assessment of shock status**

168 MAPB and heart rate were measured throughout haemorrhage and resuscitation. Blood
169 lactate was measured and used as an index of shock and tissue perfusion. Arterial lactate
170 concentrations (mmol/L) were assessed using the Accutrend Lactate monitor (Roche,
171 Mannheim, Germany). Echocardiography measured at T60 and T120 before and after

172 resuscitation, and at T180 was carried out to assess cardiac function (left ventricle SV).
173 Hematocrit (Hct; %) and hemoglobin (Hb; g/dL) were measured with a ProCyte Dx
174 Hematology Analyzer (IDEXX Europe B.V, Hoofddorp, The Netherlands).

175 **Terminal blood sampling and tissue storage**

176 Terminal exsanguination was performed via the carotid catheter only and processed for serum
177 separation (Z-Gel microtube, Starstedt. Westphalia, Germany). Serum samples were stored at
178 -80°C. The hearts were removed immediately after the end of the experiment and processed
179 accordingly to further test specifics.

180 **Cardiac biomarker assessment**

181 Serum heart fatty acid-binding protein (H-FABP) levels were assayed using commercially
182 available mouse specific enzyme-linked immunosorbent assay (ELISA Cat. No. HFABP-1)
183 kit (supplied by Life Diagnostics Inc., West Chester, PA, USA.). ELISAs were performed in
184 accordance with the manufacturer's instructions. Standard curves for all ELISAs were plotted
185 and dilution-corrected sample concentrations were interpolated from the standard curve.

186 **Immunohistochemistry analysis**

187 At 3h THS, a subset of animal (n=3 naïve, n=5 THS) were terminally anaesthetised and
188 cardiac tissue was immediately fixated (10% NBF). Naïve animals were housed together
189 under the same conditions but did not have any surgical intervention, trauma, haemorrhage or
190 fluid resuscitation. Cardiac tissue was paraffin-fixed for histology and immunohistochemistry
191 (IHC). Sections (7µm) were deparaffinised and hydrated through xylene and ethanol baths.
192 Sections were subjected to antigen retrieval (10mM of citrate buffer, pH 6.0, 10 min in
193 microwave) and then cooled at room temperature. Tissue was permeabilised with 10%
194 Triton-X in PBS for 15 min and then blocked with 10% goat or donkey serum, 1% bovine
195 serum albumin (BSA) in PBS for an hour, at room temperature (RT). The following primary
196 antibodies diluted in blocking solution were used (overnight incubation in a humid chamber
197 at 4°C): goat anti-ionized calcium binding adaptor molecule 1 (Iba-1; for macrophages and
198 monocytes 1:500; Wako Chemicals USA, Inc., Richmond, VA; Cat#ab109497), rat anti-
199 Lymphocyte antigen 6 complex, locus G (Ly6G-clone 1A8; for neutrophils 1:200;
200 BioLegend, London, UK; Cat# 127602); rabbit anti-mouse cleaved caspase-8 1:200 (Asp387;
201 Cell Signalling Tec.; Cat#8592); rabbit anti-mouse MTCO2 (1:125; Abcam plc, Cambridge,
202 UK; Cat#ab110258) or rabbit polyclonal anti-AIF (1:100; Abcam, UK; Cat#ab2086). The
203 secondary antibodies were donkey anti-goat IgG 568, goat anti-rat 594 IgG, goat anti-rabbit
204 IgG 488 or goat anti-mouse IgG 594 (Molecular Probes, Leiden, the Netherlands; 1:400 in
205 PBS). Sudan black (0.3% w/v in 70% ethanol) was used to reduce autofluorescence and
206 Hoescht 33342 stain (Sigma-Aldrich, Gillingham UK; 1µg/ml of PBS) was used to visualise
207 nuclei. Slides were mounted and cover-slipped using Vectashield mounting medium (H-1000,
208 Vector Laboratories, Burlingame, CA).

