

1 **TITLE:**

2 Reliability of Gastrointestinal Barrier Integrity and Microbial Translocation Biomarkers at
3 Rest and Following Exertional Heat Stress

4 **AUTHORS:**

5 Henry B. Ogden ¹, Joanne L. Fallowfield ², Robert B. Child ³, Glen Davison ⁴, Simon C.
6 Fleming ⁵, Robert M. Edinburgh ⁶, Simon K. Delves ², Alison Millyard ¹, Caroline S.
7 Westwood ¹. and Joseph D. Layden ¹

8 **AUTHOR AFFILIATION:**

9 ¹ School of Sport, Health and Wellbeing, Plymouth MARJON University, Plymouth, United
10 Kingdom

11 ² Institute of Naval Medicine, Alverstoke, United Kingdom

12 ³ School of Chemical Engineering, University of Birmingham, Birmingham, United Kingdom

13 ⁴ Endurance Research Group, School of Sport and Exercise Sciences, University of Kent,
14 Chatham Maritime, United Kingdom

15 ⁵ Royal Cornwall NHS trust, Truro, United Kingdom

16 ⁶ Department of Health, University of Bath, Bath, United Kingdom

17 **RUNNING TITLE:**

18 Reliability of Gut Integrity Biomarkers Around Exertional Heat Stress

19 **KEYWORDS:**

20 Gut, Exercise, Endotoxin

21 **TOTAL WORDS:**

22 8352 (Main Body, Legends, References)

23 **TOTAL REFERENCES:**

24 68

25

26 **CORRESPONDING AUTHOR:**

27 Henry B. Ogden

28 Plymouth MARJON University

29 Faculty of Sport, Health and Wellbeing,

30 Derriford Rd, Plymouth, PL6 8BH, United Kingdom

31 Telephone: 0791 454 0094

32 Email: ogden.h@pgr.marjon.ac.uk

33 **SUBJECT AREA**

34 Environmental and Exercise Physiology

35 **NEW FINDINGS:**

36 *What is the central question(s) of this study?*

37 To assess the reliability of gastrointestinal barrier integrity and microbial translocation
38 biomarkers both at rest and in response to exertional-heat stress.

39 *What is the main finding and its importance?*

40 Acceptable and defined-levels of reliability are presented for the serum Dual-Sugar
41 Absorption (lactulose/L-rhamnose) Test, Intestinal Fatty-Acid Binding Protein, Claudin-3,
42 Lipopolysaccharide Binding Protein, total 16s DNA, but not the *Bacteroides*/total 16s DNA
43 ratio at both measurement time-points.

44

45 **ABSTRACT**

46 **Purpose:** Exertional-heat stress adversely disrupts (GI) barrier integrity and, through
47 subsequent microbial translocation (MT), negatively impacts health. Despite widespread
48 application, the temporal reliability of popular GI barrier integrity and MT biomarkers is poorly
49 characterised. **Method:** Fourteen males completed two 80-minute exertional-heat stress
50 tests (EHST) separated by 7-14 days. Venous blood was drawn pre, immediately- and 1-hour
51 post both EHSTs. GI barrier integrity was assessed using the serum Dual-Sugar Absorption
52 Test (DSAT), Intestinal Fatty-Acid Binding Protein (I-FABP) and Claudin-3 (CLDN-3). MT was
53 assessed using plasma Lipopolysaccharide Binding Protein (LBP), total 16S bacterial DNA and
54 *Bacteroides* DNA. **Results:** No GI barrier integrity or MT biomarker, except absolute
55 *Bacteroides* concentration, displayed systematic trial order bias ($p \geq 0.05$). I-FABP (trial 1 = Δ
56 $0.834 \pm 0.445 \text{ ng}\cdot\text{ml}^{-1}$; trial 2 = Δ $0.776 \pm 0.489 \text{ ng}\cdot\text{ml}^{-1}$) and CLDN-3 (trial 1 = Δ 0.317 ± 0.586
57 $\text{ng}\cdot\text{ml}^{-1}$; trial 2 = Δ $0.371 \pm 0.508 \text{ ng}\cdot\text{ml}^{-1}$) were increased post-EHST ($p \leq 0.01$). All MT
58 biomarkers were unchanged post-EHST. Coefficient of variation and typical error of
59 measurement post-EHST were: 11.5% and 0.004 (ratio) for the DSAT 90-minutes post probe
60 ingestion; 12.2% and 0.004 (ratio) at 150-minutes post probe ingestion; 12.1% and 0.376
61 $\text{ng}\cdot\text{ml}^{-1}$ for I-FABP; 4.9% and $0.342 \text{ ng}\cdot\text{ml}^{-1}$ for CLDN-3; 9.2% and $0.420 \mu\text{g}\cdot\text{ml}^{-1}$ for LBP; 9.5%
62 and $0.15 \text{ pg}\cdot\mu\text{l}^{-1}$ for total 16S DNA; and 54.7% and 0.032 for *Bacteroides*/total 16S DNA ratio.
63 **Conclusion:** Each GI barrier integrity and MT translocation biomarker, except
64 *Bacteroides*/total 16S ratio, had acceptable reliability at rest and post exertional-heat stress.

65

66

67 **ABBREVIATIONS**

68	ANOVA	Analysis of variance
69	B-A	Bland-Altman Limits of Agreement
70	Bact.	<i>Bacteroides</i>
71	BactDNA	Bacterial DNA
72	CLDN-3	Claudin-3
73	CV	Coefficient of Variation
74	DSAT	Dual Sugar Absorption Test
75	EDTA	Ethylenediaminetetraacetic acid
76	EHST	Exertional Heat Stress Test
77	ELISA	Enzyme Linked Immunosorbent Assay
78	GI	Gastrointestinal
79	HPLC	High Performance Liquid Chromatography
80	HR	Heart Rate
81	I-FABP	Intestinal Fatty-Acid Binding Protein
82	ISAK	International Society for the Advancement of Anthropometric
83		Kinanthropometry
84	LBP	Lipopolysaccharide Binding Protein
85	L/R	Lactulose-to-Rhamnose
86	MT	Microbial Translocation
87	PCR	Polymerase Chain Reaction
88	RH	Relative Humidity
89	RPE	Rate of Perceived Exertion
90	SD	Standard Deviation
91	T _{core}	Core Body Temperature
92	T _{body}	Mean Body Temperature
93	TEM	Typical Error of Measurement
94	T _{skin}	Mean Skin Temperature
95	TS	Thermal Sensation
96	$\dot{V}O_{2max}$	Maximal Oxygen Uptake

97

98 **INTRODUCTION**

99 The gastrointestinal (GI) microbiota is a complex microbial ecosystem, which performs
100 numerous functions symbiotic to human health (Cani, 2018). However, to prevent immune
101 activation the microbiota must remain contained within the GI lumen, a process that is tightly
102 regulated by the multi-layered GI barrier (Wells et al., 2017). Exertional heat stress is one
103 stimulus that adversely disrupts GI barrier integrity, and in a linear manner to the severity of
104 splanchnic hypoperfusion (van Wijck et al., 2011) and core body temperature (Pires et al.,
105 2017). In severe cases, luminal microbial products are capable of transversion into the
106 systemic circulation, a response now considered to underlie multiple common athletic health
107 conditions (Costa et al., 2017). Specifically, the most concerning of these health conditions
108 include exercise-induced anaphylaxis (Christensen et al., 2019) and exertional heatstroke
109 (Lim, 2018). In non-exercise settings, research presently links GI microbial translocation (MT)
110 within the pathophysiology of numerous chronic illnesses, including GI disease (Camilleri et
111 al., 2012), cardiovascular disease (Neves et al., 2013) and degenerative disorders of the
112 central nervous system (Mulak and Bonaz, 2015). Thus, reliable biomarkers of GI barrier
113 integrity and/or MT appear important in the surveillance, diagnosis and treatment of these
114 conditions. To date, there is little evidence documenting the reliability of most commonplace
115 assessment biomarkers, which limits interpretation of their application in both laboratory and
116 field settings.

117 GI barrier integrity can be assessed *in vivo* using several biomarkers of intestinal
118 permeability, epithelial injury and tight junction integrity (Wells et al., 2017). The Dual-Sugar
119 Absorption Test (DSAT) is the gold-standard GI permeability technique (Bischoff et al., 2014).
120 The traditional endpoint of the DSAT is the 5-hour urinary recovery of pre-ingested lactulose-
121 to-L-rhamnose (L/R; Bischoff et al., 2014) and offers good test-retest reliability when applied
122 at rest (Marchbank et al., 2011). Analytical improvements have recently validated a serum
123 DSAT over a reduced (i.e. 1-3 hours) time course (van Wijck et al., 2013) and with improved
124 diagnostic sensitivity (JanssenDuijghuijsen et al., 2016; Pugh et al., 2017a). However, given
125 the transient appearance of sugar probes within the blood (Fleming et al, 1996), potentially
126 due to the wide heterogeneity in gastric emptying rates following exercise (Costa et al., 2017),
127 the reliability of this technique requires verification. Intestinal Fatty Acid-Binding Protein (I-

128 FABP) is a cytosolic protein expressed exclusively within enterocytes of the
129 duodenum/jejunum, and has a half-life of 11 minutes in the systemic circulation following
130 epithelial injury (van de Poll et al. 2007). These characteristics have popularised I-FABP as a
131 prominent biomarker of small GI epithelial injury (Wells et al., 2017), with serum
132 concentrations strongly predictive of small GI histological injury (Schellekens et al., 2014). The
133 temporal reliability of I-FABP has never been directly assessed and requires interrogation
134 given its high sensitivity to sub-clinical small GI injury. Claudin-3 (CLDN-3) is a conserved GI
135 epithelial transmembrane protein, which performs an integral role in GI paracellular
136 homeostasis (Zeissig et al., 2007). As a biomarker of GI tight junction (TJ) integrity, preliminary
137 research has shown a strong relationship between urinary CLDN-3 concentration and
138 histological GI CLDN-3 breakdown (Thuijls et al., 2010a, 2010b). Similar to I-FABP, the
139 temporal reliability of plasma CLDN-3 is currently unknown.

140 GI MT can be assessed *in vivo* through several indirect biomarkers considered to be
141 indicative of systemic microbial exposure (Wells et al., 2017). Endotoxin, a form of
142 lipopolysaccharide located on the outer membrane of gram-negative bacteria, has
143 traditionally been utilised for this purpose (Costa et al., 2017). However, the search for
144 improved GI MT biomarkers is ongoing, given endotoxin analysis is susceptible to both false-
145 positive (e.g. from exogenous contamination) and false-negative (e.g. from rapid hepatic
146 clearance) results (Dullah and Ongkudon, 2017). Lipopolysaccharide binding protein (LBP) is
147 a type-1 acute phase protein, secreted hepatically following systemic exposure to numerous
148 microbial-associated molecular patterns (Schumann, 2011). However, as an acute-phase
149 protein, its temporal reliability is likely highly subject to influence from numerous co-variables
150 (e.g. infection) (Citronberg et al., 2016). Bacterial DNA (bactDNA), through conserved 16S
151 gene sequencing, is an emerging biomarker of GI MT (Paisse et al., 2016). In comparison with
152 alternative MT measures, one major advantage of bactDNA is an apparent independence of
153 hepatic clearance (Mortensen et al., 2013). One innovative study recently proposed a
154 bactDNA methodology aimed to improve analytical specificity and reliability through
155 targeting a predominant GI bacterial genus (*Bacteroides*) and correcting for total 16S DNA
156 concentration (March et al., 2019).

158 The aim of the present study was to determine the reliability of biomarkers of GI
159 barrier integrity (DSAT, I-FABP, CLDN-3) and microbial translocation (LBP, total 16s bacterial
160 DNA, *Bacteroides* DNA) at rest and following exertional-heat stress. These data should inform
161 prospective study design, including biomarker selection and statistical power.

