

1 **Energetic costs of mounting an immune response in a coral reef damselfish (*Pomacentrus***  
2 ***amboinensis*)**

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## 26 **Abstract**

27 While immune responses can be energetically costly, quantifying these costs is challenging. We  
28 tested the metabolic costs of immune activation in damselfish (*Pomacentrus amboinensis*  
29 Bleeker, 1868) following a mass-adjusted injection of lipopolysaccharide (LPS) endotoxin.  
30 Fish were divided into eight treatments: two controls (handling and saline injection) and six  
31 LPS groups with concentrations ranging from 3 to 100mg kg<sup>-1</sup>. We used intermittent flow  
32 respirometry to measure differences in oxygen uptake ( $\Delta\dot{M}O_2$ ) 20h before versus 20h after  
33 LPS injection and changes in metabolic traits (lowest, routine and peak metabolic rates) as  
34 proxies of the aerobic costs of metabolism. Spleen somatic index (SSI) and gene expression in  
35 spleens were measured to assess immune activation. We found no difference in metabolic traits  
36 or SSI but observed different non-linear patterns of  $\Delta\dot{M}O_2$  in fish exposed to 50 and 100mg  
37 kg<sup>-1</sup> LPS compared to lower doses and controls. Fish exposed to high doses of LPS also had  
38 lower residual aerobic scope compared to controls and lower LPS doses. Fish exposed to doses  
39 of 3, 50, and 100mg kg<sup>-1</sup> showed altered gene expression compared to the handling control.  
40 Overall, our results suggest that immune activation has measurable effects on metabolic traits  
41 that are both dose and time-dependent.

42

## 43 **Keywords**

44 *Pomacentrus amboinensis*, metabolism, gene expression, spleen somatic index, generalized  
45 additive mixed model, damselfish

46

## 47 **Introduction**

48 Infection with macroparasites such as helminths and arthropods and/or microparasites such as  
49 bacteria, protozoa, fungi or viruses can be costly for hosts and decreases host fitness (Poulin  
50 and Morand, 2000; Schimdt-Hempel 2003; Goater et al. 2014). In animals, infection is often

51 linked to the activation of the host's immune system. Upon recognition of a disease-causing  
52 agent, the host's innate immune system, which is responsible for non-specific immune  
53 responses, activated along with a systemic inflammatory response and activation of the  
54 hypothalamic-pituitary-adrenal axis; together, these form part of the acute phase response  
55 (APR) to infection (Magnadottir 2010). The APR triggers a series of cellular reactions including  
56 hormone and cytokine secretion; activation of lymphocyte differentiation (B-cells) and  
57 proliferation (B cells and T-cells) as well as secretion of plasma protein in the liver (Møller and  
58 Erritzoe 2002; Reyes-Cerpa et al. 2012) resulting in an increase in metabolic demands. For  
59 example, mosquitofish (*Gambusia holbrooki* Girard, 1959) can experience a significant  
60 reduction in body mass while expressing an immune response, suggesting the use of energy  
61 reserves to sustain an increase in metabolic demands due to immune activation (Bonneaud et  
62 al. 2016). The metabolic costs of immune activation can also be quantified by estimating  
63 changes in aerobic metabolism (Powell et al. 2005; Methling et al. 2019). For instance,  
64 immune-challenged zebra finches (*Taeniopygia guttata* Vieillot, 1817) increase their resting  
65 metabolic rate in response to lipopolysaccharide (LPS) injection (Burness et al. 2010). These  
66 results suggest a trade-off between energy allocation to immune function and other fitness-  
67 enhancing activities (Sheldon and Verhulst 1996; Mills et al. 2010).

68  
69 Despite some evidence for this trade-off, several studies fail to demonstrate an energetic cost  
70 associated with host immune activation. For example, side-blotched lizard (*Uta stansburiana*  
71 Baird and Girard, 1852), and wild white-footed mice (*Peromyscus leucopus* (Rafinesque, 1818)  
72 showed no changes in their metabolic rates in the days, after being exposed to an immune  
73 stimulant (Derting and Compton 2003; Smith et al. 2017).

74 Several non-mutually exclusive factors could explain the inconsistencies in the energetic costs  
75 of mounting an immune response observed among studies and species. For example the

76 methods used to experimentally induce non-pathogenic immune reactions in hosts vary among  
77 studies. Studies commonly use phytohaemagglutinin (PHA) or LPS to trigger a host immune  
78 response. Yet, each method yields differences in terms of the immune reaction it triggers.  
79 Injection of PHA – a non-pathogenic antigen from a plant lectin – induces swelling (Martin et  
80 al. 2006). Alternatively, LPS – an endotoxin present in gram-negative bacteria – provokes an  
81 inflammatory response characterized by “sickness behaviour”, such as fever, lethargy, and  
82 reduced physiological function, often leading to a decrease in food intake and/or reproductive  
83 success (Bonneaud et al. 2003; Melhado et al. 2020; Lopes et al. 2021). As such, the energetic  
84 costs of different forms of immune stimulation may differ. For example, fish-eating myotis bats  
85 (*Myotis vivesi* Ménégaux, 1901), injected with 50  $\mu$ L of PHA show no change in body mass or  
86 routine metabolic rate in the 54 hours post-injection (Otálora-Ardila et al. 2016). In contrast,  
87 individuals injected with 50  $\mu$ L of LPS experience a decrease in body mass and a drastic  
88 increase in their routine metabolic rate (up by 140-185%) within 11 hours post-injection  
89 (Otálora-Ardila et al. 2016). This discrepancy suggests a higher energetic cost associated with  
90 the physiological reactions induced by LPS compared with PHA.

91  
92 Another factor that may contribute to discrepancies among studies examining the energetic  
93 costs of the immune response is that immune responses are species-specific, regardless of the  
94 immune stimulant used to induce immune activation (Berczi et al. 1966; Martin, Weil, and  
95 Nelson 2007). For instance, mammals and birds are extremely sensitive to endotoxins and react  
96 strongly to LPS injection (Berczi et al. 1966; Hasselquist and Nilsson 2012). For guinea pigs,  
97 (*Cavia porcellus* (Linnaeus,1758)), an injection of 10mg kg<sup>-1</sup> of LPS can lead to lethal  
98 endotoxin shock. In contrast, some teleost fishes are known to be remarkably tolerant to LPS,  
99 with doses of 50mg kg<sup>-1</sup> of LPS failing to induce sickness symptoms in coho salmon  
100 (*Oncorhynchus kisutch* (Walbaum,1792)) or rainbow trout (*Salmo gairdneri* (Walbaum,1792))

101 (Berczi et al. 1966). It is noteworthy that most studies exploring the physiological costs of  
102 immune activation focus on birds and mammals (Hasselquist and Nelson 2012). Similar studies  
103 in reptiles, amphibians or fishes remain scarce despite known differences among taxa and  
104 species (Ackerman et al. 2000; Mackenzie et al. 2006; Zanuzzo et al. 2015; Bonneaud et al.  
105 2016; Bennoit and Craig 2020). Finally, experiments often study the effects of a single dose, or  
106 limited doses, of a given immune stimulant without the necessary controls (i.e., a handling  
107 control and a saline injection control), making it difficult to understand the dose-response of  
108 hosts to endotoxin administration. Individual within species variation in metabolic rates are also  
109 not typically considered in study designs despite minimum metabolic rates sometimes varying  
110 greatly (2-3-fold) among individuals within the same population (Burton et al. 2011). This large  
111 among-individual variation in metabolic rates makes it more difficult to detect differences  
112 among treatments.

113  
114 Here, we examine the metabolic costs of mounting an immune response in the coral reef  
115 damselfish, (*Pomacentrus amboinensis* Bleeker, 1868), by measuring oxygen uptake in  
116 individuals pre- and post-immune stimulation, across a range of LPS doses. We also assess  
117 differences in spleen mass and gene expression across treatments to assess the effectiveness of  
118 an endotoxin injection in stimulating an immune response. The spleen is a major lymphoid  
119 organ in fish and plays a critical role in clearing infection with spleen size often used as an  
120 indicator of an immune response in fish (Lefebvre et al. 2004; Rauta et al. 2012). *Pomacentrus*  
121 *amboinensis* are well-studied in terms of their physiology and behaviour. This species was  
122 previously found to mount an immune response following injection of 50mg kg<sup>-1</sup> of LPS as  
123 demonstrated by an alteration in gene expression associated with immune activation (Binning  
124 et al. 2018). Thus, our objectives were to (1) assess whether individuals respond physiologically  
125 to an immune challenge with LPS through changes in oxygen uptake, spleen size and alteration

126 in immune gene expression; and if so, (2) determine whether the host's immune response varies  
127 across LPS doses. We hypothesized that higher doses of LPS would cause the most significant  
128 difference in oxygen uptake pre- and post-injection. Additionally, we hypothesized that fish  
129 exposed to LPS would have a higher spleen somatic index (SSI) and an altered immune gene  
130 expression compared to both control groups (saline injection and handling control) if the  
131 injection triggers an immune response.