209 For calculations, 2 slides per animal with at least 18 fields were captured across the short-axis
210 LV myocardium stained section. Images were viewed (x400) and photographed using a Zeiss
211 Axioskop 2 microscope (Carl Zeiss, Jena, Germany) with a Hamamatsu camera (C4742-95;
212 Hamamatsu Photonics K.K., Hamamatsu, Japan). Analysis was done using the ImageJ
213 analysis software for counting the number of positively stained cells. Total nuclei count was
214 measured to normalise percentage of positive stained cells across fields. All imaging

215 acquisition and analysis were carried out blinded to the experimental interventions, and data
216 were only allocated to the specific experimental group at the end of the analysis.

217 **Flow Cytometry full heart analysis**

218 At 3h post-study, a subset of animals (n=6 naïve, n=5 sham-subjected to vessel cannulation
219 and terminal anaesthesia only, n=5 THS) were intracardially perfused with heparinised saline
220 under terminal anaesthesia. The hearts were then immediately isolated, cut into 1 mm³ pieces
221 and dissociated by incubation with papain (Merck, UK) and DNase I (ThermoFisher
222 Scientific, UK) for 30 minutes at 37°C. Following lysis of residual red blood cells (RBC
223 Lysis Buffer, Biolegend, UK), cell suspensions were incubated with CD16/CD32-block
224 (Biolegend, UK) for 30 minutes at 4°C, followed by incubation for 30 minutes at 4°C with
225 PE-conjugated anti-CD45 to define immune cell populations, and FITC-conjugated anti-
226 Ly6C/G (clone RB6-8C5) and APC-conjugated anti-F4/80 (all ThermoFisher Scientific, UK)
227 to differentiate neutrophils (F4/80Neg, Ly6C/GHi) from pro-inflammatory (F4/80Pos,
228 Ly6C/GInt) and anti-inflammatory (F4/80Pos, Ly6C/GLow) monocytes/macrophages [19,
229 20]. Cells were then analysed by flow cytometry using a BD FACSCanto II instrument (BD
230 Biosciences) and FlowJo 8.8.1 software (TreeStar Inc., FL, USA). In all cases, 20,000 singlet
231 CD45Pos events were analysed per sample; positive staining was defined by inclusion of
232 fluorescence-minus-one controls for all antigens.

233 **Transmission electron microscopy (TEM)**

234 TEM was used to study micro-structural changes associated to specific cell injury and death
235 in the cardiomyocyte following THS. In a subset cohort of THS and sham animals from the
236 3h studies, the animals were humanely killed by overdose of anaesthesia and the heart tissue
237 immediately dissected, cut into smaller tissue specimen (~1mm³), and fixed in 2%
238 glutaraldehyde. Fixed tissue was washed three times in cacodylate buffer 0.1 M pH 7.4 and
239 then incubated in 1% osmium tetroxide in ddH₂O for 2 hrs at 4°C. After 3 washes in ddH₂O,
240 specimens were dehydrated (progressive incubation from 25% to 100% acetone) following
241 by impregnation in an increasing concentration of the epoxy resin Araldite 502 (from 25 to
242 100% of araldite in acetone) used as the embedding medium for TEM. Samples were stored
243 in fresh araldite for up to 72hrs and then stored at 60°C for 48hrs until the block was hard.

244 **Protein expression in the heart**

245 Western blotting (WB) was used to study biochemical changes associated to cardiomyocyte
246 injury (troponin I, H-FABP) and cell death pathways (Caspase 8 and Apoptosis Inducing
247 Factor-AIF) following THS. Tissue from THS (n=6) and sham (n=6) animals was used. At 3h
248 post-study, animals were deeply anaesthetised and heart tissue was immediately dissected,
249 weighted and immersed in RIPA buffer [0.1g/1ml w/v; 50 mM TrisHCl (pH 7.4), 150 mM
250 NaCl, 1% Triton X-100, 0.5% sodium deoxycholate, 0.1 % SDS, 1 mM EDTA, 10 mM NaF
251 in ddH₂O] with a cocktail of protease inhibitor and phosphatase inhibitor tablets (Pierce™
252 Protease and Phosphatase Inhibitor Mini Tablets Cat no88668, Sigma-Aldrich, UK). The
253 tissue was minced in RIPA on a cooling slide on dry ice and mash in Dounce homogenizer.
254 The sample was then sonicated (50 pulses in rounds of 10 s with 10 s rest). Samples were
255 spun (@13000 rpm in cooling centrifuge for 20 min) and the supernatant protein sample was
256 collected. Protein concentrations were determined by Bradford assay. Equal amounts of
257 protein sample were mixed with NuPAGE® MOPS SDS Running Buffer. 20X)-NP0001 was
258 made up and a tank filled. Bolt™ 4-12%, 10-well NW04122BOX) Bis-Tris Plus Gel was
259 inserted into the tank; 10ul of samples were loaded with protein ladder (RPN800E-GE health