162 **METHODS**

163 **Participants and Ethical Approval**

164 Fourteen healthy males (Table 1) volunteered to participate in the present study. All
165 participants were non-smokers, habitually active, non-endurance trained (>4 h·week⁻¹) and
166 unacclimated to hot environments. A general medical questionnaire was used to screen for
167 previous histories of gastrointestinal, cardiorespiratory and metabolic illnesses. No
168 participant took pharmacological medications (e.g. laxatives, antibiotics) or reported
169 suffering from an acute illness within 14 days prior to data collection. Informed consent was
170 obtained for each participant following a full written and oral explanation of the experimental
171 procedures. The study protocol was approved by Plymouth MARJON University Research
172 Ethics Committee (Approval Code: EP040) and was conducted in accordance with the
173 principles outlined in the Declaration of Helsinki, except for trial registration within a
174 database.

175 [Table 1 – Insert Here]

176 **Experimental Overview**

177 Participants visited the laboratory on three occasions. During the first visit, baseline
178 anthropometrics and maximal oxygen uptake ($\dot{V}O_{2max}$) were assessed. The second and third
179 visits were the main experimental trials. These were separated by 7-14 days to negate the
180 influence of prior exertional-heat stress on thermoregulatory (Barnett and Maughan, 1993)
181 and GI barrier integrity (Snipe et al., 2017) responses. During both main experimental trials,
182 participants completed an intermittent exertional-heat stress test (EHST), consisting of two
183 bouts of 40 minutes fixed-intensity treadmill walking (6 km·h⁻¹ and 7% gradient) in the heat
184 (35°C and 30% relative humidity; RH). The exercise bouts were separated by 20-minutes
185 seated recovery, including 4-minutes forearm cold water immersion. This protocol is
186 consistent with general military guidance on work/rest schedules for sustained physical

187 activity in the heat (Military Headquarters of the Surgeon General, 2017) and unpublished
188 pilot data from our laboratory showing a ~ 2-fold elevation in DSAT responses relative to rest
189 ($n= 6$; DSAT 90-minute post probe ingestion; [rest] = 0.014 ± 0.006 , [post EHST] = $0.028 \pm$
190 0.005 ; $p = 0.02$). Data collection coincided with non-summer months in Plymouth, United
191 Kingdom, where daily mean ambient temperature at a local meteorological station
192 (Camborne, United Kingdom; latitude: 50.218° N) remained below 20°C (Met Office, 2019).
193 A schematic illustration of the protocol is shown in Figure 1.

194

195 [Figure 1 – Insert Here]

196

197 **Dietary and Lifestyle Controls**

198 Dietary supplementation (e.g. glutamine, probiotics, bovine colostrum) and prolonged
199 thermal exposures (e.g. saunas, sunbeds) were prohibited from 14 days before until the end
200 of data collection (Costa et al., 2017). Alcohol, caffeine, strenuous physical activity and non-
201 steroidal anti-inflammatory drugs (e.g. ibuprofen) were all abstained for 48 hours before main
202 experimental visits (Costa et al., 2017). Participants adhered to a ≥ 10 hour overnight fast and
203 consumed 500 ml of plain water two hours prior to main experimental visits. Conformity with
204 all pre-trial controls was assessed in writing upon laboratory arrival using a pre-trial control
205 questionnaire. Participants remained fasted throughout all main experimental trials
206 (Edinburgh et al., 2018), but were permitted a $12 \text{ ml}\cdot\text{kg}^{-1}$ bolus of ambient temperature water
207 ($28\text{-}30^\circ\text{C}$) to drink over 20 minutes following both 40-minute EHST bouts.

208 **Anthropometric Measurements**

209 Participants height, weight and body fat were measured following ISAK guidelines
210 (Marfell-Jones et al. 2006). Height was measured barefoot using a stadiometer to the nearest
211 0.1 cm (Marsden HM-200, Rotherham, UK), whilst body mass was measured on an electronic
212 scale to the nearest 0.05 kg (Tanita MC 180 MA, Tokyo, Japan). Skinfold thicknesses were
213 taken in duplicate by the same researcher at the bicep, tricep, subscapular and suprailliac
214 using skinfold callipers to the nearest 0.1 cm (Harpندن, Holtain Ltd, Crymych, UK).
215 Predictions of body density were calculated using age and gender related equations (Durnin

216 and Womersley, 1974).

217 **Maximal Oxygen Uptake**

218 Maximal oxygen uptake ($\dot{V}O_{2max}$) was determined using an incremental treadmill test
219 (Desmo HP, Woodway GmbH, Weil am Rhein, Germany) to volitional exhaustion. The test was
220 undertaken in normothermic laboratory conditions (18-22°C, 40-60% RH). The test began at
221 a speed of 10 km·h⁻¹ on a fixed 1% inclination. The treadmill speed was then increased at 1
222 km·h⁻¹ increments every three minutes until reaching 13 km·h⁻¹, when inclination was then
223 increased by 2% every two minutes. Expired metabolic gases were measured continuously
224 using a breath-by-breath metabolic cart (Metalyser 3B, Cortex, Leipzig, Germany). Heart rate
225 (HR; Polar FT1, Polar Electro OY, Kempele, Finland) and rating of perceived exertion (RPE;
226 Borg, 1970) were measuring during the final ten seconds of each stage. The highest 30 second
227 average $\dot{V}O_2$ was taken to be $\dot{V}O_{2max}$.

228 **Exertional-Heat Stress Test**

229 EHSTs commenced in the morning (08:30 ± 1 hour) to avoid the influence of circadian
230 variation (Waterhouse et al., 2005). Upon laboratory arrival, participants provided a capillary
231 blood sample into a K₂EDTA microtube (Microvette®, Sarstedt, Numbrecht, Germany) for
232 duplicate hydration assessment via plasma osmolality using freeze-point depression
233 (Osmomat 3000, Gonotec, Berlin, Germany). Participants then measured their own nude
234 body mass (Tanita MC 180 MA, Tokyo, Japan). They then self-inserted a single use rectal
235 thermistor (T_{core} ; Phillips 21090A, Guildford, UK) 12 cm beyond the anal sphincter and a HR
236 monitor was positioned around their chest (EQ02, Equivital™, Cambridge UK). Next they
237 dressed in standard summer military clothing (i.e. jacket [zipped, sleeves extended], trousers,
238 boxer briefs, socks, trainers) and entered the environmental chamber that was regulated at
239 ~35°C (Trial 1: 35.2 ± 0.3°C; Trial 2: 35.4 ± 0.4°C; $p= 0.15$) and ~30% RH (Trial 1: 28 ± 4%; Trial
240 2: 28 ± 2%; $p= 0.25$). Skin thermistors (EUS-UU-VL3-O, Grant Instruments, Cambridge, UK)
241 were then affixed on the participant using one layer (5 x 5 cm) of cotton tape (KT Tape®, KT
242 Health, UT, USA) and mean skin temperature (T_{skin}) was calculated using standard equations
243 (Ramanathan, 1964).

244 Participants then undertook the pre-defined EHST. Throughout, T_{core} and T_{skin} were

245 recorded using a temperature logger (Squirrel SQ2010, Grant Instruments, Cambridge, UK)
246 and HR using a Sensor Electronics Module (SEM) unit (EQ02, Equivital™, Cambridge UK). Mean
247 whole body temperature (T_{body}) was calculated from simultaneous T_{core} and T_{skin}
248 measurements (Jay and Kenny, 2007). All data, including RPE (Borg et al., 1970) and thermal
249 sensation (TS; Toner et al., 1986) were reported at 20 minute intervals. Between the two
250 walking bouts, participants immersed their forearms in a $\sim 15^{\circ}\text{C}$ cold-water bath (Trial 1: 15.4
251 $\pm 0.8^{\circ}\text{C}$, Trial 2: $15.3 \pm 0.7^{\circ}\text{C}$; $p = 0.39$). Upon EHST termination, participants were removed
252 from heat and their post-EHST nude body mass was recorded. Absolute sweat losses were
253 calculated from the change in dry nude body mass from pre-to-post EHST after correction for
254 fluid intake and blood withdrawal.

255 **Blood Collection and Analysis**

256 Venous blood samples (12 ml) were drawn immediately pre, post and one-hour post
257 EHST. Participants stood upright for a minimum of 20 minutes before collection to allow
258 capillary filtration pressure to stabilise (Shirreffs and Maughan, 1994). Blood was drawn from
259 a forearm antecubital vein under minimal stasis (<30 seconds). Samples were collected
260 proportionally into serum-separator (SST II) and K_2 EDTA tubes (Becton Dickinson and
261 Company, Plymouth, UK). The SST II tube was allowed to clot for 30-40 minutes at room
262 temperature. A 0.5ml aliquot of K_2 EDTA blood was removed for immediate haematological
263 analysis. Samples were centrifuged at $1300g$ for 15 minutes at 4°C to separate serum and
264 plasma. Aliquots were frozen at -80°C until analyses. All blood handling was performed with
265 sterile (pyrogen, DNA free) pipette tips and microtubes.

266 **Haematology**

267 Haemoglobin was measured in duplicate using a portable photometric analyser
268 (Hemocue® Hb 201+, EFK Diagnostics, Madeburg, Germany; Duplicate) and haematocrit in
269 duplicate using the microcapillary technique following centrifugation at $14,000g$ for 4 minutes
270 at room temperature (Haematospin 1400, Hawksley and Sons Ltd, Lancing, England). Plasma
271 volume was estimated using standard equations (Dill and Costill, 1974). Post-exercise analyte
272 concentrations were left uncorrected for acute plasma volume shifts, given the similarity of
273 responses between trials and the low molecular weights of quantified analytes.

274 **Dual-Sugar Absorption Test**

275 Participants orally ingested a standard sugar probe solution containing 5 g Lactulose
276 (Lactulose Oral Solution, Sandoz, Holzkirchen, Germany) and 2 g L-Rhamnose (L-rhamnose FG,
277 99% pure, Sigma Aldrich, Missouri, USA) dissolved within 50 ml of plain water (osmolality =
278 $\sim 750 \text{ mOsm}\cdot\text{kg}^{-1}$) ten minutes into the EHST. Probe concentrations were determined from
279 serum samples collected immediately pre, 90 minutes (i.e. post-EHST) and 150 minutes (i.e.
280 1-hour post-EHST) post probe ingestion following a previously described high performance
281 liquid chromatography protocol (Fleming et al., 1996). The recovery of both sugars was
282 determined per litre serum ($\text{mg}\cdot\text{l}^{-1}$). Calculation of the L/R ratio was made corrected relative
283 (%) to the concentration of sugar consumed. The limit of detection was $0.1 \text{ mg}\cdot\text{l}^{-1}$ and the
284 laboratory reference coefficient of variation was 1.8-8.5% for both probes (Fleming et al.,
285 1996).

286 **Enzyme Linked Immunosorbent Assays**

287 I-FABP ([1:2 serum dilution]; ELH-FABP2, Raybiotech[®], Norcross, USA), CLDN-3
288 ([undiluted plasma]; EH1342, Wuhan Fine Biotech, Wuhan, China) and LBP ([1:250 plasma
289 dilution]; RK01764, ABclonal, Wuburn, USA) were measured in duplicate immediately pre
290 and post EHST using a solid-phase sandwich ELISA. Optical density was measured at 450 nm
291 using a microplate reader and sample concentrations were determined from a logarithmic
292 standard curve. The intra-assay coefficients of variation were 5.0% (I-FABP), 1.5% (CLDN-3)
293 and 2.6% (LBP).

294 **Quantitative Real-Time Polymerase Chain Reaction**

295 BactDNA was measured in duplicate plasma samples collected immediately pre- and
296 post EHST using a quantitative real-time polymerase chain reaction assay (qPCR) on a
297 LightCycler 96 instrument (LightCycler 96, Roche, Basel, Switzerland). Cell free DNA was
298 isolated from plasma using a Quick-DNA Mini Prep Plus kit (D4068, Zymo Research, Irvine, CA,
299 USA) following manufacturer's instructions. Total 16S bacterial DNA was quantified according
300 to March et al. (2019) using a universal library probe (ULP, Roche, Basel, Switzerland), with
301 standards (E2006-2, Zymo Research, Irvine, CA, USA) and primers (Eurogentec, Liège,
302 Belgium) specific to a 16S region (limit of detection $0.1 \text{ pg}\cdot\text{ul}^{-1}$). *Bacteroides* species DNA
303 (*Bact.* DNA) were quantified using a double-dye probe/primer kit (Path-Bacteroides-spp,

304 Genesig, Primerdesign Ltd, Chandler's Ford, UK). Negative controls (PCR grade water) for the
305 entire extraction process were below the limit of detection for both measures. Ratio data are
306 presented as *Bacteroides*/total bacterial DNA (*Bact./16S*). The intra-assay coefficients of
307 variation were 6.3% (total 16S) and 17.5% (*Bacteroides*).