132

## 133 **Methods**

### 134 **Fish collection and husbandry**

135 This study was conducted at the Lizard Island Research Station (LIRS), on the northern Great  
136 Barrier Reef, Australia (14° 40' S; 145° 28' E) within a two month period between July and  
137 September 2015. Field collections and experiments were conducted under permits from the  
138 Great Barrier Reef Marine Park Authority (G14/37048.1) with approval from the Queensland  
139 Government (DAFF) Animal Ethics Committee (CA 2015/07/878). A total of forty-eight  
140 females *P. amboinensis* were collected on July 25<sup>th</sup> and August 13<sup>th</sup> (non reproductive period)  
141 using hand nets and monofilament barrier nets (10 mm stretch mesh) from continuous reef sites  
142 in the Lizard Island lagoon (3-5m depth) by two divers on SCUBA.

143 Fish were transported in individual water-filled plastic Ziploc bags to the aquarium facilities at  
144 LIRS within 90 minutes of capture. To remove unencysted ecto- and endoparasites, fish were  
145 treated with a Praziquantel solution (1 tablet dissolved in 3ml ethanol, mixed in 20L water) for  
146 90 minutes, followed by 60 seconds in a freshwater bath. Each individual was then measured  
147 (mass (M), , standard length (SL)) and transferred to an individual white holding aquarium  
148 (28.0 W X 38.0 L X 18.0 H) with an opaque PVC tube (6.0 cm L X 5.5 cm diameter) for shelter.  
149 Fish were fed Nutrafin Tropical fish flakes twice a day to satiation.

150

**151    Respirometry measurements – estimate of metabolic traits**

152    Metabolic rates for all forty-eight individuals (SL:  $52.2 \pm 5.7$  mm; M:  $5.9 \pm 2.1$  g; mean  $\pm$  S.D.)  
153    were estimated using automated intermittent-flow respirometry in resting chambers  
154    (AquaResp, [www.aquaresp.com](http://www.aquaresp.com)) fitted with a recirculation loop to allow proper mixing (Roche  
155    et al. 2013; Clark et al. 2013; Rosewarne et al. 2016; Killen et al. 2021). We measured an  
156    individual's rate of oxygen uptake ( $\dot{M}O_2$ ; mg O<sub>2</sub> h<sup>-1</sup>) as a proxy for aerobic metabolic rate. Full  
157    details of the respirometry setup are given in supplemental material (Table S1), following the  
158    reporting guidelines described in Killen et al. (2021). Respirometry trials were carried out in  
159    custom made cylindrical 0.5 L resting chambers (volume includes recirculation tubes)  
160    submerged in U.V. sterilized, temperature-controlled and aerated water baths with water  
161    temperature maintained at  $24 \pm 0.3^\circ\text{C}$  (actual variation) using a TECO TC 20 water chiller /  
162    heater system. Chambers were opaque with only a small viewing window located on the top to  
163    minimize visual disturbance to the fish. Dissolved oxygen concentration was recorded with a  
164    four channel FireSting O<sub>2</sub> Optical Oxygen Meter (Pyroscience, Aachen, Germany). Every six  
165    days, the respirometry chambers were disassembled, rinsed in freshwater and left to dry for 8h  
166    to reduce background oxygen consumption due to bacteria. Changes in water oxygen tension  
167    (PwO<sub>2</sub>) due to fish respiration were continuously monitored at 1 Hz with the software  
168    AquaResp.

169    All respirometry trials began between 17.00 and 19.00 with an exhaustive chase protocol  
170    followed by air exposure to estimate the maximum metabolic rate (hereafter: MMR), the upper  
171    limit of a fish capacity to aerobically metabolize oxygen under a given environmental condition  
172    (Norin and Clark 2016; Killen et al. 2017). To estimate MMR, fish were individually placed in  
173    a circular bucket (30 cm diameter) and chased continuously by hand for three minutes, held out  
174    of the water for one minute and after that, then immediately placed in the respirometry chambers  
175    (Roche et al. 2013). The first measurement period commenced when the respirometer was

176 sealed (maximum 15 seconds). For this first measurement, the chamber remained closed for 15  
177 minutes to have continuous measurements to calculate  $\dot{M}O_{2\max}$ . Afterwards, ten minutes loops  
178 with a 240s flush, 60s wait and 300s measure cycle were taken to measure  $\dot{M}O_2$ . This continuous  
179 periodic monitoring of  $\dot{M}O_2$  in the respirometer allowed estimation of individual standard  
180 metabolic rate, the minimum energy required in a post-absorptive state to maintain homeostasis,  
181 overnight (hereafter: SMR) (Chabot et al. 2016). We calculated SMR as the 20<sup>th</sup> percentile of  
182 measures taken overnight, excluding the first five hours after transfer to the respirometer to  
183 minimise the effects of stress on estimates of SMR (Chabot et al. 2016). From these measures,  
184 we were able to determine individual's aerobic scope (hereafter: AS), the total amount of energy  
185 available to allocate to physiological systems including the immune system, calculated as  
186 MMR-SMR (Derting and Compton 2003; Claireaux and Lefrançois 2007; Chabot et al. 2016b).  
187 We fasted the fish for for 24h prior to the respirometry trials to avoid increases in  $\dot{M}O_2$  caused  
188 by a digestive state (Jobling 1981, Chabot et al. 2016b). The body mass of each fish was  
189 determined to the nearest 0.1g before the start of each trial. Two loops were run without a fish  
190 in the respirometer to measure initial and final background rates of respiration. Background  
191 respiration was accounted for by subtracting oxygen consumption in an empty chamber from  
192 concurrent  $\dot{M}O_2$  values taken in chambers containing fish (Rodgers et al. 2016). Artificial  
193 lighting was turned off, and only natural lighting was available during the rest of the trial.

194

### 195 **Respirometry - Immune challenge**

196 After approximately 20h in the respirometer, fish were given an immune challenge to estimate  
197 the metabolic costs of mounting an innate immune response. We used lipopolysaccharide (LPS;  
198 Sigma-Aldrich L2880, serotype 055:B5), a component of the *E*-coli cell wall, to elicit an  
199 immune response in fish. Fish were haphazardly assigned to one of eight treatments (6 fish per  
200 treatment) such that the mean fish mass was similar among treatment groups: 1) handling



201 control, 2) saline injection (0.9 % saline control), 3) 3mg kg<sup>-1</sup> LPS, 4) 5mg kg<sup>-1</sup> LPS, 5) 10mg  
202 kg<sup>-1</sup> LPS, 6) 30mg kg<sup>-1</sup> LPS, 7) 50mg kg<sup>-1</sup> LPS, 8) 100mg kg<sup>-1</sup> LPS. Fish were removed from  
203 the respirometers and injected intraperitoneally (i.e. abdominal cavity) using heparinized 29  
204 gauge insulin syringes (heparin concentration: 28 mg/ml<sup>-1</sup>) through a plastic bag partially filled  
205 with seawater to avoid stress caused by air exposure and to protect the fish from surface  
206 abrasions during handling. The volume of the injection was mass-adjusted for each fish  
207 resulting in a injected volume standardized across concentrations and fish. Following injection,  
208 the test fish was immediately placed back in the respirometry chamber and left for a further 20h  
209 to measure  $\dot{M}O_2$  post-injection. These 20h of measurements post-injection were taken over the  
210 same time period as the period of measurements pre-injection. With post-injection  $\dot{M}O_2$  values,  
211 we measured the peak, lowest, and routine metabolic rate to compare difference in metabolic  
212 traits among groups. We defined the lowest metabolic rate as the 20<sup>th</sup> percentile of  $\dot{M}O_2$  values  
213 post-injection and calculated the routine metabolic rate as the average  $\dot{M}O_2$  values post-  
214 injection. We selected the peak value of  $\dot{M}O_2$  as a proxy for the peak metabolic rate post-  
215 injection. Furthermore, we calculated the percentage of aerobic scope available for the fish  
216 across time within each treatment by subtracting  $\dot{M}O_2$  values post-injection from the aerobic  
217 scope calculated pre-injection. We also estimated the total integrated post-injection oxygen  
218 uptake for each individual by calculating the area under the curve (AUC) of  $\dot{M}O_2$  versus time  
219 post-injection. AUC was calculated as integral of  $\dot{M}O_2$  post-injection using the trapezoid rule  
220 (Chabot et al. 2016b). A positive AUC suggests an overall increase in energy demands  
221 following injection, whereas a negative AUC suggests a decrease in energy demands following  
222 injection. Post-respirometry experiments, the fish were returned to their holding tanks for a  
223 week and fed twice per day.