260 care) into 10 well NuPage Bis-Tris Mini gel. Optical density was determine using ImageJ
261 software (NIH). A cardiac tissue specific, mouse monoclonal troponin I antibody (Ab10231,
262 Abcam, UK; 24kDa) was used for the labelling of troponin. Caspase-8 (rabbit monoclonal
263 anti-cleaved caspase 8; 8592S; Cell Signalling, UK; 41kDa) expression was used as a marker
264 of activation of death receptor initiated cell death pathway. AIF (rabbit polyclonal anti-AIF;
265 Ab2086; Abcam, UK; 67 kDa) expression and location was assessed by WB analysis, with
266 the tissue homogenates being processed for differential centrifugation to assess the sub-
267 cellular compartmentation, removing the large organelles nuclei, cellular debris and intact
268 cells, and allowing for selective extraction of the cytosol and mitochondria using Cytosol and
269 Mitochondrial Extraction Buffer Mix containing DTT and Protease Inhibitors. MTCO2
270 (Anti-MTCO2; Ab198286; Abcam, UK; 1/150 - 25 kDa) and GAPDH (Anti-GAPDH
271 antibody; Ab8245; Abcam, UK) were used as a mitochondrial and cytosol markers,
272 respectively.

273 **Statistical analysis**

274 Lactate levels (mmol/L), MABP measurements (mmHg), resuscitation fluids (ml),
275 echocardiographic LV data (CO=ml/min; SV= μ l; LDEDV= μ l), % of inflammatory cells
276 (neutrophils / monocytes / macrophages) and H-FABP serums levels = ng/ml) were
277 expressed using mean values, with standard deviation (SD). Statistical analysis was
278 performed using GraphPad Prism 8 (GraphPad Software, San Diego, USA). Normally
279 distributed data was analysed using Student's two-tailed t test or one-way ANOVA, followed
280 by Tukey's multiple comparison test. Non-normally distributed data was analysed with
281 Kruskal-Wallis and Mann-Whitney U test analysis. P<0.05 was taken to represent
282 significance. All analysis was performed blinded to the experimental interventions. (Further
283 details in supplementary Table 2).

284 **Results:**

285 **Myocardial injury after trauma and haemorrhage**

286 The trauma haemorrhage model demonstrated progressive myocardial damage and cardiac
287 dysfunction following an hour of trauma and haemorrhage without resuscitation (Figure 1).
288 We examined cardiac effects of trauma alone, trauma and haemorrhage to target MABP of
289 60-70mmHg (THS 60-70); and to a target an MABP of 30-40mmHg (THS 30-40), achieving
290 different depths of shock severity (Figure 1A, supplemental Table 3) Myocardial damage, as
291 measured by serum H-FABP concentrations, increased with increasing model severity, and
292 were significantly elevated for the most severe THS group (THS 30-40 = $166.69 \pm$
293 38.64 ng/ml), compared to sham, trauma only and 60-70 mmHg THS group ($p<0.01$). (Fig
294 1C). At end-experiment, stroke volume and cardiac output were significantly decreased from
295 baseline in all groups (Fig 1D-1F). In the THS 30-40 group, with over 30% blood loss,
296 cardiac output was 87% lower than baseline (Fig 1E) at 1-hour post trauma compared to
297 sham and trauma-only groups ($p<0.001$, Figure 1E).

298 **Cardiac dysfunction following myocardial damage**

299 In order to determine the functional effects of myocardial damage, we extended our TH 30-
300 40mmHg model for 3 hours to include two resuscitation phases at 60 and 120 minutes post-
301 haemorrhage (Figures 2A and 2B). Shock severity worsened over time despite the two
302 resuscitation steps (Figures 2B & C), with serum lactate reaching a median of 7.9 ± 5.2
303 mmol/L at 180 minutes (from 1.4 ± 0.37 mmol/l at baseline $p=0.004$, supplemental Table 4).