308 **Statistics**

309 All statistical analyses were performed using Prism Graphpad software (Prism V.8, La
310 Jolla, California, USA). Comparisons were made after determining normal distribution using a
311 Shapiro-Wilk test ($p \geq 0.05$). A two-way analysis of variance (ANOVA) with repeated measures
312 (time x trial) was used to identify differences between the two trials for whole-body
313 physiological, GI barrier integrity and MT data. If Mauchly's test for sphericity was violated,
314 Greenhouse Geiser corrections were applied for epsilon < 0.75 , while the Huynh-Feldt
315 correction was used for less severe asphericity. When there was only a single comparison, a
316 paired t-test or non-parametric Wilcoxon signed-ranks test was used to determine between-
317 trial differences. Statistical significance was accepted at the alpha level of $p \leq 0.05$. Data are
318 presented as mean \pm standard deviation (SD).

319 A composite *a priori* battery of statistical tests was conducted to determine inter-trial
320 reliability (Atkinson and Nevill, 1998). The DSAT was compared at each 90- and 150-minutes
321 following sugar-probe ingestion, whilst each GI biomarker was compared at rest, post EHST
322 and the delta (Δ). Systematic bias was assessed using a paired t-test or non-parametric
323 Wilcoxon signed-ranks test. Meaningful differences were evaluated using Cohen's *d* (Lakens,
324 2013). Effect sizes were categorised as trivial (≤ 0.19), small (0.20-0.49), medium (0.50-0.79)
325 and large (≥ 0.8). Relative reliability was assessed using a Pearson's product-moment
326 correlation coefficient or non-parametric Spearman's rank correlation coefficient.
327 Correlations were classified as small (≤ 0.69), moderate (0.70-0.89) and high (≥ 0.90) (Vincent
328 and Weir, 1995). Absolute reliability was assessed using each the: coefficient of variation
329 ($[SD/mean]*100$), typical error of the measurement (TEM; SD of difference between
330 scores/ $\sqrt{2}$) and Bland-Altman (B-A) plots with mean difference (bias) and 95 % Limits of
331 Agreement (LoA; Bland and Altman, 1986). CVs were classified as very good ($\leq 10\%$) and
332 acceptable ($\leq 20\%$). Relationships between biomarkers were compared using a Pearson's
333 product-moment correlation coefficient or non-parametric Spearman's rank correlation

334 coefficient. Heteroscedasticity was examined from the non-parametric correlational
335 coefficient between absolute differences and individual means presented on B-A plots.
336 Outliers were defined as ± 2.4 SD units (normally distributed) or ± 4.0 SD units (non-normally
337 distributed) outside of the mean and were removed from subsequent analysis (Aguinis et al.,
338 2013).

339 **Power Analysis**

340 Given the novelty of the dependent variables being evaluated and statistical approach
341 to undertake a battery of reliability statistical tests, it was determined infeasible to perform
342 an *a priori* sample size calculation. Instead, general guidance on appropriate sample sizes (n
343 = 12) for pilot studies were followed, whilst accounting for a ~20% anticipated participant
344 drop-out rate (Julious, 2005).

345

346 **RESULTS**

347 **Thermoregulatory, Cardiovascular and Perceptual Strain**

348 T_{core} (Figure 2A; time x trial $p = 0.63$), T_{skin} (Figure 2B; time x trial $p = 0.13$) and T_{body}
349 (Figure 2C; time x trial $p = 0.43$) all increased over time to a similar extent between trial one
350 and two. The reliability of peak, mean and Δ in T_{core} , T_{skin} and T_{body} were all good (Table 2).
351 Pre-trial plasma osmolality (trial one: 293 ± 7 mOsmol \cdot kg $^{-1}$, trial two: 294 ± 7 mOsmol \cdot kg $^{-1}$; p
352 = 0.67), Δ plasma volume (trial 1: $-0.61 \pm 5.15\%$, trial 2: $-0.02 \pm 3.69\%$; $p = 0.67$), mean sweat
353 rate (trial 1: 1.53 ± 0.38 , trial 2: 1.56 ± 0.45 l \cdot h $^{-1}$; $p = 0.61$) and percentage body mass loss (trial
354 1: 1.15 ± 0.48 ; trial 2: $1.21 \pm 0.52\%$; $p = 0.31$) were all similar between trial one and two. HR
355 (Figure 2D; time x trial $p = 0.11$), RPE (Figure 2E; time x trial $p = 0.38$) and TS (Figure 2F; time
356 x trial $p = 0.56$) all increased over time to a similar extent between trial one and two. The
357 reliability of peak, mean and Δ HR, RPE and TS were all good (Table 2).

358

359 [Figure 2 – Insert Here]

360

361 [Table 2 – Insert Here]

362

363

364 **Dual-Sugar Absorption Test**

365 Lactulose and L-rhamnose were both undetectable in all participants' basal sample
366 prior to probe ingestion. Inter-trial DSAT responses displayed no systematic bias between
367 trials at both 90- (Figure 3A) and 150-minutes (Figure 3C). There was moderate relative
368 reliability and acceptable absolute reliability at both the 90- and 150-minute time-points. B-A
369 plots displayed bias for both the 90- (Figure 3B) and 150-minute (Figure 3D) time-points.
370 Individual Lactulose and L-rhamnose concentrations had worse reliability than the combined
371 L/R ratio (Table 3). Heteroscedasticity was not present for any analyses.

372 **Intestinal Fatty Acid Binding Protein**

373 I-FABP displayed no trial order systematic bias at either rest, post- or the Δ time-point
374 (Figure 3E). Following EHSTs, I-FABP was elevated above rest (trial 1: $\Delta = 0.834 \pm 0.445 \text{ ng}\cdot\text{ml}^{-1}$
375 $[56 \pm 31\%]$; trial 2: $\Delta = 0.776 \pm 0.489 \text{ ng}\cdot\text{ml}^{-1}$ $[46 \pm 26\%]$; $p \leq 0.01$; Figure 3E). At all time-
376 points, I-FABP displayed moderate relative and acceptable absolute reliability (Table 3). B-A
377 plots are presented to illustrate bias for post EHST concentrations (Figure 3F).
378 Heteroscedasticity was not present for any analyses. One participant was excluded as an
379 outlier. Two participants' I-FABP responses displayed unexplainably poor reliability both at
380 rest and post EHSTs. These data were retained given where verbal adherence to pre-trial
381 controls was verbally confirmed. However, removal of these data would have notably
382 improved the reliability of I-FABP both at rest ($r = 0.97$; $\text{CV} = 6.1\%$; $\text{TEM} = 0.200 \text{ ng}\cdot\text{ml}^{-1}$; $\text{B-A} \pm$
383 $\text{LoA} = -0.046 \pm 0.308 \text{ ng}\cdot\text{ml}^{-1}$) and post the EHST ($r = 0.97$; $\text{CV} = 7.2\%$; $\text{TEM} = 0.221 \text{ ng}\cdot\text{ml}^{-1}$; B-
384 $\text{A} \pm \text{LoA} = 0.078 \pm 0.467 \text{ ng}\cdot\text{ml}^{-1}$).

385 **Claudin-3**

386 CLDN-3 displayed no trial order systematic bias at either rest, post- or the Δ time-point
387 (Figure 3G). Following EHSTs CLDN-3 was elevated above rest (trial 1: $\Delta = 0.317 \pm 0.586 \text{ ng}\cdot\text{ml}^{-1}$
388 $[11 \pm 17\%]$; trial 2: $\Delta = 0.371 \pm 0.508 \text{ ng}\cdot\text{ml}^{-1}$ $[9 \pm 13\%]$; $p \leq 0.01$; Figure 3G). At all time-points,
389 CLDN-3 displayed high relative and very good absolute reliability (Table 3). B-A plots are

390 presented to illustrate bias for post EHST concentrations (Figure 3H). Heteroscedasticity was
391 not present for any analyses.

392

393 [Table 3 – Insert Here]

394

395 [Figure 3 – Insert Here]

396

397 **Lipopolysaccharide Binding Protein**

398 LBP displayed no trial order systematic bias at either rest, post EHSTs or the Δ time-
399 point (Figure 4A). There was no influence of the EHST on LBP concentration ($p = 0.41$). At all-
400 time-points, LBP displayed moderate relative and very good absolute reliability (Table 4). B-A
401 plots are presented to illustrate bias for post EHST concentrations (Figure 4B).
402 Heteroscedasticity was not present for any analyses.

403

404

405 **Bacterial DNA**

406 Total 16s (Figure 4C) and *Bact./16S* (Figure 4G) displayed no systematic bias at either
407 rest, post EHSTs or the Δ . *Bacteroides* concentrations (Figure 4E) were systematically lower in
408 trial 2 versus trial 1 ($p = 0.04$). At rest, total 16s displayed moderate relative and very good
409 absolute reliability, whereas *Bacteroides* displayed poor relative and absolute reliability. The
410 combined *Bact./16S* ratio subsequently showed poor relative and absolute reliability at rest
411 (Table 4). There was no influence of the EHST on either total 16s ($p = 0.39$), *Bacteroides* ($p =$
412 0.33) or *Bact./16S* ($p = 0.18$) responses. B-A plots are presented to illustrate bias for post EHST

413 concentrations (Figure 4D, 4F and 4H). Heteroscedasticity was not present for any analyses.
414 One participant was excluded from all bactDNA analysis as an outlier.

415

416 [Table 4 – Insert Here]

417 [Figure 4 – Insert Here]

418

419 **Association between Biomarkers**

420 Validation of the DSAT at the 90 and 150- minutes time-points across both trial one
421 and trial two (n=28), found responses to be systematically greater at 150- (0.034 ± 0.015)
422 compared with 90-minutes (0.027 ± 0.013 ; $p = 0.05$, ES 0.50). There was poor relative ($r =$
423 0.08) and absolute (CV = 31.8%, TEM = 0.014) reliability between the sample time-points,
424 suggestive of inter-individual variability in sugar probe kinetics. Few statistically significant
425 correlations were reported when comparing GI barrier integrity and MT biomarkers. Small
426 positive correlations were reported between absolute post EHST concentrations for: I-FABP
427 and CLDN-3 ($r = 0.41$, $p = 0.04$), LBP and total 16s DNA ($r = 0.48$, $p = 0.02$), LBP and *Bacteroides*
428 ($r = 0.38$; $p = 0.05$), *Bacteroides* and total 16s DNA ($r = 0.40$, $p = 0.04$). When displayed as pre-
429 to-post DELTA, small positive correlations were reported between: LBP and DSAT at 150
430 minutes ($r = 0.54$; $p < 0.01$).

431

432 **DISCUSSION**

433 The aim of this study was to determine the short-term (one-two weeks) temporal
434 reliability of several empirical biomarkers of GI barrier integrity (DSAT, I-FABP, CLDN-3) and
435 MT (LBP, total 16s bacterial DNA, *Bacteroides* DNA) following exertional-heat stress. The main
436 findings of this study were that the serum DSAT, I-FABP, CLDN-3, LBP and total 16s bacterial
437 DNA all displayed moderate-to-strong relative and acceptable absolute reliability between

438 repeat EHSTs. In comparison, absolute *Bacteroides* DNA and *Bact./total* 16s DNA ratio
439 displayed weak relative and unacceptable absolute reliability between repeat EHSTs.