224

## 225 **Tissue collection**

226 To ensure sufficient time for the systemic and the adaptive immune response to establish a  
227 seven day period was observed between the LPS injection and the tissue collection. This time  
228 period is sufficient for changes in gene expression to be detected in this species (Binning et al.  
229 2018). Fish were euthanized in an overdose of AQUI-S<sup>®</sup> solution (AQUI-S New-Zeland Ltd).  
230 Fish weight and length were measured again. The spleen was then dissected out and weighed.  
231 Spleen tissue was preserved in RNAlater<sup>®</sup> (R0901 Sigma-Aldrich), frozen, and stored at -20°C  
232 until they were shipped to the University of Greenwich for PCR analysis.

### 233 **Spleen tissue and gene expression analysis**

234 The spleen somatic index (SSI) was calculated as the proportion of the spleen in relation to the  
235 total fish wet mass. The protocol used for immune gene analysis is described in Binning et al.  
236 (2018). Briefly, RNA was extracted from the spleen with ReliaPrep<sup>™</sup> RNA Cell Miniprep  
237 System (Promega). RNA concentration was determined using a nanodrop 2000 (Thermo  
238 Scientific). The samples were then normalized to a common concentration of 50ng  $\mu\text{l}^{-1}$  using  
239 nuclease-free water (Promega). 425 ng RNA were treated with DNase (Quanta PerfeCta DNase  
240 I Kit (QuantaBio)) and 212 ng of the treated RNA were transcribed using Quanta qScript cDNA  
241 Synthesis Kit (QuantaBio) with a total volume of 10  $\mu\text{l}$  following manufacturer instructions. A  
242 1:10 dilution with Nuclease-free water (Promega) was performed and samples were stored at  
243 4°C overnight. Gene expression analysis was performed the next day. Immunological response  
244 to LPS challenge was quantified using gene polydom/svep1 (Binning et al. 2018). To ensure  
245 specificity melting curves of the qPCR products were checked between 60 and 90°C with 1°C  
246 intervals. Gene expression data (ct-values) were normalised against the house keeping gene  
247 12s, to give dct. One fish in the LPS30 was not included in the SSI and gene expression analysis  
248 due to an inability to retrieve the spleen. Two fish in the handling control, one fish in the saline  
249 group, one fish exposed to 3mg  $\text{kg}^{-1}$  of LPS, one fish exposed to 5mg  $\text{kg}^{-1}$  of LPS, and one fish

250 exposed to 100mg kg<sup>-1</sup> of LPS had a spleen too small to perform RNA extraction preventing  
251 the analysis of immune gene expression.

252

### 253 **Statistical analysis**

254 All statistical analyses and data visualization were performed using R v.4.0.3 (R Core Team,  
255 2016). We examined the relationship between metabolic traits (lowest, routine, and peak  
256 metabolic rates) and treatment with a linear model (LM). The lowest, routine, and peak  
257 metabolic rates were used as response variables, with treatment, fish mass and their interaction  
258 as predictors. This allowed us to assess group differences in metabolic traits while accounting  
259 for mass as a covariate. Body mass and metabolic rates were both log<sub>10</sub>-transformed to  
260 normalize the data. Percentage of residual aerobic scope available at each time step was  
261 calculated for each fish by estimating individual AS pre-injection, from which we subtracted  
262 the AS estimated post-injection. Then, we compared the average available aerobic scope among  
263 treatments. Values above 100% indicate that oxygen uptake post-injection was lower than pre-  
264 injection whilst values below 100% indicate an increase in oxygen uptake post-injection. To  
265 observe changes in the  $\dot{M}O_2$  patterns among treatments through time, we measured the  
266 difference in  $\dot{M}O_2$  ( $\Delta\dot{M}O_2$ ). We calculated  $\Delta\dot{M}O_2$  by subtracting the  $\dot{M}O_2$  pre-injection  
267 from the  $\dot{M}O_2$  post-injection taken at the exact same time. For example, fish  $\dot{M}O_2$  value taken  
268 pre-injection at 7:00:00 pm were subtracted to its  $\dot{M}O_2$  value taken post-injection at 7:00:00 pm  
269 the following day. By doing so, we aimed to determine the change in an individual's oxygen  
270 uptake relative to its own baseline over time. The first hour of  $\dot{M}O_2$  measurement pre- and post-  
271 injection were excluded from  $\dot{M}O_2$  measurement to exclude any change in oxygen uptake linked  
272 to either, the MMR measurement pre-injection or the handling/injection post-injection (Fig.S1).  
273 To capture the nonlinear pattern observed for changes in  $\dot{M}O_2$  over time among treatments, we  
274 used a generalized additive mixed model (GAMM; function "gam" in R package "mgcv", Wood

275 and Wood 2015). GAMMs are an extension of generalized linear mixed models using smooth  
276 terms alongside parametric terms. Smooth terms are a set of non-linear functions (i.e., basis  
277 functions), which are multiplied each by a coefficient estimated from the data and summed.  
278 Using the `gam.check()` function, we determined the number of basis functions needed for the  
279 smooth term "time" in our model. The `k` parameter (i.e., number of basis functions) was set to  
280 30 basis functions for the smooth term "time". Our model contained the parametric terms  
281 " $\Delta\dot{M}O_2$ " (response variable), "treatment" (explanatory variable), "mass" (explanatory  
282 variable), and the smooth term "time" (explanatory variable) representing patterns related to the  
283 non-linear change in  $\Delta\dot{M}O_2$  over time (in minutes). We included the random effect 'fish ID'  
284 in the model to allow smoothers to vary in intercept and slope. We used restricted maximum  
285 likelihood (REML) for parameter estimation. Following diagnostic plots assessment, we used  
286 the scaled  $t$  family in place of Gaussian as our data appeared heavy tailed (Wood 2017).  
287 We used estimation statistics to determine differences in the spleen somatic index (SSI),  
288 immune gene expression, group mean, standard deviations and treatment effect sizes with 95%  
289 confidence intervals are presented using Gardner-Altman estimation plots (*R package "dabestr"*,  
290 Ho et al. 2019).  
291 We used a Bayesian multivariate model (*R package MCMCglmm*; Hadfield et al. 2010) with a  
292 Gaussian error distribution to examine covariation among three immune activation parameters:  
293 SSI, immune gene expression, and the AUC. Treatment and fish mass were included in the  
294 model as fixed effects. Predictor variables were mean-centered (mean = 0) and standardised to  
295 one standard deviation. Visual inspection of diagnostic plots (parameters traces) indicated chain  
296 convergence. All statistical analyses and data visualization were performed using R v. 4.0.3 (R  
297 Core Team, 2016).