304 MABP increased immediately after resuscitation, but then progressively declined, despite
305 transient increases with volume resuscitation. End-experiment MAP was 44% lower than
306 post Resus-1 levels (43.3 ± 8 vs 77.4 ± 5 mmHg Figure 2B; supplemental Table 4). Hb
307 concentration and Hct were not significantly different between the THS group and sham, at
308 3h (Hb 14.5 ± 1.6 vs 13.2 ± 1.4 and Hct 47.5 ± 8.7 vs 43.1 ± 5.4 , in sham and THS groups,
309 respectively; supplemental Table 2 and Fig 1).

310 On functional echocardiographic assessment, volume resuscitation to normalise stroke
311 volume did not maintain LVEDV, stroke volume or cardiac output. Mean LVEDV was only
312 23% of baseline at end-experiment ($16.40 \mu\text{l}$ vs $76.17 \mu\text{l}$; $p < 0.0001$); stroke volume was 22%
313 of baseline ($10.5 \mu\text{l}$ vs $47.1 \mu\text{l}$, $p < 0.0001$) and total cardiac output was 30% from baseline
314 ($5.8 \mu\text{l}$ vs $22.4 \mu\text{l}$, $p < 0.0001$, Figures 2E-G, Table S2). In line with these findings, H-FABP
315 concentrations continued to increase from T60 min post-THS, rising to a mean of 340 ng/ml
316 at end-experiment compared to 116.7 ng/ml at T60 ($p = 0.003$) and 10.3 ng/ml compared to
317 sham animals at end-experiment, $p < 0.001$) (Table 2). Despite resuscitation, 50% of animals
318 had died by 3h post-haemorrhage. (Figure 2H).

319 **Myocardial inflammation and structural damaged following THS impacts on** 320 **cardiomyocyte survival**

321 From our in situ IHC-analysis in the myocardium, a significantly higher % of neutrophils
322 (Ly6G stained cells) was identified in the heart of the THS group compared to naïve animals
323 (0.15 ± 0.07 vs. 0.8 ± 0.1 % of positive LyG6 cells, respectively; $P = 0.02$) (Fig.3A); a high
324 presence of macrophages and monocytes were also identified in the THS group (1.7 ± 0.01 vs.
325 0.3 ± 0.01 % positive Iba-1 cells in THS vs naïve, respectively; $P = 0.001$). (Fig. 3B).

326 From the flow cytometry analysis carried out following ex vivo heart cells disaggregation, a
327 significantly higher number of neutrophils were identified in THS group compared to naïve
328 non-injured animals (11.7 ± 5.2 vs. 3.4 ± 2.7 % F4/80Neg, Ly6C/GHi cells; one way ANOVA
329 with Tukey's multiple Comparison $P = 0.01$) (Fig3.D). Sham animals did not differ from naïve
330 or THS (6.4 ± 3.9 % F4/80Neg, Ly6C/GHi cells). When investigating monocyte cell
331 population, we identified a statistically higher ratio of pro-inflammatory (F4/80Pos,
332 Ly6C/GInt) vs anti-inflammatory (F4/80Pos, Ly6C/GLow) monocytes in the THS group
333 compared to the naïve and sham animals (1.14 ± 0.8 for THS group vs 0.25 ± 0.1 for naïve
334 group and 0.32 ± 0.1 for sham group; $P = 0.006$ and $P = 0.01$, respectively) (Fig.3H). There was
335 also a trend to shift the presence of pro vs. anti-inflammatory monocytes in the hearts of THS
336 animals, compared to that for the naïve and sham groups; but these data did not reach the
337 threshold for statistical significance (Fig. 3F,G).