440 The serum DSAT is a valid alternative of the traditional urine DSAT (Fleming et al, 1996;
441 van Wijck et al., 2011), which offering improved sensitivity to detect transient losses in GI
442 barrier integrity following exercise (JanssenDuijghuijsen et al., 2016; Pugh et al., 2017a).
443 Despite this, the temporal reliability of the serum DSAT has never been previously assessed.
444 Potential sources of variability with the serum DSAT might relate to both the transient time
445 course of sugar probes in the blood and low absolute lactulose concentrations that challenge
446 the detection limits of common analytical techniques (Fleming et al., 1996; van Wijck et al.,
447 2013). In this study, we show for the first time that the serum DSAT can be utilised with
448 acceptable reliability, which is comparative to that previously reported with the urine DSAT
449 over both a three-day (van Elburg et al. 1995) and two-week period (Marchbank et al., 2011).
450 The optimal time-point for blood collection with the serum DSAT is an unresolved issue that
451 concerns the methodological implementation of this measure. Herein, blood was collected at
452 both 90-minutes post probe ingestion as this provides the most valid estimate of the urine
453 DSAT in basal conditions (Fleming et al., 1996), and at 150-minutes post as this is where peak
454 responses arose following similar exercise stress (van Wijck et al., 2011). Remarkably, the
455 temporal reliability of both time-points assessed was almost identical, though given large
456 inter-individual variation in probe kinetics, the magnitude of responses at the two time-points
457 had poor validity. Together, these findings advocate the use of the serum DSAT at either 90-
458 or 150- minutes following probe ingestion (where logistically most convenient) as a reliable
459 alternative to the urine DSAT. There appears little requirement to correct for basal sugar
460 probe concentrations (pre-probe ingestion) following a ≥ 10 hour overnight fast given that all
461 participants samples were returned negative.

462 I-FABP is the principal biomarker of GI epithelial injury (Wells et al., 2017). Despite
463 growing popularity, the temporal reliability of circulating I-FABP has never been previously
464 assessed. In the present study, resting I-FABP concentrations were consistently at the upper
465 end of the general healthy reference range for studies utilising an human ELISA kit (0.1-2.0
466 ng·ml⁻¹; Treskes et al., 2017). These concentrations must be considered when evaluating the
467 absolute reliability thresholds reported herein. The rationale for large between-study

468 discrepancies in absolute I-FABP concentrations are poorly understood, though are more
469 likely attributable to analytical discrepancies (e.g. ELISA antibody, ELISA wash procedure,
470 sample storage), than participant demographic (Treskes et al., 2017). The reliability of I-FABP
471 at rest displayed moderate relative and acceptable absolute reliability. Following both EHSTs,
472 I-FABP increased by approximately 50% or $0.800 \text{ ng}\cdot\text{ml}^{-1}$. This response is comparable to
473 numerous similar duration/intensity exercise protocols, such as: 45-to-60 minutes of $\sim 70\%$
474 watt_{max} normothermic cycling (van Wijck et al., 2011, [61%, $\Delta 0.306 \text{ ng}\cdot\text{ml}^{-1}$] 2012, [61%; Δ
475 $0.179 \text{ ng}\cdot\text{ml}^{-1}$] and 20-30 minutes of $\sim 80\%$ $\text{VO}_{2\text{max}}$ running (Barberio et al., 2015 [46%, $\Delta 0.297$
476 $\text{ng}\cdot\text{ml}^{-1}$]; March et al., 2017 [72%; $\Delta 0.350 \text{ ng}\cdot\text{ml}^{-1}$]). In comparison, far greater elevations in I-
477 FABP have been shown following 90-120 minutes of moderate-intensity running performed
478 in the heat (30°C ; Morrison et al., 2014 [663%; $\Delta 0.203\text{-}0.806 \text{ ng}\cdot\text{ml}^{-1}$]; Snipe et al., 2017
479 [288%, $\Delta 0.897 \text{ ng}\cdot\text{ml}^{-1}$]; 2018 [432%, $\Delta 1.230 \text{ ng}\cdot\text{ml}^{-1}$]). Given the high sensitivity of I-FABP to
480 even minor GI injury, it is vital that known extraneous variables (e.g. prandial/hydration
481 status, prior exercise) are tightly controlled prior to investigation. Whilst participants in the
482 present study provided written conformity to all pre-trial controls, two participants' resting I-
483 FABP concentrations appeared suspect to prior GI injury in one trial, which interestingly was
484 unable to be detected by any other analyte.

485 CLDN-3 is the principle biomarker of GI TJ integrity (Wells et al., 2017). Despite
486 introduction as a TJ biomarker almost a decade ago, the biological relevance of elevated
487 circulating CLDN-3 is still poorly understood. This includes the assessment of temporal
488 reliability, which is currently unknown. In the present study, resting CLDN-3 concentrations
489 were consistent with previous evidence ($0.5\text{-}15 \text{ ng}\cdot\text{ml}^{-1}$) in healthy populations (Yeh et al.,
490 2013; Typpo et al., 2015). At rest, large inter-individual variation in CLDN-3 concentration was
491 evident, meaning that relative reliability was almost uniform. Following both EHSTs, plasma
492 CLDN-3 consistently increased by approximately 8-10%. This finding compares well to the only
493 previous exercise study, where concentrations increased directly following a one hour
494 moderate-intensity ($70\% \text{ VO}_{2\text{max}}$) run in both temperate (22°C ; $6.7 > 7.6 \text{ ng}\cdot\text{ml}^{-1}$) and hot
495 (33°C ; $6.6 > 8.2 \text{ ng}\cdot\text{ml}^{-1}$) ambient environments (Yeh et al., 2013). The clinical relevance of this
496 small, transient increase in CLDN-3 following exercise is poorly understood, though is modest
497 in comparison with the magnitude of increase (4-20 fold) shown acutely following major non-
498 abdominal surgery (Typpo et al., 2015; Habes et al., 2017). Promisingly, of all the GI barrier

499 integrity biomarkers compared, CLDN-3 displayed the strongest relative and absolute
500 reliability.

501 LBP is a type-1 acute phase protein that responds to a wide-variety of microbial-
502 associated molecular patterns and is widely considered a stable indirect biomarker of
503 bacterial endotoxin exposure (Dullah and Ongkudon, 2017). In the present study, resting LBP
504 concentrations displayed showed moderate relative and good absolute reliability. These
505 results are in support of one previous study, which found short-term (≤ 7 day) basal LBP
506 responses to display moderate relative reliability (intraclass correlation coefficient = 0.61;
507 Citronberg et al., 2016). In comparison, direct assessment of endotoxin appears to have weak
508 basal temporal reliability, with an intra-individual CV of 22% reported over a similar 7-day
509 period in basal conditions (Guy et al., 2017). Following the EHST, LBP was unchanged in both
510 trials, with concentrations offering comparable levels of reliability compared to rest. Whilst
511 the evidence is sparse regarding LBP responses to exercise, previous evidence has shown a
512 minor elevation in LBP of 10-15% immediately following a fatiguing treadmill walk (4.5 km·h⁻¹
513 ¹) in the heat (40°C; 106 minutes; Selkirk et al., 2008), and 1 hour of moderate intensity (70%
514 VO_{2max}) treadmill running (Jonvik et al., 2019). A potential explanation for these discrepant
515 findings likely relate to greater thermoregulatory/cardiovascular strain in previous studies.

516 BactDNA is an emerging GI MT biomarker, given the recent characterisation of the
517 blood microbiome and improvements in 16S PCR sensitivity (Paisse et al., 2016). In the
518 present study, resting total 16S DNA concentrations displayed moderate relative and good
519 absolute reliability. This finding is promising, given previous concerns that plasma bactDNA
520 concentrations are susceptible to background sample contamination (Glassing et al., 2016).
521 Quantification of total plasma 16S bacterial DNA in exercise settings has never been
522 previously examined, though consistent with other MT biomarkers, the present results show
523 total 16S bactDNA to be stable following moderate intensity exertional-heat stress. One
524 criticism of total 16S bactDNA assessment, particularly in exercise settings, is a lack of GI
525 specificity, with total concentrations influenced by factors including: DNase concentration
526 (Velders et al., 2014) and 16S DNA contamination from other body/blood compartments
527 (Paisse et al., 2016). To account for this error, one hypothetically improved method involves
528 targeting a highly abundant GI genus such as *Bacteroides* (~30% of GI microbiota) and

529 correcting for total 16S concentration (March et al., 2019). This method is particularly
530 favourable given that the phyla *Firmicutes* and *Bacteroidetes* comprise >90% of the GI
531 microbiome (*Bacteroides*; Huttenhower et al., 2012) and <5% of the plasma microbiome
532 (Paisse et al., 2016). Utilising this hypothesis, March *et al* (2019) reported that the plasma
533 *Bacteroides*/16S DNA ratio tended to increase (~25%; $p = 0.07$) following a one-hour
534 moderate intensity (70% VO_{2max}) run in the heat (30°C), though large inter-individual
535 variability in responses were evident. In the present study, the *Bacteroides*/16S DNA ratio was
536 unchanged following the EHST and appeared to be systematically lower post the EHST in trial
537 two (but not the Δ). This systematic bias was unexpected given the uniformity of all other
538 analytes assessed and the poor analytical reliability of this biomarker (e.g. mean duplicate CV
539 = 17.5%). It is presently unclear whether the poor reliability of this measure has obscured a
540 true effect of the EHST and/or the meaningfulness of this variability during more severe MT.

541 Evidence directly comparing correlations between GI barrier integrity and/or MT
542 biomarkers in exercise settings has been limited to date. Given general logistical constraints
543 of the urine/plasma DSAT, the majority of relevant evidence has attempted to validate
544 (correlate) this method against more practical GI barrier integrity biomarkers. These studies
545 have generally shown significant, though weak correlations ($r = 0.4-0.6$) between basal
546 corrected (Δ) DSAT (urine 5 hour) and I-FABP responses directly following minor exercise-
547 induced GI barrier integrity loss (van Wijck et al., 2011, 2012; March et al., 2017). In the
548 present study, the DSAT did not correlate with any other GI integrity biomarkers. A potential
549 explanation for this null finding might result from the lack of basal DSAT correction or the low
550 overall severity of GI barrier integrity loss. In comparison, a small positive correlation was
551 reported between I-FABP and CLDN-3. This finding is supportive of previous evidence showing
552 urinary I-FABP and CLDN-3 to weakly correlate ($r = 0.38$) in patients with major non-abdominal
553 surgery (Habes et al., 2017). The expression of CLDN-3 across multiple tissues might partially
554 explain why this correlation was not stronger (Thuijls et al., 2010a). In general, no GI barrier
555 integrity and MT biomarkers, except DSAT 150 and Δ LBP, were found to correlate. Previous
556 exercise gastroenterology research has shown various combinations of these biomarkers to
557 weakly correlate ($r = 0.1-0.6$; Yeh et al., 2013; Sessions et al., 2016; March et al., 2019) or not
558 correlate (Karhu et al., 2017; Snipe et al., 2018). Several physiological (e.g. hepatic/immune
559 microbial clearance, transcellular microbial translocation, GI microbial density) and analytical

560 (e.g. exogenous sample contamination, inconsistent biomarker kinetics, DSAT/I-FABP limited
561 to small GI integrity) factors all likely weaken this association (Wells et al, 2017).