298

299 **Results**

### 300 *Change in metabolic traits post-injection*

301 We found no significant difference in the peak metabolic rate attained by fish across groups  
302 post-injection (LM,  $F_{(7,32)} = 0.93$ ,  $p = 0.491$ ) (Fig.1 (a)). There were also no differences in the  
303 lowest metabolic rate (LM,  $F_{(7,32)} = 1.130$ ,  $p = 0.369$ ) and routine metabolic rate (LM,  $F_{(7,32)} =$   
304  $0.76$ ,  $p = 0.62$ ) (Fig.1 (b-c)). The treatment-mass interaction was not significant for any of the  
305 metabolic traits (peak metabolic rate:  $F_{(7,32)} = 0.802$ ,  $p = 0.591$ ; lowest metabolic rate:  $F_{(7,32)} =$   
306  $1.120$ ,  $p = 0.374$ ; routine metabolic rate:  $F_{(7,32)} = 0.605$ ,  $p = 0.746$ ). Fish mass had a significant  
307 positive effect on fish routine metabolic rate ( $F_{(7,32)} = 7.150$ ,  $p = 0.011$ ) and was marginally  
308 non-significant for both the peak and lowest fish metabolic rate (repectively:  $F_{(7,32)} = 4.053$ ,  $p$   
309  $= 0.052$  ;  $F_{(7,32)} = 4.088$ ,  $p = 0.051$ ). Fish exposed to  $50\text{mg kg}^{-1}$  and  $100\text{mg kg}^{-1}$  had, on average,  
310 the lowest aerobic scope post-injection with, respectively, 92.57 % and 95.96% of their aerobic  
311 scope available (Fig.2 (g-h)). In comparison, both control groups approached nearly 100% of  
312 their aerobic scope (99,30% for the handling control and 100.26% for the saline) post-injection  
313 (Fig.2 (a-b)). Fish injected with doses of  $30\text{mg kg}^{-1}$  and lower, all had greater than 95% of their  
314 aerobic scope available post-injection (dose of  $3\text{mg kg}^{-1} = 96.35\%$ ; dose of  $5\text{mg kg}^{-1} = 96,84\%$ ;  
315 dose of  $10\text{mg kg}^{-1} = 98.09\%$ ; dose of  $30\text{mg kg}^{-1} = 102.02\%$ ) (Fig.2 (c-f)). The peak level of  
316  $\dot{M}O_2$  post injection occurred between 11 and 12h for the control groups and the LPS doses  
317 ranging from  $3\text{mg kg}^{-1}$  to  $30\text{mg kg}^{-1}$  (Fig.2 (a-f)). For fish exposed to  $50\text{mg kg}^{-1}$  and  $100\text{mg}$   
318  $\text{kg}^{-1}$ , the peak  $\dot{M}O_2$  occurred between 8 and 9 h post injection (Fig.2 (g-h)). Finally, control  
319 groups and LPS injected fish showed no difference in AUC (Table 2, Fig.3).

320

### 321 *Changes in oxygen uptake overtime*

322 In our GAMM model, the estimated degrees of freedom ranged from 12.30 to 23.22 suggesting  
323 a highly nonlinear relationship between the oxygen uptake through time and LPS doses (Table  
324 1).  $\Delta\dot{M}O_2$  tended to vary through time in fish in both control groups, and in the LPS-

325 injection treatment groups with dose of 3, 5, 10, and 30mg kg<sup>-1</sup> (all  $p < 0.05$ ; Table 1, Fig.4  
326 (a-f)) but not in LPS treatment groups with doses of 50, and 100mg kg<sup>-1</sup> ( LPS50:  $p = 0.53$  and  
327 LPS100:  $p = 0.65$ ; Table 1, Fig.4 (g-h)). In other words, there were changes in  $\dot{M}O_2$  uptake in  
328 fish exposed to 3, 5, and 10mg kg<sup>-1</sup> LPS: fish in these groups initially had lower  $\dot{M}O_2$  post-  
329 injection compared to pre-injection, followed by oxygen uptake above pre-injection values 10  
330 hours post-injection, which occurred around sunrise (time, Fig.4 ((c, e)). This pattern was also  
331 observed in fish exposed to a saline injection (Fig.4 (b)). The handling control group and fish  
332 injected with an LPS dose of 30mg kg<sup>-1</sup> showed a similar pattern, exhibiting a steady increase  
333 in  $\dot{M}O_2$  over time (Fig.4 (a, f)). Interestingly, fish exposed to the two highest LPS doses (50  
334 and 100mg kg<sup>-1</sup>) had patterns of  $\Delta\dot{M}O_2$  through time that did not differ from 0, suggesting a  
335 relatively constant oxygen uptake both pre and post-injection.

336

### 337 *Changes in SSI and gene expression*

338 We found no difference in SSI between the two control groups (handling and saline injection)  
339 and between the handling control (the contrast group) and the different LPS doses (Table 2,  
340 Fig.5). There was no difference in immune gene expression between the handling and saline  
341 control groups (Table 2, Fig.6). However, we observed a dose-dependent effect of LPS injection  
342 on immune gene expression, with gene expression being altered in fish injected with the lowest  
343 dose of LPS (3mg kg<sup>-1</sup>) and the two highest LPS doses (50mg kg<sup>-1</sup> and 100mg kg<sup>-1</sup>) compared  
344 to the handling control group (Table 2, Fig.6). There was also an indication of a similar pattern  
345 for fish injected with 30mg kg<sup>-1</sup> of LPS (mean difference in gene expression compared to the  
346 handling control: 1.18 dct, 95% *CI*: -0.0031– 3.04).

347

### 348 *Covariation between SSI, gene expression, and metabolism*

349 We found no evidence of a strong association between SSI, gene expression, and metabolism  
350 following immune stimulation (AUC–SSI  $r = -0.22$ ; AUC–gene expression  $r = -0.26$ ; SSI–gene  
351 expression  $r = 0.28$ ; Fig.S2). A more detailed examination of correlations within individual  
352 treatment groups supports this finding (Fig.S3 and Fig.S4).

353

## 354 Discussion

355 Our objectives were to determine if immune stimulated coral reef fish showed difference in  
356 metabolic traits and oxygen uptake over time following immune activation and if these  
357 differences are dose dependent. We observed no significant difference in metabolic traits post-  
358 injection across treatments. However, we found that fish exposed to LPS doses of 3mg kg<sup>-1</sup>,  
359 5mg kg<sup>-1</sup>, 10mg kg<sup>-1</sup> and 30mg kg<sup>-1</sup> as well as fish in both control treatments had a  $\Delta\dot{M}O_2$   
360 through time relationships that differed significantly from zero. In most cases,  $\Delta\dot{M}O_2$  was  
361 lower in the first 10 hours post-injection (night time) followed by a sharp increase in  $\Delta\dot{M}O_2$   
362 around dawn (day time), which remained elevated for the rest of the day. Fish exposed to 50mg  
363 kg<sup>-1</sup> and 100mg kg<sup>-1</sup> LPS did not follow this trend, and instead tended to have similar  $\Delta\dot{M}O_2$   
364 values both overnight and in the daytime. Fish exposed to these doses of LPS also showed a  
365 lower residual aerobic scope on average compared to both controls and lower doses of LPS.  
366 We also observed altered immune gene expression in fish injected with the lowest dose (3mg  
367 kg<sup>-1</sup>) as well as the two highest doses (50mg kg<sup>-1</sup> and 100mg kg<sup>-1</sup>) compared with fish in the  
368 handling control. Fish SSI did not differ across treatments. Overall, these results highlight  
369 complex responses to immune system activation that depend on the dose of LPS administered.

370

371 Immune stimulation and metabolic traits

372 Our results are consistent with previous findings observed in ectotherms that showed no  
373 difference in metabolic traits following exposure to LPS (Smith et al. 2017; Stahlschmidt and

374 Glass 2020). We hypothesized that triggering an immune response would result in increased  
375 oxygen consumption as a result of upregulation in genes associated with immune defense.  
376 However, it is possible that an individual may maintain similar metabolic demands and instead  
377 reallocate energy away from other activities such as growth and reproduction in order to fuel  
378 an immune response (Uller et al. 2006; Lind et al. 2020). Interestingly, across all treatments the  
379 remaining aerobic scope was above 90% of that measured pre-injection (Fig.2), suggesting that  
380 mounting a mild immune response does not impose a large constraint on the capacity of an  
381 individual to perform other aerobic activities simultaneously. Indeed, the overall integrated  
382 energy use as measured by the AUC showed no difference among treatments (Table 2; Fig.3).  
383 Nonetheless, residual aerobic scope was lowest for the two highest doses of LPS. Indeed,  
384 Binning et al. (2018) found that 50mg kg<sup>-1</sup> of LPS induced an immune response in Ambon  
385 damselfish. Whether a 5% or 7% reduction in AS represents a significant cost to fish in natural  
386 settings, especially when combined with other biotic or abiotic stressors, remains to be  
387 explored.