338 TEM analysis demonstrated significant morphological changes within the myocardium, with
339 interstitial oedema and widespread disorganisation of the cardiac myofibrillar ultrastructure,
340 and a relaxation of the sarcomere in the cardiomyocytes (Fig. 4A). We identified the presence
341 of poorly circumscribed mitochondria distributed in a disorderly fashion across cardiac
342 muscle fibres, with significant structural changes. In THS animals, 33% of the mitochondria
343 contained amorphous dense bodies, in comparison to 23% in the sham animals ($p = 0.006$).
344 Mitochondrial swelling, with loss of electron dense material from the matrix and breakdown
345 of the cristae and vacuolation, was also observed in THS animals (Fig. 4B). The nuclei of
346 this THS group also exhibited signs of irreversible structural changes, such as margination of
347 chromatin, associated with surrounding oedema, chromatin condensation and rarefaction of
348 the nucleoplasm, where there is loss of the chromatin (Fig.4C-E), compare to sham animals

349 (Fig4.F-H). Most of these lesions are consistent with presence of myocardial ischaemia injury
350 the THS hearts (Fig 4).

351 Myocardial damaged was confirmed by the reduced levels of H-FABP in the myocardium of
352 the THS hearts ($p=0.06$ NS; $p=0.04^*$, respectively; two-tailed unpaired t-test; means \pm SD),
353 indicating the loss /release of these proteins, attributable to myocardial injury. (Fig.4I,J).

354 A significant decrease in caspase 8 expression in THS was seen on IHC (Fig 4 K) is
355 suggestive that the cardiomyocyte damage seen is not dependent upon death receptor initiated
356 or protease dependent apoptosis. The difference in caspase 8 on WB although downward
357 trending was not statistically significant (Fig 4L). The total AIF expression was similar
358 between Sham and THS models (0.20 vs 0.17 respectively, NS; Fig 5A), but a
359 compartmentalisation of AIF outside the mitochondria was observed in the TH models. The
360 decrease in the mitochondrial AIF in comparison to cytosolic AIF in the THS hearts
361 (mitochondria vs. total cell $P=0.04$ and mitochondria vs. supernatant/cytosol $P=0.04$; Fig 5B)
362 suggests a leakage of the mitochondrial AIF into the cytosol, with the potential activation of a
363 mitochondrial driven cell death pathway (Fig.5A-D).

364 Discussion

365 Our study confirms that the preclinical modelling approach that we have developed
366 physiologically and biochemically mirrors the cardiac dysfunction seen in bleeding trauma
367 patients. Traumatic injury and controlled haemorrhage induced significant acute cardiac
368 damage despite subsequent echocardiography-guided volume resuscitation. These changes
369 were associated with demonstrable myocardial cell death and inflammation leading to
370 reduced survival.

371 In our unresuscitated model, we identified cardiac functional changes and myocardial damage
372 with comparable elevations in the cardiomyocyte damage molecule H-FABP [21] to those
373 reported in clinical studies [8, 22]. The associated functional decreases in cardiac output were
374 at least in part related to the loss of stroke volume. However, our resuscitated model, with
375 confirmed restoration of volume preload, demonstrated a persistent and progressive loss of
376 cardiac function associated with increasing myocardial injury indicated by further elevations
377 in H-FABP. Catecholamine release with increased inotropy will undoubtedly compensate for
378 some degree of cardiomyocyte loss. However in our model, volume resuscitation to baseline
379 LVEDV was not able to restore cardiac function to pre-shock levels. This resultant cardiac
380 dysfunction is increasingly recognised as a key determinant of critical care utilization and
381 survival in trauma patients [10, 23, 24].

382 Cardiac dysfunction is well known in sepsis [12, 25], but the aetiology of trauma induced
383 myocardial injury likely has different underlying pathophysiology [1-16, 26]. We observed
384 an acute cardiac inflammatory response, with an increase in monocytes/macrophages and
385 neutrophils infiltration in the heart is identified following THS injury. This may be driven by
386 the systemic inflammatory response to trauma [15, 26], or may be a direct response of the
387 immune system to cardiomyocyte cell death. The acute recruitment of monocytes and
388 neutrophils from circulation into the myocardium has also been reported in cardiac stress
389 ischemic conditions [27]. The persistence of this myocardial inflammation response may lead
390 to further endogenous cytokine production and leukocyte recruitment and infiltration,
391 increasing oxidative stress, cell damage and cardiac dysfunction [28].