562 **LIMITATIONS**

563 Despite implementation of a tightly controlled methodological design, which
564 accounted for the majority of extraneous variables, the presented results were not without
565 some limitations. First, the EHST was only able to evoke moderate GI barrier integrity loss and
566 did not influence MT, potentially limiting the application of these findings in severe situations
567 of GI barrier integrity loss. A previous systematic review has suggested an exercise induced
568 hyperthermia threshold of 38.6°C T_{core} for GI barrier integrity loss (DSAT, I-FABP and
569 endotoxin) to be commonplace (>50% incidence) and of 39.0°C for GI barrier integrity loss to
570 be universal (100% incidence; Pires et al., 2017). Consistently, previous research supports the
571 notion that MT biomarkers (endotoxin) are less responsive to subtle alterations in GI barrier
572 integrity that were otherwise detected by the DSAT or I-FABP following exercise (Snipe et al.,
573 2017, 2018; March et al., 2019). Positively, no GI barrier integrity or MT biomarker displayed
574 statistical heteroscedasticity in this EHST model, suggestive that absolute reliability was not
575 dependent upon the magnitude of biomarker response. Next, biomarker analysis was limited
576 to a single time-point after the EHST (at termination), though this can be justified in that peak
577 responses have been consistently shown to occur at this instance in comparable exertional-
578 heat stress interventions (e.g. *I-FABP*, Snipe et al., 2017; *CLDN-3*, Yeh et al., 2013; *LBP*,
579 Moncada-Jimenez et al., 2010; *Bact./16S*, March et al., 2019). Third, there was statistically
580 significant systematic bias for peak T_{core} and T_{body} responses, which were lower (0.17°C and
581 0.18°C) following implementation of trial two. This result was not anticipated, given
582 numerous previous studies showing a one week washout period to be sufficient in preventing
583 carry-over (heat acclimation) effects following exertional-heat stress exposure of comparable
584 severity (Barrett and Maughan, 1993; Willmott et al., 2015). The meaningfulness of this
585 systematic bias did not appear to statistically influence any GI barrier integrity or MT
586 biomarker. Finally, given neither a basal DSAT or urinary DSAT were performed, it was not
587 possible to directly determine either the impact of the EHST on DSAT results or make
588 comparisons between DSAT responses between biofluids. This decision was made to
589 minimise participant time burden.

590 **CONCLUSION**

591 This is the first study to comprehensively assess the reliability of GI barrier integrity
592 and/or microbial translocation biomarkers both at rest and following exertional (-heat) stress.
593 Quantifying biomarker reliability is a vital step required to inform marker selection for
594 application in laboratory and field settings. Each of the GI barrier integrity biomarkers
595 assessed displayed moderate-to-good relative and acceptable absolute reliability both at rest
596 and post the EHST. Serum DSAT responses had comparable reliability at two-separate time-
597 points following sugar-probe ingestion (90- and 150-minutes), though response kinetics
598 displayed inconsistent time courses. I-FABP and CLDN-3 both increased following the EHST
599 and their responses were found to weakly correlate. None of the selected MT biomarkers
600 were elevated following the EHST, suggestive that a greater severity of GI barrier integrity loss
601 is required for MT. LBP and total 16S DNA both demonstrated moderate-to-good relative and
602 acceptable absolute reliability at both time-points. There was a weak correlation between
603 LBP and total 16S post-EHST responses. Despite offering superior methodological rationale,
604 *Bacteroides* DNA had unacceptable reliability. The findings of the present study have direct
605 relevance for evaluating the efficacy of interventions to attenuate the rise in GI barrier
606 integrity/MT when exercising in the heat. Such interventions might include exercise training,
607 heat acclimatisation and nutritional supplementation. The findings of this study might also
608 have value to the pharmaceutical industry, to quantify the efficacy of drugs to maintain GI
609 barrier integrity, or to evaluate improvements in drugs that traditionally resulted in GI
610 integrity loss.

611

612

613 **COMPETING INTERESTS**

614 No competing interests

615 **AUTHOR CONTRIBUTION**

616 HO, JF, RC, SD, CW and JL concepted and designed the research; HO, AM, CW performed the
617 experiments; HO, GD, SF and RE acquired data; HO, JF, RC, GD, SD and JL interpreted the
618 results; HO wrote the manuscript; HO, JF, RC, GD, SF, RE, AM, CW, AM and JL edited, revised
619 and agree to accountability of the accuracy and integrity of the manuscript. Data were
620 collected at School of Sport, Health and Wellbeing, Plymouth MARJON University. All persons
621 designated as authors qualify for authorship, and all those who qualify for authorship are
622 listed.

623 **FUNDING**

624 No funding was received for this research

625 **ACKNOWLEDGMENTS**

626 We wish to thank the participants for volunteering their time and effort to take part in this
627 research

628

629 **REFERENCES**

- 630 Aguinis, H., Gottfredson, R.K. and Joo, H. (2013). Best-practice recommendations for defining,
631 identifying, and handling outliers. *Organizational Research Methods*, 16(2), pp.270-301.
- 632 Armed Forces Surveillance Branch. (2018). Update: Heat illness, active component, US Armed
633 Forces, 2017. *Medical Surveillance Monthly Report*, 25(4), pp.6.
- 634 Atkinson, G. and Nevill, A.M. (1998). Statistical methods for assessing measurement error
635 (reliability) in variables relevant to sports medicine. *Sports Medicine*, 26(4), pp.217-238.
- 636 Barberio, M.D., Elmer, D.J., Laird, R.H., Lee, K.A., Gladden, B. and Pascoe, D.D. (2015).
637 Systemic LPS and inflammatory response during consecutive days of exercise in
638 heat. *International Journal of Sports Medicine*, 36(03), pp.262-270.
- 639 Barnett, A. and Maughan, R.J. (1993). Response of unacclimatized males to repeated weekly
640 bouts of exercise in the heat. *British Journal of Sports Medicine*, 27(1), pp.39-44.
- 641 Bischoff, S.C., Barbara, G., Buurman, W., Ockhuizen, T., Schulzke, J.D., Serino, M., Tilg, H.,
642 Watson, A. and Wells, J.M. (2014). Intestinal permeability—a new target for disease
643 prevention and therapy. *BMC Gastroenterology*, 14(1), p.189.
- 644 Bland, J.M. and Altman, D. (1986). Statistical methods for assessing agreement between two
645 methods of clinical measurement. *The Lancet*, 327(8476), pp.307-310.
- 646 Borg, G. (1970). Perceived exertion as an indicator of somatic stress. *Scandinavian Journal of*
647 *Rehabilitation Medicine*, 2, pp.92-98.
- 648 Camilleri, M., Madsen, K., Spiller, R., Van Meerveld, B.G. and Verne, G.N. (2012). Intestinal
649 barrier function in health and gastrointestinal disease. *Neurogastroenterology &*
650 *Motility*, 24(6), pp.503-512.
- 651 Cani, P.D. (2018). Human gut microbiome: hopes, threats and promises. *Gut*, 67(9), pp.1716-
652 1725.

653 Christensen, M.J., Eller, E., Kjaer, H.F., Broesby-Olsen, S., Mortz, C.G. and Bindselev-Jensen, C.
654 (2019). Exercise-induced anaphylaxis: causes, consequences, and management
655 recommendations. *Expert Review of Clinical Immunology*, 15(3), pp.265-273.

656 Citronberg, J.S., Wilkens, L.R., Lim, U., Hullar, M.A., White, E., Newcomb, P.A., Le Marchand,
657 L. and Lampe, J.W. (2016). Reliability of plasma lipopolysaccharide-binding protein (LBP)
658 from repeated measures in healthy adults. *Cancer Causes and Control*, 27(9), pp.1163-1166.

659 Costa, R.J.S., Snipe, R.M.J., Kitic, C.M. and Gibson, P.R. (2017). Systematic review: exercise-
660 induced gastrointestinal syndrome—implications for health and intestinal
661 disease. *Alimentary Pharmacology and Therapeutics*, 46(3), pp.246-265.

662 Dill, D.B. and Costill, D.L. (1974). Calculation of percentage changes in volumes of blood,
663 plasma, and red cells in dehydration. *Journal of Applied Physiology*, 37(2), pp.247-248.

664 Dullah, E.C. and Ongkudon, C.M. (2017). Current trends in endotoxin detection and analysis
665 of endotoxin–protein interactions. *Critical Reviews in Biotechnology*, 37(2), pp.251-261.

666 Durnin, J.V. and Womersley, J.V.G.A. (1974). Body fat assessed from total body density and
667 its estimation from skinfold thickness: measurements on 481 men and women aged from 16
668 to 72 years. *British Journal of Nutrition*, 32(1), pp.77-97.

669 Edinburgh, R.M., Hengist, A., Smith, H.A., Travers, R.L., Koumanov, F., Betts, J.A., Thompson,
670 D., Walhin, J.P., Wallis, G.A., Hamilton, D.L. and Stevenson, E.J. (2018). Preexercise breakfast
671 ingestion versus extended overnight fasting increases postprandial glucose flux after
672 exercise in healthy men. *American Journal of Physiology-Endocrinology and*
673 *Metabolism*, 315(5), pp.1062-1074.

674 Fleming, S.C., Duncan, A., Russell, R.I. and Laker, M.F. (1996). Measurement of sugar probes
675 in serum: an alternative to urine measurement in intestinal permeability testing. *Clinical*
676 *Chemistry*, 42(3), pp.445-448.

677 Glassing, A., Dowd, S.E., Galandiuk, S., Davis, B. and Chiodini, R.J. (2016). Inherent bacterial
678 DNA contamination of extraction and sequencing reagents may affect interpretation of
679 microbiota in low bacterial biomass samples. *Gut Pathogens*, 8, pp.24-24.

680 Guy, J.H., Edwards, A.M., Miller, C.M., Deakin, G.B. and Pyne, D.B. (2017). Short-term
681 reliability of inflammatory mediators and response to exercise in the heat. *Journal of Sports*
682 *Sciences*, 35(16), pp.1622-1628.

683 Habes, Q.L., Linssen, V., Nooijen, S., Kiers, D., Gerretsen, J., Pickkers, P., Scheffer, G.J. and
684 Kox, M. (2017). Markers of intestinal damage and their relation to cytokine levels in cardiac
685 surgery patients. *Shock*, 47(6), pp.709-714.

686 Huttenhower, C., Gevers, D., Knight, R., Abubucker, S., Badger, J.H., Chinwalla, A.T., Creasy,
687 H.H., Earl, A.M., FitzGerald, M.G., Fulton, R.S. and Giglio, M.G. (2012). Structure, function
688 and diversity of the healthy human microbiome. *Nature*, 486(7402), p.207.

689 JanssenDuijghuijsen, L.M., Mensink, M., Lenaerts, K., Fiedorowicz, E., Protégé Study Group,
690 van Dartel, D.A., Mes, J.J., Luiking, Y.C., Keijer, J., Wichers, H.J. and Witkamp, R.F. (2016).
691 The effect of endurance exercise on intestinal integrity in well-trained healthy
692 men. *Physiological Reports*, 4(20), p.e12994.

693 Jay, O. and Kenny, G.P. (2007). The determination of changes in body heat content during
694 exercise using calorimetry and thermometry. *Journal of the Human-Environment*
695 *System*, 10(1), pp.19-29.

696 Julious, S. A. (2005). Sample size of 12 per group rule of thumb for a pilot study.
697 *Pharmaceutical Statistics*, 4 (4), pp. 287–291.

698 Jonvik, K.L., Lenaerts, K., Smeets, J.S., Kolkman, J.J., Van Loon, L.J. and Verdijk, L.B. (2019).
699 Sucrose but Not Nitrate Ingestion Reduces Strenuous Cycling-induced Intestinal
700 Injury. *Medicine and Science in Sports and Exercise*, 51, pp.436-444.

701 Karhu, E., Forsgård, R.A., Alanko, L., Alfthan, H., Pussinen, P., Hämmäläinen, E. and Korpela, R.
702 (2017). Exercise and gastrointestinal symptoms: running-induced changes in intestinal
703 permeability and markers of gastrointestinal function in asymptomatic and symptomatic
704 runners. *European Journal of Applied Physiology*, 117(12), pp.2519-2526.

705 Lim, C. (2018). Heat sepsis precedes heat toxicity in the pathophysiology of heat stroke—a
706 new paradigm on an ancient disease. *Antioxidants*, 7(11), p.149.

707 March, D.S., Jones, A.W., Thatcher, R. and Davison, G. (2019). The effect of bovine
708 colostrum supplementation on intestinal injury and circulating intestinal bacterial DNA
709 following exercise in the heat. *European Journal of Nutrition*, 58(4), pp.1441-1451.

710 March, D.S., Marchbank, T., Playford, R.J., Jones, A.W., Thatcher, R. and Davison, G. (2017).
711 Intestinal fatty acid-binding protein and gut permeability responses to exercise. *European*
712 *Journal of Applied Physiology*, 117(5), pp.931-941.