388  
389 An acute inflammatory response from LPS injection can affect host metabolic traits in the  
390 minutes to hours post-injection (Novoa et al. 2009; Bonneaud et al. 2016; Otálora-Ardila et al.  
391 2016). In the present study doses of 3mg kg<sup>-1</sup>, 5mg kg<sup>-1</sup>, and 10mg kg<sup>-1</sup> elicited  $\dot{M}O_2$  responses  
392 similar to control treatments. Indeed, oxygen uptake was lower overnight in fish from these  
393 groups during the second day in the respirometers, which may be indicative of habituation to  
394 the experimental set-up (Fig.4 (a-f)). Delta $\dot{M}O_2$  values in these treatments also trended above 0  
395 around 10 hours post-injection, at the onset of dawn, suggesting increased diurnal activity in  
396 fish on the second day of experiments. Conversely, fish from the highest-dose treatments (50mg  
397 kg<sup>-1</sup> and 100mg kg<sup>-1</sup>) tend to have opposite delta $\dot{M}O_2$  trends. Delta $\dot{M}O_2$  values were constantly  
398 above 0 within the first 10 hours of the trial before decreasing below or near 0 after 10 hours



399 post-injection. This suggests that these individuals do not show decreased  $\dot{M}O_2$  overnight nor  
400 increased diurnal activity during the daytime which may be linked to an immune activation  
401 beginning within the first 2-4 hours post injection. Otálora-Ardila et al. (2016) observed that  
402 time was a key component to consider due to its interacting effect with LPS treatment on the  
403 routine metabolic rate of fish-eating bat. Indeed, our results demonstrate that temporal trends  
404 in oxygen uptake associated with the magnitude of the immune response can be present despite  
405 no significant mean differences in metabolic traits averaged across time. Thus, we believe that  
406 constant measures of oxygen uptake over time post immune stimulation is necessary to capture  
407 these complex non-linear dynamics.

408  
409 The influence of circadian rhythms on fish metabolism is well-documented, and several studies  
410 show a tight relationship between variation in oxygen uptake and the light-dark cycle (Svendsen  
411 et al. 2014; McKenzie et al. 2015). For instance, lake sturgeon, (*Acipenser fulvescens*  
412 Rafinesque, 1817), exhibit variation in  $\dot{M}O_2$  over 24 hours, with a peak at dusk, and at dawn,  
413 depending on the light cycle the fish are exposed to (Svendsen et al. 2014). The interactions  
414 between the circadian cycle and the immune system can lead to increased immune functions  
415 during the resting phase (Halberg et al. 1960; Scheiermann et al. 2013). In our experimental  
416 timeline, the resting phase occurred directly after the LPS injection (between 0 and 10 hours;  
417 Fig.4). We suggest that the increase in oxygen demands above 0 for 50mg kg<sup>-1</sup> and 100mg kg<sup>-1</sup>  
418 during the fish resting phase is a sign of immune activation. The opposite pattern for  $\Delta\dot{M}O_2$   
419 was observed for control groups and doses ranging from 3mg kg<sup>-1</sup> to 30mg kg<sup>-1</sup> of LPS with  
420 fluctuations in  $\Delta\dot{M}O_2$  below 0 values during the night time and consistently above it after  
421 sunrise likely linked to natural biological rhythms of the fish rather than reflecting an immune  
422 activation.

423

**424 Immune stimulation effect on the spleen somatic index and gene expression**

425 In fish, the spleen is a crucial secondary lymphoid organ with a central role in systemic  
426 immunity due to its functions in coordinating innate and adaptative immune responses (Mebius  
427 and Kraal 2005; Uribe et al. 2011). The immunological function of the spleen in fighting off  
428 infection results in changes in its structure and morphology (Bjørngen and Koppang 2021).  
429 Spleen enlargement as an adaptative response to immune activation has been observed across  
430 fish species, including damselfish (Lefebvre et al. 2004; Seppänen et al. 2009; Binning et al.  
431 2018). We did not find evidence for spleen enlargement in the present study. This absence  
432 suggests that the doses of endotoxins may not have been sufficient to trigger an immune  
433 response, or that the timing of sampling following injection (7 days post-injection) was not  
434 appropriate to capture differences in SSI among treatments. Interestingly, the SSI values  
435 measured in the fish injected with saline and 50mg kg<sup>-1</sup> in the present study were similar to  
436 those obtained for fish in the same treatments in Binning et al. (2018). Notably, the fish used  
437 here and in Binning et al. (2018) were from the same population and collected at the same time  
438 of year as in the present study. Additionally, we had a much smaller sample size per treatment  
439 (for both saline and LPS 50mg kg<sup>-1</sup> injection  $n = 6$  in the present study vs.  $n = 16$  in Binning et  
440 al. (2018)). These factors may explain the lack of clear differences observed in SSI between our  
441 groups vs. those found in Binning et al. (2018). Spleen enlargement alone may not be an  
442 accurate measure to characterize the occurrence of an immune response given the large among-  
443 individual variation that exists among treatments (Petitjean et al. 2021)

444  
445 We found altered immune gene expression in fish injected with the two highest doses of LPS  
446 (50mg kg<sup>-1</sup> and 100mg kg<sup>-1</sup>) and with the lowest one (3mg kg<sup>-1</sup>). Responses to immune  
447 stimulants are dose and time dependent. The initial phase of the inflammatory response can be  
448 followed by either a down-regulation or an up-regulation of the immune system, which can last

449 for days (Novoa et al. 2009; Abdel-Mageid et al. 2020). A non-lethal dose of LPS can cause a  
450 temporary tolerance state to LPS which causes a suppression of genes linked with the  
451 inflammatory immune response (Foster and Medzhitov 2009). Doses closer to a lethal  
452 concentration can cause a hyperactive innate immune state to prevent an endotoxin shock  
453 caused by the accumulation of LPS within the organism (Novoa et al. 2009). In our study, the  
454 intermediate doses of LPS (5mg kg<sup>-1</sup>, 10mg kg<sup>-1</sup>, 30mg kg<sup>-1</sup>) may have caused a tolerant state,  
455 which might explain the absence of immune gene alteration. Conversely, the highest doses,  
456 which might be closer to lethal concentration, may have caused a hyperactive innate immune  
457 state. The mechanism behind the significant alteration of immune gene expression observed at  
458 the lowest dose of LPS (3mg kg<sup>-1</sup>) could result from a cross-immunity response to waterborne  
459 pathogenic agents. Over a lifetime, individuals can be exposed to the same pathogen multiple  
460 times. During the first encounter, immune cells specialized for fighting this pathogen are  
461 created and stored to provide immunity in the event of another encounter later in life. This  
462 acquired immunity gives subsequent protection against this pathogen and, through cross-  
463 immunity, can equally protect against other similar types of pathogens (Schmid-Hempel 2003;  
464 Swennes et al. 2007; Ardia et al. 2011). Previous studies observed that repeated exposure to  
465 extremely low doses of LPS through feeding enhanced the immune response of grey mullet,  
466 (*Mugil cephalus* (Linnaeus, 1758) and striped catfish, (*Pangasianodon hypophthalmus*  
467 Sauvage, 1878) (Bich Hang et al. 2016; Abdel-Mageid et al. 2020). In the present study, we  
468 used wild-caught fish. *P. amboinensis* may have been exposed in the wild to endotoxin, similar  
469 to LPS. Because of a cross-immunity, it is possible that the low dose of LPS caused a reaction  
470 similar to grey mullet and catfish, which strongly responded to a very low dose of LPS after  
471 repeated exposure.

472

473 Overall, our results suggest that there are some measurable energetic costs to mounting an  
474 immune response in Ambon damselfish. However, these costs vary in a dose- and time  
475 dependent manner, which can be difficult to capture when comparing average metabolic traits  
476 measurement without considering changes overtime. The higher doses of LPS caused an  
477 increase in  $\dot{M}O_2$  uptake above pre-injection values in the first few hours post-injection with  
478 only a marginal effect on the available aerobic scope and no effect on the overall integrated  
479 energy use. Furthermore, both higher doses and the lowest dose resulted in altered immune gene  
480 expression. To avoid underestimating the energetic costs of mounting an immune response, it  
481 appears critical for future studies to consider measuring traits at different levels of biological  
482 organisation as well as to account for variation among individuals through time in traits such  
483 as oxygen uptake rate. Additionally, the costs associated with an immune challenge can differ  
484 depending on the time of the day, making the injection timing a critical element to accurately  
485 attribute changes in energy demands to the onset of an immune response.

486

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491

#### 492 **Competing interests statement**

493 The authors declare no competing interests.

494

#### 495 **Contribution statement**

496 SAB, DGR, SSK and RB conceived the study. SAB, SC, DGR and JJM collected the data. ML,  
497 DGR and SSK analysed the data and produced the figures. ML, SAB, DGR and SSK wrote the  
498 manuscript. All authors contributed to the drafts and gave final approval for publication.

499

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#### 506 **Data availability statement**

507 The data and analytical code for this study are available to reviewers at <https://osf.io/xdsyn> and  
508 will be shared publicly at the time of publication.