392 We observed severe cardiomyocyte ultrastructural and organelle dysfunctional damage by 3h
393 post trauma, with myofibrillary disarray, relaxation of sarcomeric proteins, mitochondrial
394 vacuole formation, membrane disruption and chromatin features consistent with irreversible
395 damage. Such cellular stress features, augurs of cell death, have been described in rodent
396 cardiomyocytes following ischemia-reperfusion [29]. Furthermore, the increased
397 translocation of mitochondrial AIF into the cytosol, and then into the nucleus, confirms the
398 activation of cell death pathways associated with cell death. This presence of AIF leakage in
399 addition to the reduced expression of caspase 8, may indicate the involvement of the alternate
400 cell death pathway of necroptosis as a principle mode of cardiomyocyte cell death in trauma
401 [30, 31]. Many alarmins released following trauma (ATP, DNA, histones, HMGB-1, HSP70)
402 have been associated with necroptosis signalling [32], and such necroptosis has already been
403 identified as central to other sterile inflammation conditions such as acute pancreatitis [33]
404 and organ injury[34]. Our data raises the possibility of mitochondrial mediated necroptosis
405 triggered by specific extracellular alarmins as the underlying cardiomyocyte cell death
406 pathways, leading to cardiac dysfunction in trauma patients. However, the role of AIF and
407 necroptosis is still poorly understood in the realm of organ injury and unexamined in the
408 context of trauma. Therefore, it is pertinent to conduct further studies to modulate AIF
409 translocation and examine its effect on necroptosis, cell survival and therefore cardiac
410 dysfunction in models prior to translation to humans. Our modelling approach will support
411 further mechanistic studies on the role of inflammatory mediators in driving specific
412 tissue/organ dysfunction after trauma, particularly allowing for the use of transgenic animals
413 for inflammatory pathways.

414 There are several limitations to this study. We limited our model to 3 hours, and longer
415 durations will be required to examine the longer term impacts of injury on cardiac function.
416 The implementation of serial echocardiography to guide resuscitation is unique in trauma
417 models and delivers a new in-vivo understanding of cardiac dysfunction in trauma
418 haemorrhage. However, it was difficult to assess myocardial contractility with ultrasound
419 and further advanced imaging approaches such as speckle tracking or MRI tagging could be
420 used to explore this, as well as ex-vivo isolated heart techniques. Monitoring of other
421 clinically relevant resuscitation parameters like urine output, arterial blood gas or central
422 venous pressure could also have been explored, but their implementation in this echo-guided
423 trauma mouse model remain challenging due to low blood volumes and technical limitations.
424 Our inflammatory and biochemical analyses identified cardiomyocyte cell death suggesting,
425 but not definitive of, activation of necroptosis pathways. Further work will be required, to
426 fully characterise the cell death mechanism and its relationship with the sterile inflammatory
427 response.

428 Cardiac dysfunction is now a major mode of trauma haemorrhage death after admission. We
429 have identified the development of severe and irreversible myocardial damage, despite fluid
430 resuscitation, leading to cardiac dysfunction and death. We pose AIF-driven necroptosis as a
431 possible underlying mechanistic pathway for the cell death. Myocardial protection through
432 novel management strategies and therapeutic approaches represents a major opportunity for
433 improving survival after major trauma.

434 **Ethics Statement**

435 This study was carried out in accordance with the recommendations of Home Office
436 guidance on the operation of Animals (Scientific Procedure Act, 1986) in accordance with the
437 EU Directive 2010/63/EU and the Guide for the Care and Use of Laboratory Animals of the

438 National Research Council. The protocols were approved by the Animal Welfare and Ethics
439 Review Board of Queen Mary University of London and conducted under the UK home
440 office license number PC5F29685. Experimental planning and design was performed in
441 accordance with the ARRIVE guidelines for data randomization, blinding for results analysis
442 and sample size calculation. All studies were carried out under non-recovery terminal
443 anaesthesia. Animals never regain conscious state, been constantly monitored (MABP, HR,
444 Resp. rate, body temperature) throughout the whole study. Survival state is defined as the
445 animal's ability to maintain a MAP>15mmHg with measurable respiratory and cardiac
446 function. Once any animal reaches any physiological state below the survival threshold, the
447 animals is humanely killed by exsanguination to collect terminal tissue / blood samples.

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451 members from the Centre for Trauma Sciences for their advice.

452 **Conflict of Interest Statement**

453 The authors declare that the research was conducted in the absence of any commercial or
454 financial relationships that could be construed as a potential conflict of interest.