713 Marchbank, T., Davison, G., Oakes, J.R., Ghatei, M.A., Patterson, M., Moyer, M.P. and
714 Playford, R.J. (2010). The nutraceutical bovine colostrum truncates the increase in gut
715 permeability caused by heavy exercise in athletes. *American Journal of Physiology-*
716 *Gastrointestinal and Liver Physiology*, 300(3), pp.477-484.

717 Marfell-Jones, M., Olds, T., Stewart, A. and Carter, L. (2006). *ISAK manual: international*
718 *standards for anthropometric assessment*. South Africa: Potchefstroom.

719 Met Office. (2019). Cambourne. [Online]. Available at:
720 [https://www.metoffice.gov.uk/pub/data/weather/uk/climate/stationdata/cambornedata.tx](https://www.metoffice.gov.uk/pub/data/weather/uk/climate/stationdata/cambornedata.txt)
721 [t](https://www.metoffice.gov.uk/pub/data/weather/uk/climate/stationdata/cambornedata.txt) [Accessed 14/10/2019]

722 Military Headquarters of the Surgeon General. (2017). [Online]. Available at:
723 [https://assets.publishing.service.gov.uk/government/uploads/system/uploads/attachment_](https://assets.publishing.service.gov.uk/government/uploads/system/uploads/attachment_data/file/793094/JSP_539_Part_2_V3.1_Updated_04-19_.pdf)
724 [data/file/793094/JSP_539_Part_2_V3.1_Updated_04-19_.pdf](https://assets.publishing.service.gov.uk/government/uploads/system/uploads/attachment_data/file/793094/JSP_539_Part_2_V3.1_Updated_04-19_.pdf) [Accessed 14/10/2019]

725 Moncada-Jiménez, J., Plaisance, E.P., Araya-Ramírez, F., Taylor, J.K., Ratcliff, L., Mestek, M.L.,
726 Grandjean, P.W. and AragonVargas, L.F. (2010). Acute hepatic response to diet modification
727 and exercise-induced endotoxemia during a laboratory-based duathlon. *Biology of*
728 *Sport*, 27(2), pp. 111-118.

729 Morrison, S.A., Cheung, S.S. and Cotter, J.D. (2014). Bovine colostrum, training status, and
730 gastrointestinal permeability during exercise in the heat: a placebo-controlled double-blind
731 study. *Applied Physiology, Nutrition, and Metabolism*, 39(9), pp.1070-1082.

732 Mortensen, C., Karlsen, S., Grønbaek, H., Nielsen, D.T., Frevert, S., Clemmesen, J.O., Møller,
733 S., Jensen, J.S. and Bendtsen, F. (2013). No difference in portal and hepatic venous bacterial
734 DNA in patients with cirrhosis undergoing transjugular intrahepatic portosystemic shunt
735 insertion. *Liver International*, 33(9), pp.1309-1315.

736 Mulak, A. and Bonaz, B. (2015). Brain-gut-microbiota axis in Parkinson's disease. *World*
737 *Journal of Gastroenterology*, 21(37), p.10609.

738 Neves, A.L., Coelho, J., Couto, L., Leite-Moreira, A. and Roncon-Albuquerque, J.R., 2013.
739 Metabolic endotoxemia: a molecular link between obesity and cardiovascular risk. *Journal of*
740 *molecular endocrinology*, 51(2), pp.R51-64.

741 Païssé, S., Valle, C., Servant, F., Courtney, M., Burcelin, R., Amar, J. and Lelouvier, B. (2016).
742 Comprehensive description of blood microbiome from healthy donors assessed by 16 S
743 targeted metagenomic sequencing. *Transfusion*, 56(5), pp.1138-1147.

744 Pires, W., Veneroso, C.E., Wanner, S.P., Pacheco, D.A., Vaz, G.C., Amorim, F.T., Tonoli, C.,
745 Soares, D.D. and Coimbra, C.C. (2017). Association between exercise-induced hyperthermia
746 and intestinal permeability: a systematic review. *Sports Medicine*, 47(7), pp.1389-1403.

747 Pugh, J.N., Impey, S.G., Doran, D.A., Fleming, S.C., Morton, J.P. and Close, G.L. (2017a).
748 Acute high-intensity interval running increases markers of gastrointestinal damage and
749 permeability but not gastrointestinal symptoms. *Applied Physiology, Nutrition, and*
750 *Metabolism*, 42(9), pp.941-947.

751 Ramanathan, N.L. (1964). A new weighting system for mean surface temperature of the
752 human body. *Journal of Applied Physiology*, 19(3), pp.531-533.

753 Schellekens, D.H., Grootjans, J., Dello, S.A., van Bijnen, A.A., van Dam, R.M., Dejong, C.H.,
754 Derikx, J.P. and Buurman, W.A. (2014). Plasma intestinal fatty acid-binding protein levels
755 correlate with morphologic epithelial intestinal damage in a human translational ischemia-
756 reperfusion model. *Journal of Clinical Gastroenterology*, 48(3), pp.253-260.

757 Schumann, R.R. (2011). Old and new findings on lipopolysaccharide-binding protein: a
758 soluble pattern-recognition molecule. *Biochemical Society Transactions*, 39(4), pp.989-993.

759 Selkirk, G.A., McLellan, T.M., Wright, H.E. and Rhind, S.G. (2008). Mild endotoxemia, NF- κ B
760 translocation, and cytokine increase during exertional heat stress in trained and untrained
761 individuals. *American Journal of Physiology-Regulatory, Integrative and Comparative*
762 *Physiology*, 295(2), pp.611-623.

763 Sessions, J., Bourbeau, K., Rosinski, M., Szczygiel, T., Nelson, R., Sharma, N. and Zuhl, M.
764 (2016). Carbohydrate gel ingestion during running in the heat on markers of gastrointestinal
765 distress. *European Journal of Sport Science*, 16(8), 1064-1072.

766 Shirreffs, S.M. and Maughan, R.J. (1994). The effect of posture change on blood volume,
767 serum potassium and whole body electrical impedance. *European Journal of Applied*
768 *Physiology and Occupational Physiology*, 69(5), pp.461-463.

769 Snipe, R.M., Khoo, A., Kitic, C.M., Gibson, P.R. and Costa, R.J. (2017). Carbohydrate and
770 protein intake during exertional heat stress ameliorates intestinal epithelial injury and small
771 intestine permeability. *Applied Physiology, Nutrition, and Metabolism*, 42(12), pp.1283-
772 1292.

773 Snipe, R.M., Khoo, A., Kitic, C.M., Gibson, P.R. and Costa, R.J. (2018). The impact of
774 exertional-heat stress on gastrointestinal integrity, gastrointestinal symptoms, systemic
775 endotoxin and cytokine profile. *European Journal of Applied Physiology*, 118(2), pp.389-400.

776 Thuijls, G., Derikx, J.P., de Haan, J.J., Grootjans, J., de Bruïne, A., Masclee, A.A., Heineman, E.
777 and Buurman, W.A. (2010a). Urine-based detection of intestinal tight junction loss. *Journal*
778 *of Clinical Gastroenterology*, 44(1), pp.e14-e19.

779 Thuijls, G., Derikx, J.P., van Wijck, K., Zimmermann, L.J., Degraeuwe, P.L., Mulder, T.L., Van
780 der Zee, D.C., Brouwers, H.A., Verhoeven, B.H., van Heurn, L.E. and Kramer, B.W. (2010b).
781 Non-invasive markers for early diagnosis and determination of the severity of necrotizing
782 enterocolitis. *Annals of Surgery*, 251(6), pp.1174-1180.

783 Toner, M.M., Drolet, L.L. and Pandolf, K.B. (1986). Perceptual and physiological responses
784 during exercise in cool and cold water. *Perceptual and Motor Skills*, 62(1), pp.211-220.

785 Treskes, N., Persoon, A.M. and van Zanten, A.R. (2017). Diagnostic accuracy of novel
786 serological biomarkers to detect acute mesenteric ischemia: a systematic review and meta-
787 analysis. *Internal and Emergency Medicine*, 12(6), pp.821-836.

788 Typpo, K.V., Larmonier, C.B., Deschenes, J., Redford, D.T., Kiela, P.R. and Ghishan, F.K.
789 (2015). Clinical characteristics associated with postoperative intestinal epithelial barrier
790 dysfunction in children with congenital heart disease. *Pediatric Critical Care Medicine*, 16(1),
791 pp.37-44.

792 Van Elburg, R.M., Uil, J.J., Kokke, F.T.M., Mulder, A.M., Van De Broek, W.G.M., Mulder, C.J.J.
793 and Heymans, H.S.A. (1995). Repeatability of the sugar-absorption test, using lactulose and
794 mannitol, for measuring intestinal permeability for sugars. *Journal of Pediatric*
795 *Gastroenterology and Nutrition*, 20(2), pp.184-188.

796 Van Wijck, K., Lenaerts, K., Van Bijnen, A.A., Boonen, B., Van Loon, L.J., Dejong, C.H. and
797 Buurman, W.A. (2012). Aggravation of exercise-induced intestinal injury by Ibuprofen in
798 athletes. *Medicine and Science in Sports and Exercise*, 44(12), pp.2257-2262.

799 Van Wijck, K., Lenaerts, K., Van Loon, L.J., Peters, W.H., Buurman, W.A. and Dejong, C.H.,
800 (2011). Exercise-induced splanchnic hypoperfusion results in gut dysfunction in healthy
801 men. *PloS One*, 6(7), pp.e22366.

802 Van Wijck, K., Verlinden, T.J., van Eijk, H.M., Dekker, J., Buurman, W.A., Dejong, C.H. and
803 Lenaerts, K. (2013). Novel multi-sugar assay for site-specific gastrointestinal permeability
804 analysis: a randomized controlled crossover trial. *Clinical Nutrition*, 32(2), pp.245-251.

805 Velders, M., Treff, G., Machus, K., Bosnyák, E., Steinacker, J. and Schumann, U. (2014).
806 Exercise is a potent stimulus for enhancing circulating DNase activity. *Clinical*
807 *Biochemistry*, 47(6), pp.471-474.

808 Vincent, W.J. and Weir, J.P. (1995). *Quantifying reliability*. In *Statistics in Kinesiology* (5th
809 Ed.). Champaign, IL, USA: Human Kinetics , pp.214-228.

810 Waterhouse, J., Drust, B., Weinert, D., Edwards, B., Gregson, W., Atkinson, G., Kao, S.,
811 Aizawa, S. and Reilly, T. (2005). The circadian rhythm of core temperature: origin and some
812 implications for exercise performance. *Chronobiology International*, 22(2), pp.207-225.

813 Wells, J.M., Brummer, R.J., Derrien, M., MacDonald, T.T., Troost, F., Cani, P.D., Theodorou,
814 V., Dekker, J., Méheust, A., De Vos, W.M. and Mercenier, A. (2017). Homeostasis of the gut
815 barrier and potential biomarkers. *American Journal of Physiology-Gastrointestinal and Liver*
816 *Physiology*, 312(3), pp.171-193.

817 Yeh, Y.J., Law, L.Y.L. and Lim, C.L., 2013. Gastrointestinal response and endotoxemia during
818 intense exercise in hot and cool environments. *European Journal of Applied*
819 *Physiology*, 113(6), pp.1575-1583.