509

510

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760



761 **Figure captions**

762

763 Figure 1. Relationship between mass-adjusted whole organism metabolic traits (a) highest  
764 metabolic rate, (b) lowest metabolic rate and (c) routine metabolic rate, two control groups  
765 (handling control (HC) and saline injection) and six different doses of LPS ranging from 3mg  
766 kg<sup>-1</sup> to 100mg kg<sup>-1</sup>. Each dot represents a single fish. The boxplots indicate the medians, 25<sup>th</sup>  
767 and 75<sup>th</sup> percentile.

768

769 Figure 2. Percentage of residual aerobic scope in *Pomacentrus amboinensis* damselfish across  
770 treatment post-injection. Residual aerobic scope is based on absolute value of  $\dot{M}O_2$  for each fish  
771 expressed relative to their own aerobic scope calculated pre-injection. Y-axis represents the  
772 percentage of residual aerobic scope for each fish within each treatment (a) handling control, (b)  
773 saline injection, (c) LPS3 (3mg kg<sup>-1</sup>), LPS5 (5mg kg<sup>-1</sup>), LPS10 (10mg kg<sup>-1</sup>), LPS30 (30mg kg<sup>-1</sup>)  
774 LPS50 (50mg kg<sup>-1</sup>) and LPS100 (100mg kg<sup>-1</sup>). X-axis represents the time in hours post-  
775 injection.

776

777 Figure 3. Mean differences in area under the curve of *Pomacentrus amboinensis* damselfish  
778 among LPS treatment groups (a) Area under the curve (AUC) of the handling control group  
779 (purple), the saline control group (dark blue) and the LPS injected group with LPS3 (3mg kg<sup>-1</sup>)  
780 (indigo), LPS5 (5mg kg<sup>-1</sup>) (teal), LPS10 (10mg kg<sup>-1</sup>) (forest green), LPS30 (30mg kg<sup>-1</sup>)  
781 (harlequin green), LPS50 (50mg kg<sup>-1</sup>) (mustard) and LPS100 (100mg kg<sup>-1</sup>) (yellow). Vertical  
782 bars represent the standard deviation around the group means. (b) Estimation plots indicate the  
783 mean difference in AUC between treatments and control group. Black circles represent the  
784 effect sizes and the distribution of these effect obtained through nonparametric bootstrap

785 sampling (5,000 samples) is represented by the shaded area. Vertical bars represent the 95%  
786 confidence intervals.

787 Figure 4. Partial effects of the relationship between changes in oxygen uptake ( $\Delta\dot{M}O_2$ ) over  
788 time for 8 different doses of LPS injection (a-h) estimated from a generalized additive mixed-  
789 effects model (GAMM).  $\Delta\dot{M}O_2$  is calculated by subtracting pre-injection  $\dot{M}O_2$  from post-  
790 injection  $\dot{M}O_2$  at comparable time periods pre- and post-injection. Values above the red line  
791 indicate an increase in  $\dot{M}O_2$  relative to pre-injection values, and values below the red line  
792 indicate a decrease in  $\dot{M}O_2$  relative to pre-injection values. The red solid line represents the  
793 model-estimated mean centered on zero for the smooth term and the shaded area denotes a 95%  
794 confidence interval. The vertical dotted line represents the separation between nighttime 1 to  
795 10 hours after insertion in the respirometer (left side of the line) and daytime 10 to 20 hours  
796 after insertion in the respirometer (right side of the line).

797  
798 Figure 5. Mean differences in the spleen somatic index of *Pomacentrus amboinensis* damselfish  
799 among LPS treatment groups (a) Spleen somatic index (SSI) of the handling control group  
800 (purple), the saline control group (dark blue) and the LPS injected group with LPS3 (3mg kg<sup>-1</sup>)  
801 (indigo), LPS5 (5mg kg<sup>-1</sup>) (teal), LPS10 (10mg kg<sup>-1</sup>) (forest green), LPS30 (30mg kg<sup>-1</sup>)  
802 (harlequin green), LPS50 (50mg kg<sup>-1</sup>) (mustard) and LPS100 (100mg kg<sup>-1</sup>) (yellow). The  
803 vertical black lines indicate the standard deviation around the group means. (b) Estimation plots  
804 indicate the mean difference in SSI between treatments and the control group. Black circles  
805 represent the effect sizes and the distribution of these effects obtained through nonparametric  
806 bootstrap sampling (5,000 samples) is represented by the shaded area. Vertical bars represent  
807 the 95% confidence intervals.

808

809 Figure 6. Mean differences in the immune gene expression of *Pomacentrus amboinensis*  
810 damselfish among LPS treatment groups (a) Gene expression (polydom/svep1) of the handling  
811 control group (purple), the saline control group (dark blue) and the LPS injected group with  
812 LPS3 (3mg kg<sup>-1</sup>) (indigo), LPS5 (5mg kg<sup>-1</sup>) (teal), LPS10 (10mg kg<sup>-1</sup>) (forest green), LPS30  
813 (30mg kg<sup>-1</sup>) (harlequin green), LPS50 (50mg kg<sup>-1</sup>) (mustard) and LPS100 (100mg kg<sup>-1</sup>)  
814 (yellow). Vertical bars represent the standard deviation around the group means. (b) Estimation  
815 plots indicate the mean difference in gene expression between treatments and control group.  
816 Black circles represent the effect sizes and the distribution of these effect obtained through  
817 nonparametric bootstrap sampling (5,000 samples) is represented by the shaded area. Vertical  
818 bars represent the 95% confidence intervals.

Table 1 – Parameter estimates for  $\Delta\dot{M}O_2$  through time and doses of LPS; 3mg kg<sup>-1</sup> (LPS3), 5 mg kg<sup>-1</sup> (LPS5), 10mg kg<sup>-1</sup> (LPS10), 30mg kg<sup>-1</sup> (LPS30), 50mg kg<sup>-1</sup> (LPS50) and 100mg kg<sup>-1</sup> (LPS100) using a generalized additive mixed model with fish ID as a random effect. Time is smoothed across treatment to detect specific trend overtime for each treatment. Effective degrees of freedom (edf) estimated residual degrees of freedom (ref.df) and Chi.sq are given for smoothed terms. *P* values indicate whether the estimated function (s) reject or not the null hypothesis (i.e., a flat constant function). Estimate, standard error (SE), Chi.sq and *p*-values are given for the parametric variable treatment.

Variable	edf	Ref.df	Chi.sq	<i>p</i> -values
s (Time: handling control)	12.30	29	8022.4	0.005302**
s (Time: saline)	21.82	29	25067.8	0.000943***
s (Time: LPS3)	23.22	29	18760.7	0.003492**
s (Time: LPS5)	18,19	29	7679.0	0.044357*
s (Time: LPS10)	16.44	29	10785.7	0.009976**
s (Time: LPS30)	11.03	29	7472.0	0.003179**
s (Time: LPS50)	14.82	29	562.3	0.536626
s (Time: LPS100)	16.70	29	357.0	0.654444

Table 2 – Spleen somatic index (SSI) expressed in percentage, immune gene expression (polydom/svep1) and area under the curve (AUC) post-injection expressed in  $\dot{M}O_2$  (mg  $O_2$  h<sup>-1</sup>) of damselfish for saline injection and across the 6 doses of LPS compared to the handling control group.

	Saline vs. handling control ( $\Delta$ [95% CI])	LPS3 vs. handling control ( $\Delta$ [95% CI])	LPS5 vs. handling control ( $\Delta$ [95% CI])	LPS10 vs. handling control ( $\Delta$ [95% CI])	LPS30 vs. handling control ( $\Delta$ [95% CI])	LPS50 vs. handling control ( $\Delta$ [95% CI])	LPS100 vs. handling control ( $\Delta$ [95% CI])
Spleen somatic index (SSI)	-0.005 [0.025 – 0.006]	0.009 [0.016 – 0.027]	-0.006 [0.025 – 0.004]	0.005 [0.016 – 0.022]	0.006 [0.014 – 0.022]	-0.0002 [0.019 – 0.012]	0.021 [0.0009 – 0.0373]
Gene expression (polydom/svep1)	0.254 [-0.583 – 1.26]	2.21 [0.553 – 4.19]	0.159 [-0.859 – 1.43]	0.158 [-0.690 – 1.31]	1.18 [-0.031 – 3.07]	1.04 [0.217 – 2.1]	1.26 [0.476 – 2.32]
Area under the curve (AUC)	51.6 [-339 – 401]	80.3 [-151 – 208]	66.5 [-169 – 213]	33.8 [-199 – 166]	-71.6 [-321 – 79.3]	226 [-32.9 – 495]	63.2 [-191 – 214]

SSI, gene expression and area under the curve data values are mean differences between handling control and treatment groups with 95% bootstrapped confidence interval ( $\Delta$  [95% CI]).