455 **Author Contributions**

456 KB, JW, VN and JLT designed the overall study and experimental programme, together with
457 CT. JW, VN, BP, OFB, SMcA and JLT performed the experiments including animal studies,
458 cell sorting experiments, microscopic studies, and ELISAs. JW, VN, BP, JLT, and KB
459 contributed to experimental design and data analysis and coordinated the study and
460 supervised financial support for the studies. JW, VN and JLT produced initial drafts of the
461 manuscript. CT is the project license holder for the animal work carried out. All authors
462 contributed and revise the drafting of the article and gave final approval of the version to be
463 published.

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574

575 **Figure legends:**

576 **Figure 1: Characterisation of the shock status and cardiac dysfunction at 60 min post**
577 **traumatic haemorrhagic shock injury (1h post-THS).** A) Modelling strategy describing the
578 experimental set up with vessel cannulations, traumatic soft and hard tissue injury and
579 pressure-controlled haemorrhage. B) Median values, interquartile range and range of arterial
580 blood lactate levels at 60min post-THS between the experimental groups (S= sham with only
581 vessel cannulation only; T=Traumatic injury alone; TH 60-70=THS with 60-70 mmHg
582 MABP and TH 30-40=THS with 30-40 mmHg MABP). ****P<0.001** between TH 30-40 and S,
583 T and TH 60-70. C) Serum H-FABP levels as a myocardial injury biomarker at 60min post-
584 THS between the S, T and THS 60-70 and 30-40mmHg groups, showing a significant
585 difference in the THS-30-40 group vs. the others (****P<0.001**). D) Representative
586 echocardiographic images across the longitudinal heart axis showing the collapsed LV in
587 THS animals at 60 min post-injury, in comparison with baseline non-no injured heart; yellow
588 line depicts the endocardium; red line depicts epicardium. E) Cardiac outcome (CO; ml/min)
589 at 60 min post-injury, showing the impact of pre-load effects on the THS 60-70 and the THS
590 30-40 groups compared to S and T groups (*P<0.05, **P<0.001 between groups; **P<0.01
591 TH 60-70 versus TH 30-40). D) Stroke volume (SV; uL) at 60min post-injury, showing the
592 impact of pre-load effects on the THS 60-70 and the THS 30-40 groups (*P<0.05, **P<0.001
593 between groups).

594 **Figure 2: Characterisation of the 3h THS cardiac dysfunction (CD) model .** A)
595 Modelling strategy: mice subjected to trauma haemorrhage are held at a MAP of 30-40mmHg
596 for an hour and then resuscitated with shed blood & crystalloid (Resus1) to their original
597 stroke volume (SV). Mice are then left for a further hour and resuscitated again (Resus 2) and
598 then left to complete a 3-hour experiment. Animals undergo two echocardiographic-guided
599 resuscitations at 60 minutes and 120 minutes to baseline SV. B) Progressive decrease of
600 MABP despite pre-load resuscitation fluids. C) Serum lactate levels increase steadily after
601 the 1st resus (3.93 ±0.93mmol/L) to T180 post-THS (7.9 ±5.19mmoL). D) Left ventricular
602 end-diastolic volume (LVEDV) does not regain baseline levels despite resuscitation pre-loads
603 (Baseline: 76.17 ±10.15 vs. 180 min 16.40 ±12.98 at T180 post-THS; P<0.0001 one-way
604 ANOVA). E) Cardiac Output (CO; ml/min) and D) SV (uL) indicate the progressive loss of

605 LV function despite pre-load resuscitation. This is observed already following the 1st resus
606 after T60 THS insult, despite temporary increases in CO and SV immediately following
607 resuscitations (at T60 and T120; see Table 4). G) Following the initial dosing of shed blood
608 (SB) during the 1st resus at T60min, the crystalloid fluid require to reach baseline SV is
609 steadily increased during the 1st and 2nd resuscitation (THS60 min and THs120min,
610 respectively). E) Impact of CD in the median survival, with 50% decrease survival at
611 THS180min.