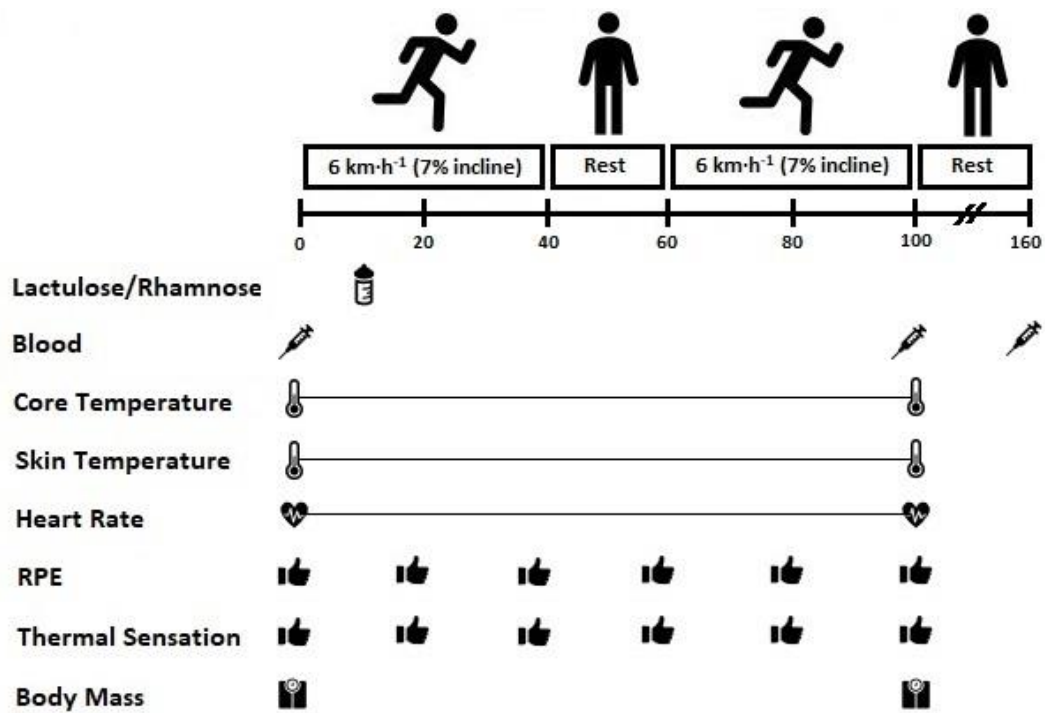
820 Zeissig, S., Bürgel, N., Günzel, D., Richter, J., Mankertz, J., Wahnschaffe, U., Kroesen, A.J.,
821 Zeitz, M., Fromm, M. and Schulzke, J.D. (2007). Changes in expression and distribution of
822 claudin 2, 5 and 8 lead to discontinuous tight junctions and barrier dysfunction in active
823 Crohn's disease. *Gut*, 56(1), pp.61-72.

824 **Table 1.** Participant demographic characteristics

Measure	Mean ± SD
Age (years)	26 ± 5

Height (m)	1.78 ± 0.06
Body Mass (kg)	83.4 ± 12.6
Physical Activity (h·week ⁻¹)	6 ± 3
Body Fat (%)	16.1 ± 4.0
$\dot{V}O_{2max}$ (ml·kg ⁻¹ ·min ⁻¹)	49 ± 4

825



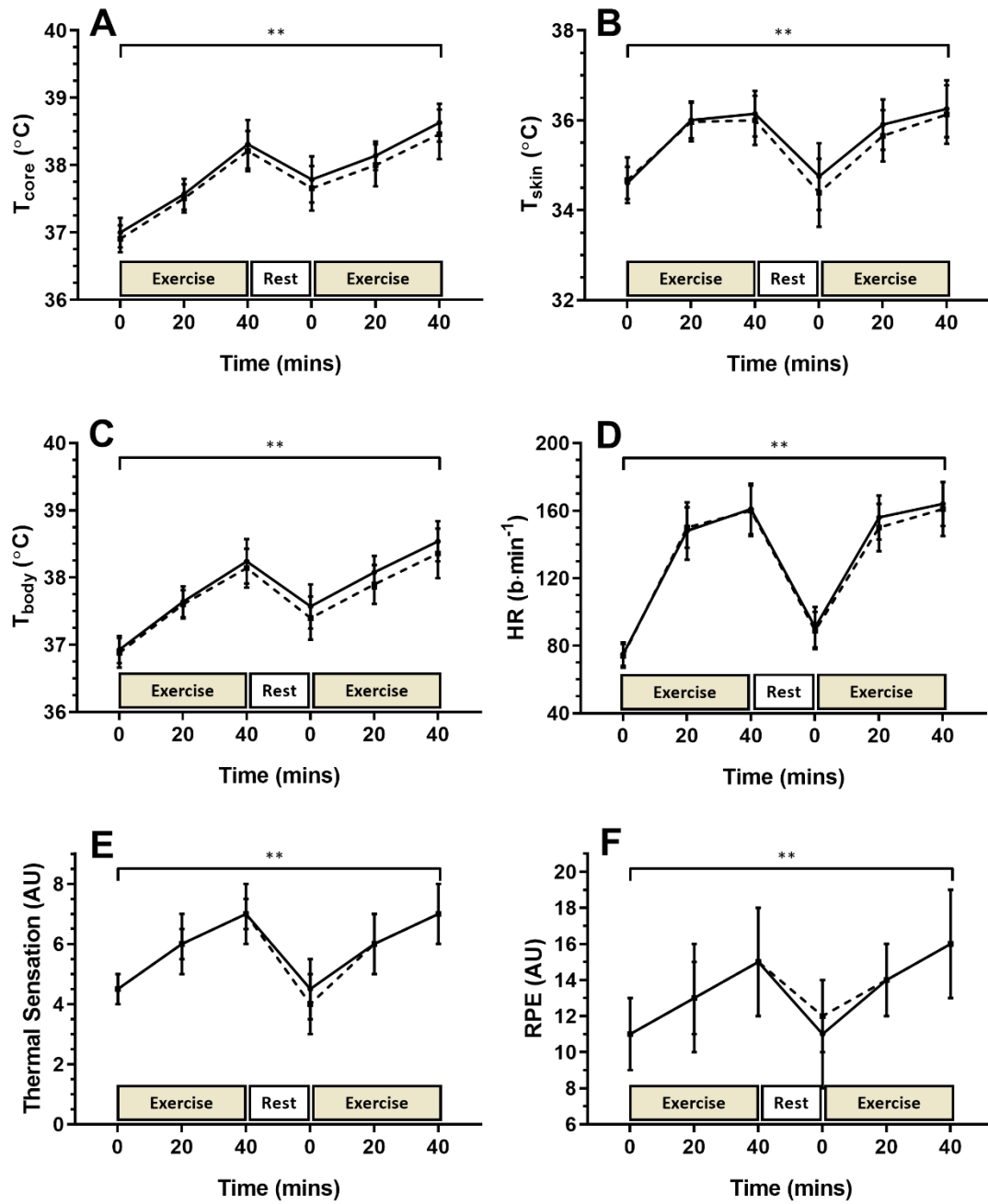
826

827 **Figure 1.** Schematic illustration of the experimental measurement timings

828

829

830



831

832 **Figure 2.** Whole-body physiological responses to repeated EHSTs: (A) = core temperature;
 833 (B) = mean skin temperature (n=13); (C) = mean body temperature (n=13); (D) = heart rate;
 834 (E) = thermal sensation; and (F) = rate of perceived exertion. Solid line = trial 1, broken line =
 835 trial 2. Significant overall effect of time ($*p \leq 0.05$; $** p \leq 0.01$).

836

837

Table 2. Relative and absolute reliability of whole-body physiological responses

	Trial 1 (SD)	Trial 2 (SD)	<i>p</i>	<i>d</i>	<i>r</i>	CV	TEM	Bias (LoA)
T _{core} (°C) Peak	38.63 ± 0.28	38.46 ± 0.37	0.02	0.49	0.78**	0.5	0.14	0.17 ± 0.46
T _{core} (°C) Mean	37.87 ± 0.19	37.78 ± 0.23	0.09	0.43	0.68**	0.3	0.13	0.09 ± 0.43
T _{core} (°C) Δ	1.62 ± 0.29	1.55 ± 0.44	0.59	0.19	0.48	-	0.26	0.07 ± 0.77
T _{skin} (°C) Peak	36.27 ± 0.63	36.13 ± 0.65	0.23	0.22	0.83**	0.6	0.27	0.13 ± 0.74
T _{skin} (°C) Mean	35.75 ± 0.44	35.64 ± 0.49	0.13	0.24	0.83**	0.4	0.17	0.11 ± 0.46
T _{skin} (°C) Δ	1.65 ± 0.63	1.46 ± 0.65	0.18	0.30	0.71**	-	0.35	0.20 ± 0.97
T _{body} (°C) Peak	38.54 ± 0.30	38.36 ± 0.37	0.01	0.53	0.79**	0.4	0.16	0.18 ± 0.44
T _{body} (°C) Mean	37.83 ± 0.20	37.74 ± 0.23	0.08	0.42	0.70**	0.3	0.12	0.09 ± 0.33
T _{body} (°C) Δ	1.62 ± 0.30	1.49 ± 0.40	0.18	0.37	0.60*	-	0.23	0.13 ± 0.63
HR (b·min ⁻¹) Peak	164 ± 13	162 ± 14	0.19	0.15	0.93**	2.0	4	2 ± 11
HR (b·min ⁻¹) Mean	150 ± 14	148 ± 13	0.29	0.14	0.84**	2.8	5	2 ± 15
HR (b·min ⁻¹) Δ	90 ± 11	90 ± 13	0.99	0.00	0.88**	-	4	0 ± 12
RPE (AU) Peak	16 ± 3	16 ± 3	0.99	0.00	0.92**	4.1	1	0 ± 3
RPE (AU) Mean	13 ± 2	13 ± 2	0.88	0.00	0.80**	5.2	1	0 ± 2
RPE (AU) Δ	6 ± 5	5 ± 3	0.95	0.24	0.72**	-	2	0 ± 4
TS (AU) Peak	7.0 ± 0.5	7.0 ± 0.5	0.99	0.00	0.84**	4.8	0.5	0.0 ± 1.0
TS (AU) Mean	6.0 ± 0.5	5.5 ± 0.5	0.02	0.00	0.88**	3.2	0.0	0.5 ± 0.5
TS (AU) Δ	2.5 ± 1.0	2.5 ± 0.5	0.12	0.00	0.61*	-	0.5	-0.5 ± 1.0

* = significant correlation ($p \leq 0.05$); ** = significant correlation ($p \leq 0.01$)

Table 3. Relative and absolute reliability of all GI barrier integrity biomarkers

	Trial 1 (SD)	Trial 2 (SD)	<i>p</i>	<i>d</i>	<i>r</i>	CV	TEM	Bias (LoA)
Lactulose (mg·l ⁻¹) 90	1.06 ± 0.38	0.90 ± 0.43	0.33	0.38	0.60*	21.3	0.263	0.151 ± 0.730
L-Rhamnose (mg·l ⁻¹) 90	15.89 ± 3.91	15.85 ± 3.13	0.29	0.29	0.53	12.9	2.601	1.036 ± 6.930
DSAT (L/R) 90	0.028 ± 0.012	0.025 ±0.014	0.17	0.23	0.77**	11.5	0.004	-0.003 ± 0.011
Lactulose (mg·l ⁻¹) 150	0.97 ± 0.48	0.95 ± 0.52	0.53	0.05	0.71**	13.0	0.132	0.023 ± 0.364
L-Rhamnose (mg·l ⁻¹) 150	12.01 ± 2.95	11.24 ± 2.96	0.09	0.27	0.86**	7.6	1.149	0.772 ± 3.060
DSAT (L/R) 150	0.033 ± 0.015	0.034 ± 0.016	0.37	0.06	0.69**	12.2	0.004	0.001 ± 0.011
I-FABP (ng·ml ⁻¹) Rest	1.560 ± 0.506	1.691 ± 0.555	0.11	0.25	0.75**	11.1	0.304	-0.180 ± 0.746
I-FABP (ng·ml ⁻¹) Post	2.394 ± 0.731	2.467 ± 0.875	0.63	0.09	0.80**	12.1	0.376	-0.073 ± 1.040
I-FABP (ng·ml ⁻¹) Δ	0.834 ± 0.445	0.776 ± 0.489	0.65	0.12	0.65**	-	0.278	0.058 ± 0.772
CLDN-3 (ng·ml ⁻¹) Rest	6.205 ± 4.382	5.971 ± 4.062	0.17	0.06	0.99**	6.8	0.423	0.233 ± 1.172
CLDN-3 (ng·ml ⁻¹) Post	6.592 ± 4.770	6.323 ± 4.270	0.34	0.06	0.99**	4.9	0.485	0.181 ± 1.341
CLDN-3 (ng·ml ⁻¹) Δ	0.317 ± 0.586	0.371 ± 0.508	0.68	0.10	0.62*	-	0.342	-0.055 ± 0.948