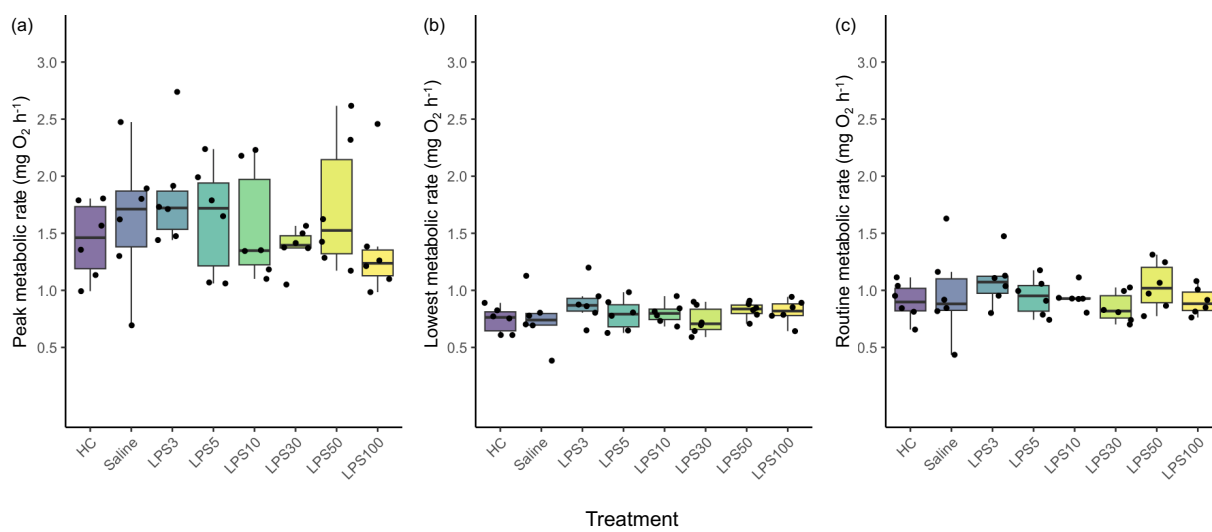


Figure 1

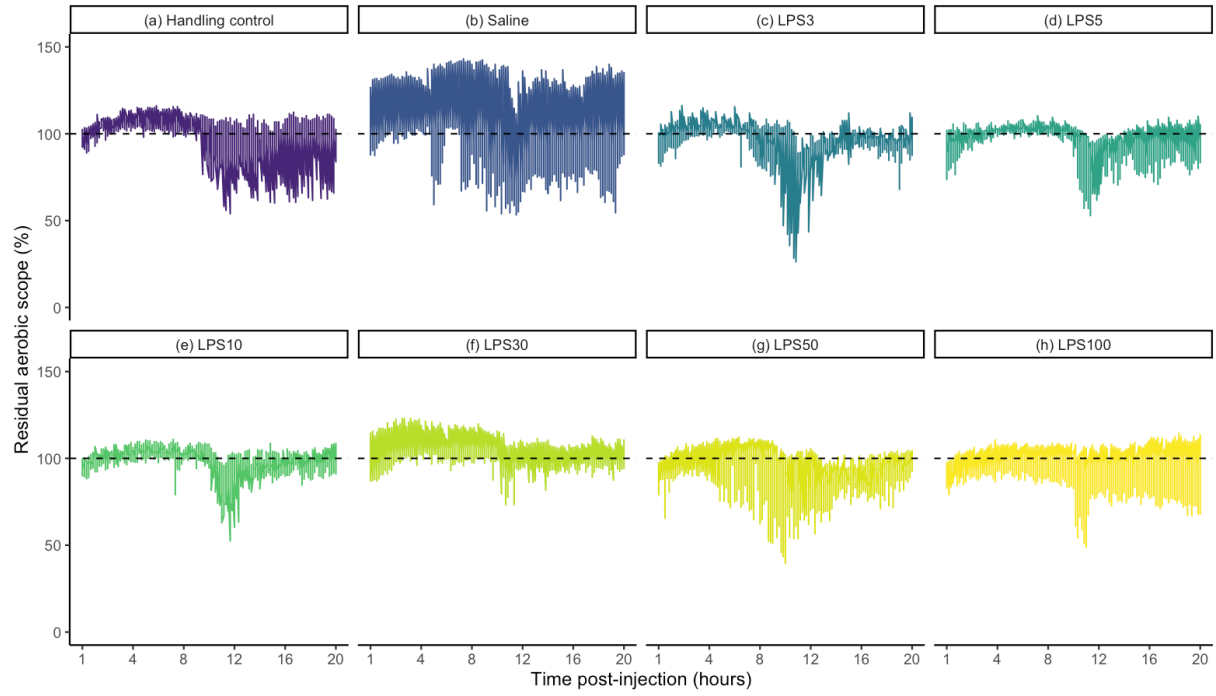


Figure 2

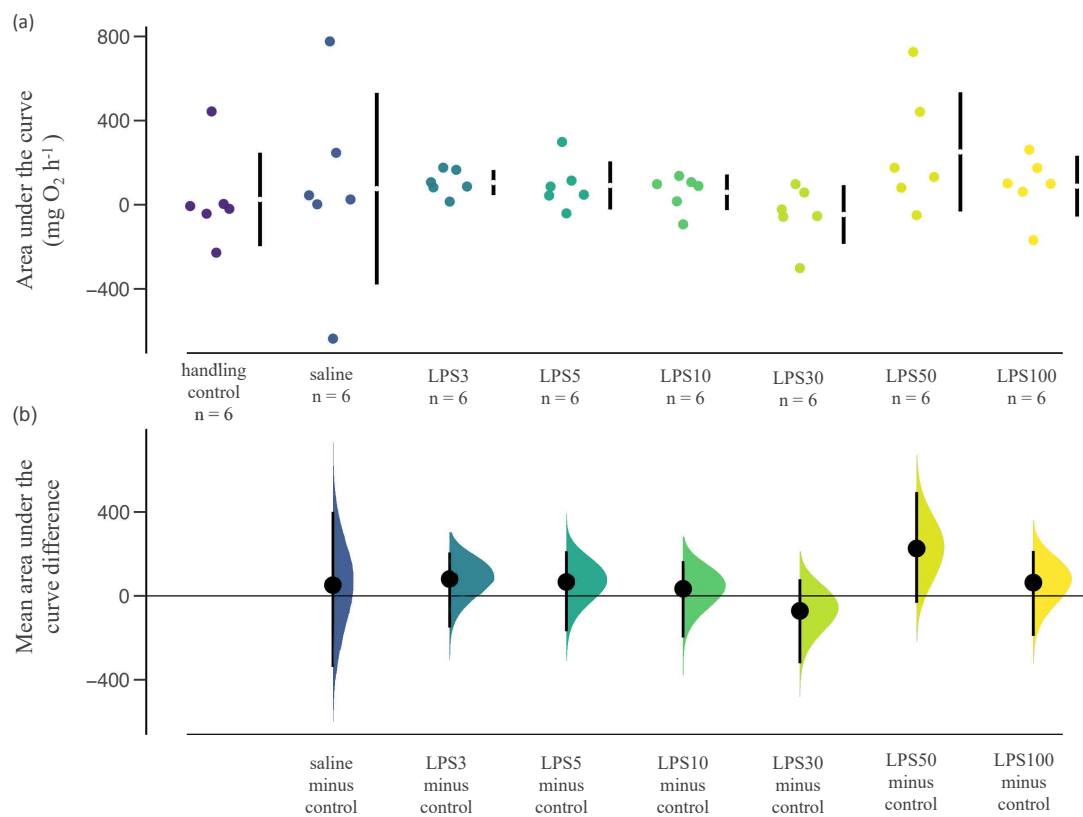


Figure 3