612 **Figure 3: Characterisation of acute cardiac inflammation response in the Trauma**
613 **Haemorrhage induced CD murine model (THS).** Immunohistochemistry (IHC)
614 myocardial analysis shows A) a significant increase in neutrophils (% LyG6 cells) and B)
615 Monocytes/ Macrophages (% Iba-1 cells) in the THS model (T180 min) compared to naïve
616 ($p \leq 0.001^{**}$, $p=0.02^{*}$; two-tailed unpaired t-test). Data is presented as means \pm SD. C)
617 Representative IHC microphotographs in heart section in a THA and naïve animal (x400) and
618 zoom insets of the LY6G and Iba-1 staining (red) for the neutrophils and
619 monocyte/macrophage, respectively, with the nuclei Hoechst staining (blue) in a THS model.
620 D-H) Flow cytometry assessment of heart-infiltrating leukocytes. D) Representative forward
621 scatter/side scatter profile and a live cell gate. Following a CD45/side scatter plot to obtain
622 the CD45⁺ leukocyte2, Ly6G Hi or low expression with F4/80 plots was used to derived
623 percentages of pro-inflammatory monocytes (Ly6G^{Hi}, F4/80⁺: M1 phenotype) and anti-
624 inflammatory monocytes/macrophages (Ly6G^{Low}, F4/80⁺: M2 phenotype). The heart from
625 THS animals showed a significant increase in the M1:M2 phenotype ratio, compared to the
626 naïve animals. E) A significant presence of neutrophils (% Ly6G⁺ F4/80⁻ cells) and a high
627 ratio of inflammatory monocytes (F, G, H; Higher ratio of Ly6G^{Hi}, F4/80⁺ inflammatory
628 monocytes vs. F4/80⁺: M2 anti-inflammatory macrophages/monocytes; $p \leq 0.001^{**}$, $p=0.02^{*}$;
629 one-way ANOVA with post-hoc analysis). Data all presented as means \pm SD.

630 **Figure 4: Myocardial injury in the trauma haemorrhage induced CD murine model.**
631 TEM shows severe myocardial injury in the heart of THS mice, with interstitial oedema,
632 widespread disorganisation of the myocardium with relaxation of the sarcomere and poorly
633 circumscribed mitochondria distributed in a disorderly fashion (A). Significant amounts of
634 mitochondrial oedema (B) Nuclei shows margination of chromatin, rarefaction of the
635 nucleoplasm (C) and glycogen depletion (D); and some cardiomyocytes dead cells with
636 margination of chromatin (E). TEM of sham mice heart tissue shows a well-organised
637 myofibrils, with mitochondria evenly organised along the cristae sarcomere (F) with intact
638 nucleus (G) and mitochondria (H). WB analysis of Troponin-I (cTnI) (I) and H-FABP (J)
639 expression shows elevated levels in the myocardium of THS mice ($p=0.06$ NS; $p=0.04^{*}$,
640 respectively; two-tailed unpaired t-test; means \pm SD). K, L) Low caspase 8 expression is
641 observed in heart of THS mice (; IHC $p < 0.0001$; WB $p > 0.05$ compared to sham animals).

642 **Figure 5: AIF localisation in the trauma haemorrhage induced CD murine model.** A)
643 AIF expression as demonstrated by western blot analysis. The total AIF expression is similar
644 in the Sham and THS models (0.20 vs 0.17, respectively). B) Cellular compartmentalisation
645 of AIF as demonstrated by WB analysis. Higher proportion of AIF is found outside the
646 mitochondria in the THS models The mean ratio of mitochondrial AIF (mAIF) to supernatant
647 AIF (sAIF) is 1.5 in sham compared to 0.4 in THS group, $p=0.16$. When comparing total AIF
648 which includes the nucleus, a lower proportion is still found in the mitochondria. Ratio of
649 mitochondrial AIF (mAIF) to total AIF (tAIF) in sham group is 0.6 compared to 0.3 in THS
650 animals, $p=0.04$. This is suggestive of its translocation of the AIF out of the mitochondrion,
651 potentially to the nucleus, triggering initiation of cells death. IHC sections from TH animals,

652 DAPI (blue), AIF (green), MTCO2 (red) and combined images. In the combined images,
653 localisation of AIF in the nucleus can be seen in the THS (D) which is absent in the sham
654 animals (C).

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656