841 * = significant correlation ($p \leq 0.05$); ** = significant correlation ($p \leq 0.01$)

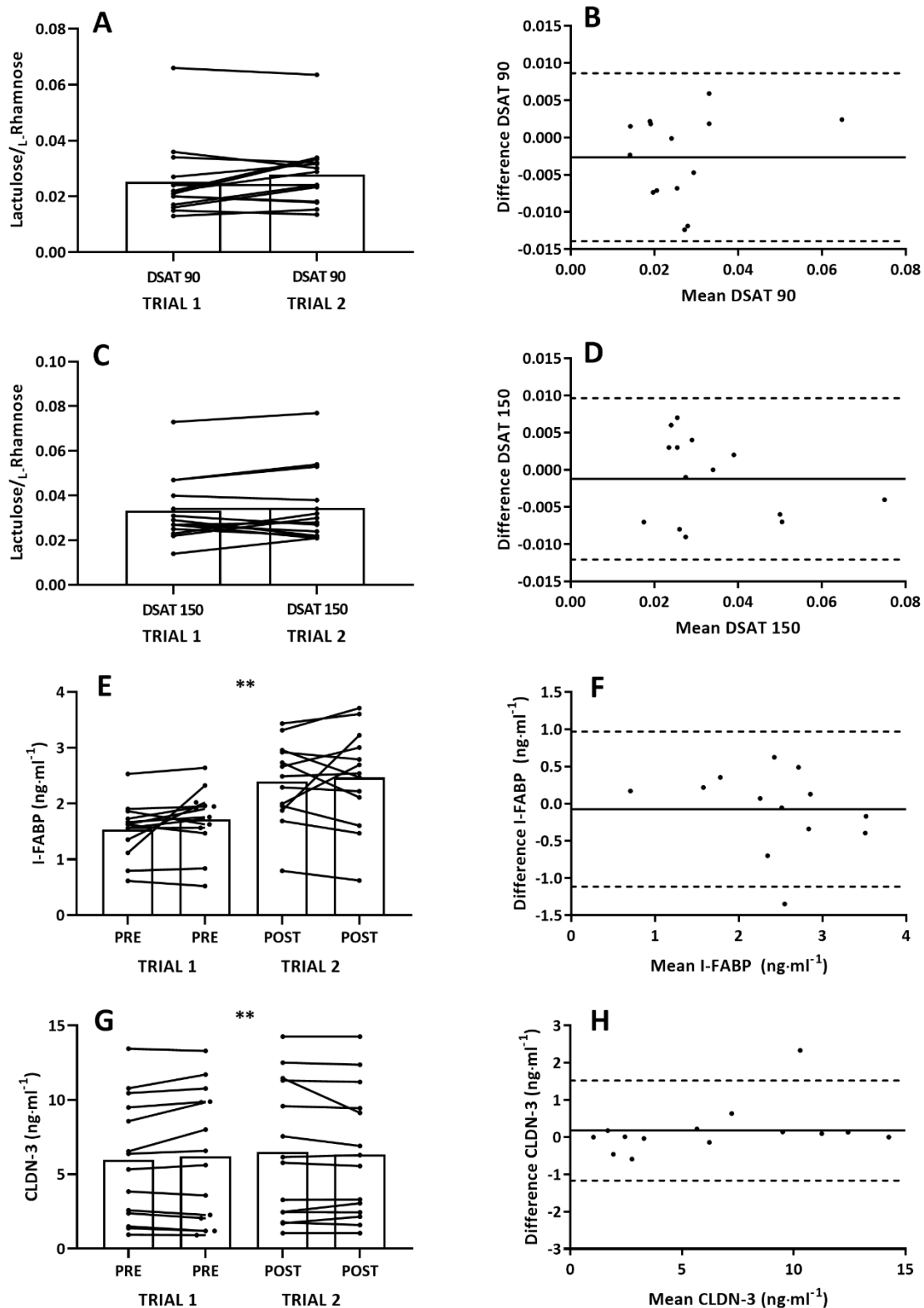
842

843

844

845

846



847

848

849

850

851

852

Figure 3. GI barrier integrity responses to EHST trial 1 and trial 2: (A) = L/R ratio at 90 minutes; (C) = L/R ratio at 150 minutes; (E) I-FABP (n=13); and (G) = CLDN-3. Significant overall effect of time (* $p \leq 0.05$; ** $p \leq 0.01$). Bland-Altman mean bias and 95% LoA between post EHST trial 1 and trial 2: (B) = L/R ratio at 90- and (D) 150- minutes; (F) I-FABP (n=13); and (H) = CLDN-3

Table 4. Relative and absolute reliability of all GI barrier integrity biomarkers

	Trial 1 (SD)	Trial 2 (SD)	<i>p</i>	<i>d</i>	<i>r</i>	CV	TEM	Bias (LoA)
LBP ($\mu\text{g}\cdot\text{ml}^{-1}$) Rest	2.625 \pm 0.993	2.564 \pm 0.871	0.79	0.07	0.85**	10.0	0.378	0.061 \pm 1.048
LBP ($\mu\text{g}\cdot\text{ml}^{-1}$) Post	2.617 \pm 1.080	2.650 \pm 0.885	0.59	0.03	0.85**	9.2	0.420	-0.033 \pm 1.166
LBP ($\mu\text{g}\cdot\text{ml}^{-1}$) Δ	-0.008 \pm 0.250	0.086 \pm 0.232	0.38	0.39	-0.16	-	0.260	-0.094 \pm 0.721
16S DNA ($\text{pg}\cdot\mu\text{l}^{-1}$) Rest	1.43 \pm 0.60	1.44 \pm 0.65	0.79	0.02	0.87**	8.1	0.19	-0.02 \pm 0.49
16S DNA ($\text{pg}\cdot\mu\text{l}^{-1}$) Post	1.47 \pm 0.70	1.52 \pm 0.73	0.45	0.07	0.82**	9.5	0.15	-0.05 \pm 0.40
16S DNA ($\text{pg}\cdot\mu\text{l}^{-1}$) Δ	0.043 \pm 0.28	0.08 \pm 0.40	0.72	0.10	0.56*	-	0.24	-0.03 \pm 0.65
<i>Bact.</i> DNA (copies $\cdot\mu\text{l}^{-1}$) Rest	0.142 \pm 0.116	0.102 \pm 0.089	0.19	0.39	0.14	55.0	0.067	0.040 \pm 0.186
<i>Bact.</i> DNA (copies $\cdot\mu\text{l}^{-1}$) Post	0.202 \pm 0.132	0.115 \pm 0.106	0.04*	0.73	0.14	56.3	0.104	0.087 \pm 0.287
<i>Bact.</i> DNA (copies $\cdot\mu\text{l}^{-1}$) Δ	0.059 \pm 0.149	0.013 \pm 0.140	0.22	0.32	0.61*	-	0.091	0.047 \pm 0.250
<i>Bact.</i> /16S DNA Rest	96.08 \pm 59.31	74.10 \pm 52.82	0.31	0.39	0.13	60.2	52.49	-0.019 \pm 0.348
<i>Bact.</i> /16S DNA Post	143.41 \pm 90.73	91.83 \pm 124.76	0.07	0.47	0.20	54.7	88.18	0.032 \pm 0.347
<i>Bact.</i> /16S DNA Δ	47.32 \pm 95.74	17.74 \pm 138.78	0.35	0.25	0.69*	-	76.88	0.052 \pm 0.378

854 * = significant correlation ($p \leq 0.05$); ** = significant correlation ($p \leq 0.01$)

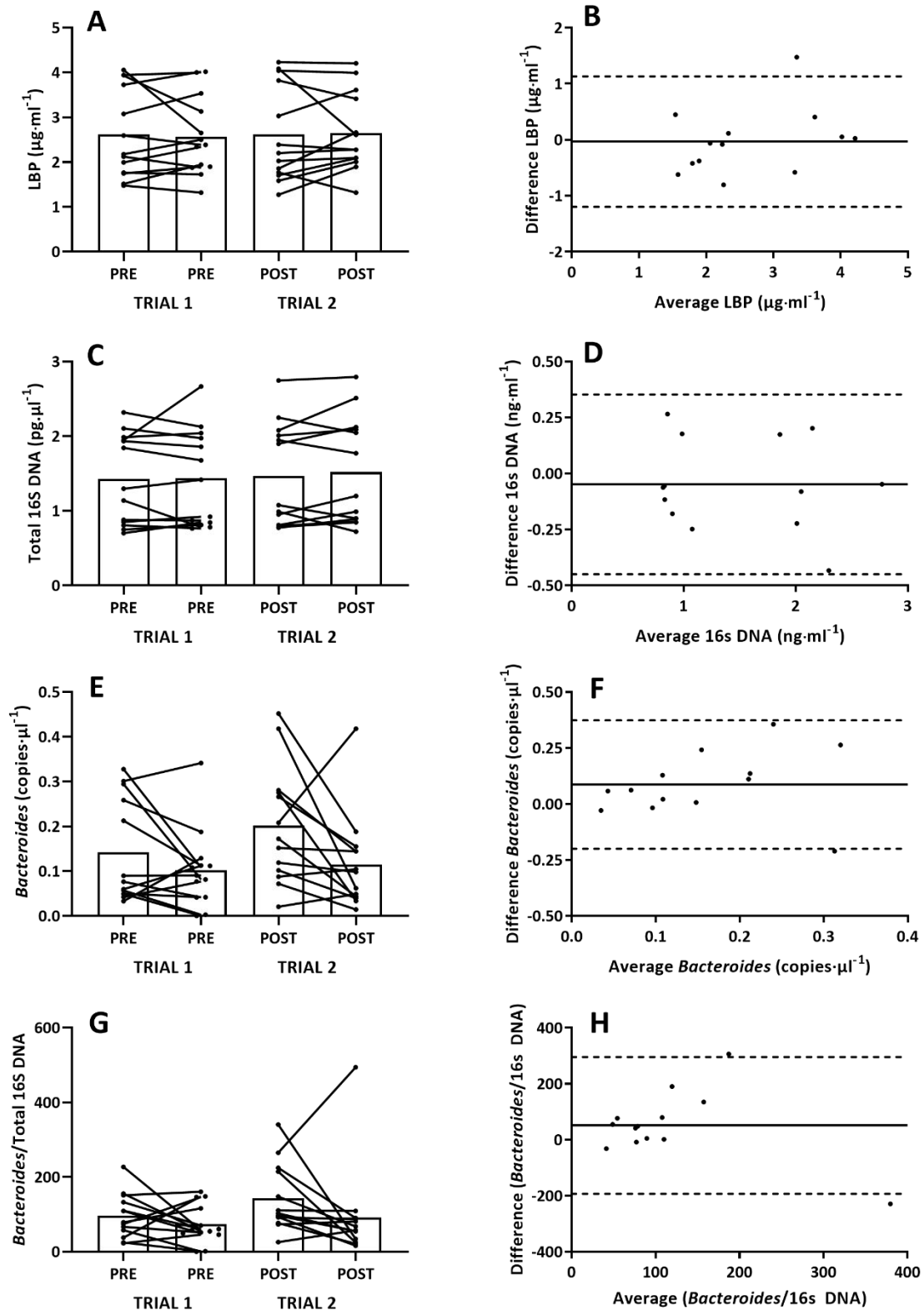
855

856

857

858

859



860

861 **Figure 4.** GI MT responses to EHST trial 1 and trial 2: (A) = LBP; (C) total 16S bacterial DNA
 862 (n=13); (E) = *Bacteroides* DNA; and (G) = *Bacteroides*/total 16s bacterial DNA (n=13).
 863 Significant overall effect of trial (+ $p \leq 0.05$). Bland-Altman mean bias and 95% LoA between
 864 post EHST trial 1 and trial 2: (B) = LBP; (D) = total 16S bacterial DNA (n=13); (F) *Bacteroides*
 865 DNA (n=13); and (H) = *Bacteroides*/total 16s bacterial DNA (n=13).