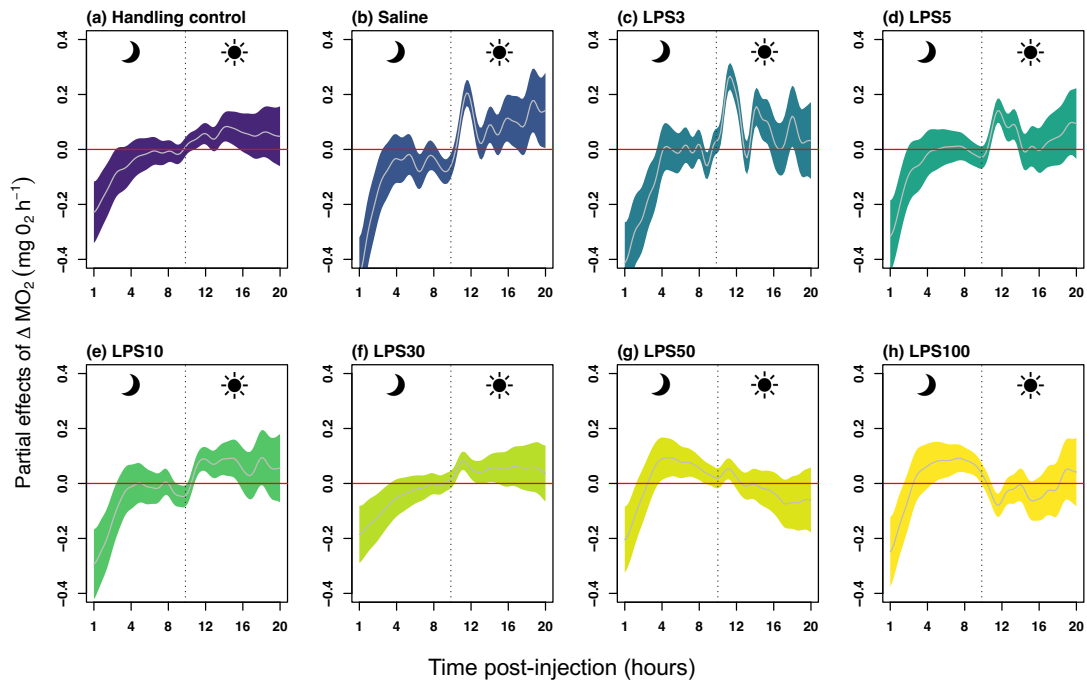


Figure 4

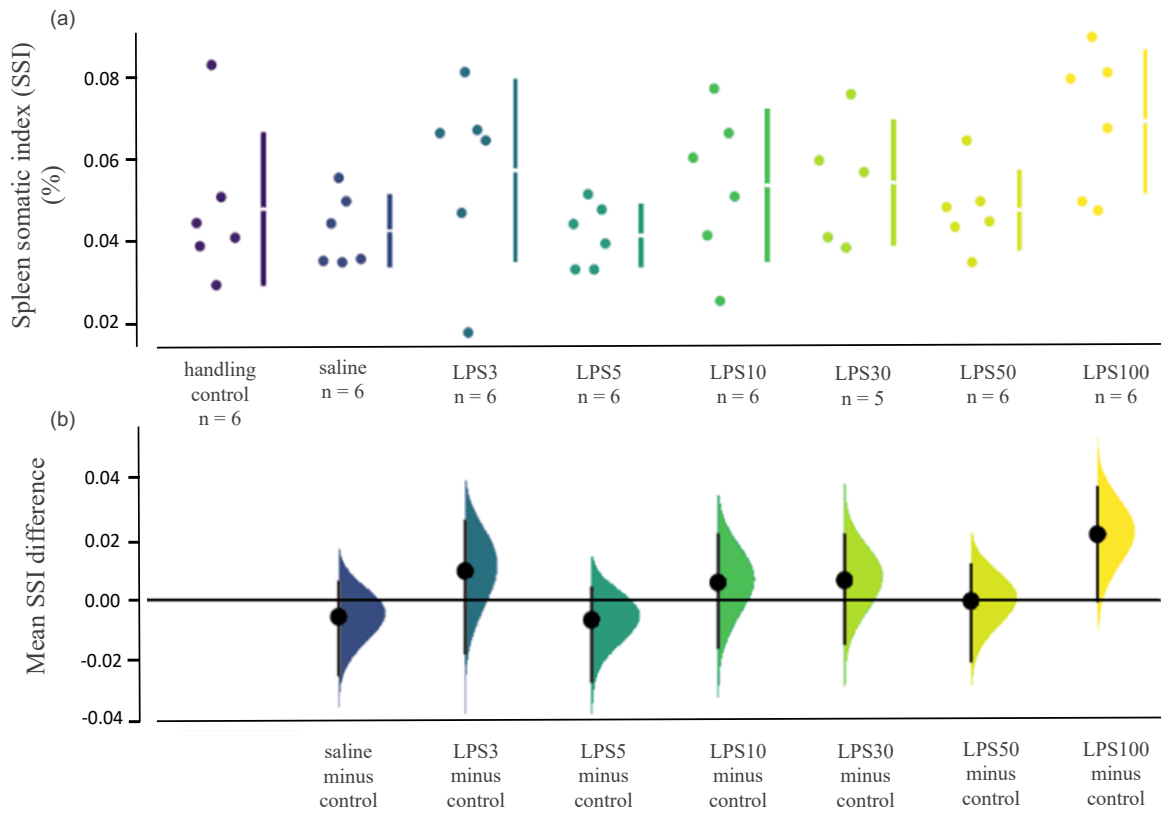


Figure 5

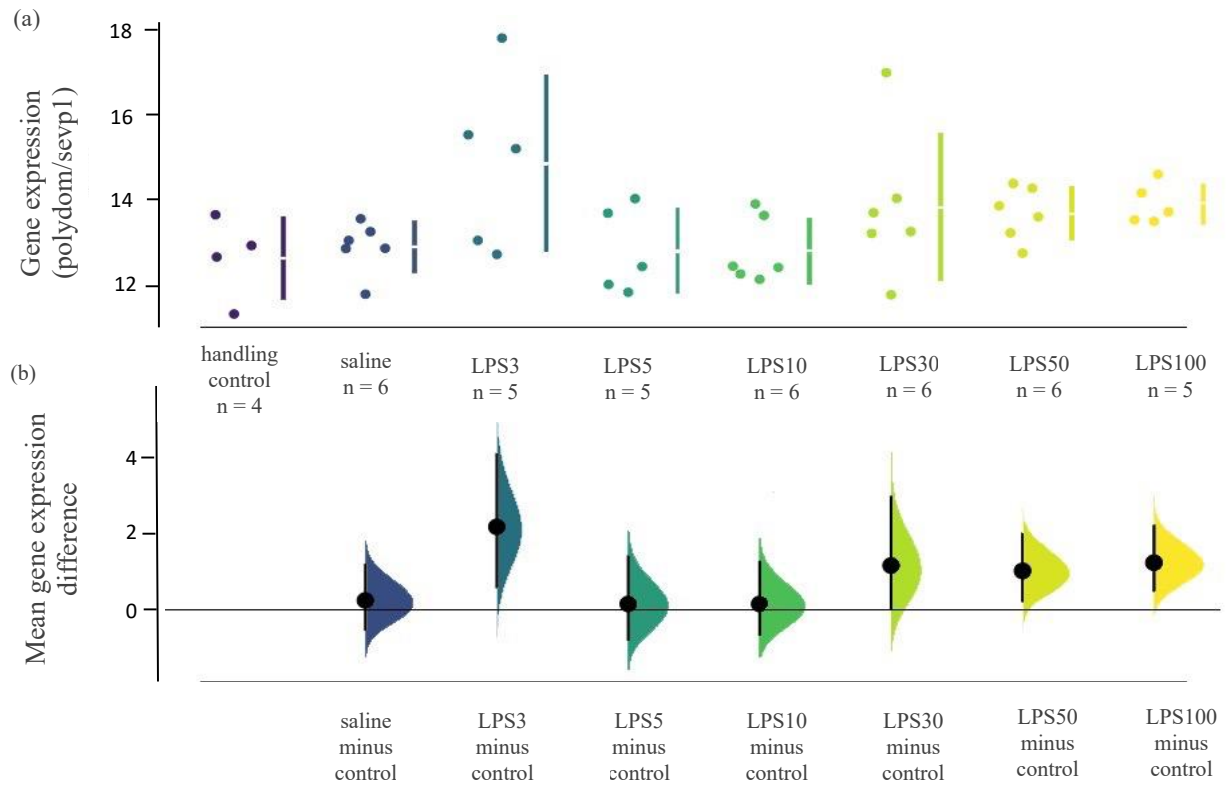


Figure 6