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1 Immunological Responses of Customised Probiotics-Fed Marron, *Cherax tenuimanus*, (Smith
2 1912) When Challenged with *Vibrio mimicus*

3
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10
11
12 **Abstract**

13
14 A two-phased experiment was conducted to investigate the effects of dietary
15 supplementation of customised probiotics on marron physiology. During the first phase
16 marron were fed probiotic supplemented feed for 70 days, while in phase two the same
17 marron were challenged with *Vibrio mimicus* and their physiological responses were
18 investigated for 4 days post-challenged.

19 The experiment was carried out in a purpose-built room, designed for aquaculture research,
20 using 18 of 250 L cylindrical plastic tanks. Five species of isolated probiotic bacteria from
21 commercial probiotic products and marron's intestine were tested in this experiment. The
22 probiotic bacteria were (*Bacillus* sp.); A10 (*Bacillus mycoides*); A12 (*Shewanella* sp.); PM3
23 (*Bacillus subtilis*); and PM4 (*Bacillus* sp.), which were added to the formulated basal marron
24 diet (34% crude protein, 8% crude lipid, 6% ash) at a concentration of 10^8 cfu/g of feed.
25 Immune responses of marron fed probiotics were evaluated by investigating organosomatic
26 indices, growth rate, survival, intermoult period, total haemocytes counts (THC), proportion
27 of granular cells (GC), bacteraemia, bacteria load in the intestine and water quality. The
28 results showed that dietary supplementation of probiotics in marron had no significant impact
29 on growth, intermoult period and survival of the marron. However, their supplementation
30 improved the physiological condition of marron in terms of significantly higher tail muscle
31 indices, THC and proportion of granular cells (GC) and reduced bacterial load in the
32 haemolymph. The addition of probiotics in marron diets also increased the bacteria load in
33 the marron intestine.

34 In addition, dietary supplementation of the customised probiotics was effective in improving
35 the resistance of marron against *V. mimicus* as they had higher THC, higher proportion of
36 GC and lower presence of bacteria in their haemolymph, after marron were challenged with
37 *V. mimicus*. The results also showed that probiotic *Bacillus mycoides* (A10) and PM4 are the
38 most beneficial dietary probiotics for marron health.

39
40 Key words: probiotics, physiological condition, immunological responses, marron
41

1. Introduction

Since the abuse of antibiotics and other chemicals for disease management in many aquaculture facilities were uncovered, safety in seafood products has received public attention. The research has shown that mismanagement of antibiotics use can lead to the emergence of bacterial resistance species [1]. Consequently, the research now is more focused in finding an alternative to antibiotics for disease management. Recently, probiotics have emerged as an alternative to antibiotics for disease management [2, 3]. The increasing demand for environment-friendly aquaculture has also led to probiotics becoming more popular as prophylactic agents and providers of improved nutrition [4, 5].

In recent years, studies in shrimp aquaculture have demonstrated that probiotics are beneficial for enhancing growth [6-9] and the immune system [10-15], by combating the pathogens through competitive exclusion mechanism [16-20]. The use of probiotics has also resulted in improving water and sediment quality [6, 10, 21, 22] and thus, leading to higher survival rates of healthier animals [1, 23-25]. Some species of probiotics have the ability to protect against viruses, although the mechanism of combating the virus is not fully understood [6, 26-28]. In aquaculture, probiotics can be applied either as a food supplement or as a water additive [21].

Marron, *Cherax tenuimanus* (Smith 1912) is one of the important freshwater crayfish species native to Western Australia [29]. Marron have attracted global interest as a potential aquaculture species due to their positive attributes, such as large harvest-size (up to 2 kg), higher price, non-burrowing behaviour, simple life cycle and ease of live transport [29-32].

Though fungus and parasites are dominant disease causing agents in marron aquaculture, yet prevalence of infection and incidence of mortality are relatively low [33, 34]. *Epistylis* and *Temnocephala* are two epibionts which are commonly found in marron, caused by poor water quality, particularly in unaerated ponds containing excessive organic matter [29]. These epibionts can decrease growth rates and reduce consumer appeal [31]. Although there is no current report on the losses in marron aquaculture caused by bacterial infection, the threat of marron getting bacterial infection are realistic as the industry grows and expands. *Vibrio mimicus* has emerged as a dominant bacterial pathogen of freshwater crayfish in aquaculture [35, 36].

The marron health can ultimately be assessed by growth and accepted survival rates, however, other physiological indicators, such as organosomatic indices, moisture content and osmoregulatory capacity, total haemocyte counts (THC), proportion of granular cells, bacteraemia and haemolymph clotting time [37], can also be used to understand the underlying mechanism for marron health. There has been no study on the effect of supplementation of probiotics in marron diet, therefore it is important to evaluate the effectiveness of dietary probiotics for the benefit of marron health under aquaculture environment.

The present study was designed to examine the effects of different sources of customised probiotic-supplemented feeds on the growth, survival, intermolt period, physiological, immune responses and bacteria load in the intestine of marron.

2. Materials and Methods

An experiment with two continuous phases was conducted. During the first phase, marron were fed different sources of probiotics supplemented diets and the second phase, involved the challenge test wherein, marron were injected with pathogenic bacteria *V. mimicus* under the laboratory conditions.

2.1. Experimental system

The experimental system was setup in a purpose-built laboratory designed for aquaculture research in the indoor aquarium facility of the Curtin Aquatic Research Laboratory (CARL), Curtin University, Perth, Western Australia. The experimental system

96 consisted of three standing units of steel racks with three shelves in each unit. Each rack
97 held six experimental units. The experimental units were cylindrical plastic tanks (80 cm
98 diameter and 50 cm high and 250 L in capacity). The tanks were filled up with freshwater
99 and supplied with constant aeration. Each tank was equipped with a submersible thermostat
100 set to 24°C and a recirculating biological filtration system (Fluval 205, Askoll, Italy). The
101 water in the tank was running continuously at a rate of approximately 3 L/min. The tanks
102 were also provided with sufficient number of marron shelters in the form of PVC pipes of
103 appropriate diameters.

104

105 2.2 Experimental animals

106 The marron juveniles (33 – 65 g) were purchased from Aquatic Resource Management
107 Pty Ltd., Manjimup, Western Australia. Before commencement of the experiments, all
108 juvenile healthy marron were kept for two weeks in holding tanks at CARL for acclimation.
109 The holding tanks were provided with aerated recirculating filtered freshwater. A commercial
110 pelleted diet (26% protein, 47- 50% carbohydrate, 9% fats and 8.9% ash) from Enviroplus
111 Pty Ltd., Perth Australia was fed to marron, at a rate of 3% body weight on alternative days.

112

113 2.3. Test diets

114 During the experimental period, the marron were fed a basal diet (34% crude protein,
115 8% crude lipid, 6% ash), formulated at CARL. The feed ingredients were passed through a
116 100 µm mesh sieve and thoroughly mixed to obtain uniform particle size. The largest
117 proportions of ingredients were mixed first before the smaller ones, to ensure all of the
118 ingredients were mixed well. A mince mixer was then used to make pellets. The pellets were
119 air dried, packed and stored at 4 °C until used.

120 Five species of probiotic bacteria isolated from the various sources were selected for
121 their growth inhibition capabilities against *Vibrio mimicus* and *V. cholera* non-01 and then
122 tested for pathogenicity by feeding to marron in a tank trial (unpublished results). *Bacillus*
123 *mycooides* (A10) and *Shewanella* sp. (A12) were selected from a number of healthy farmed
124 marron intestines. ; *Bacillus* sp. (AQ2) was selected a commercial product from Aquasonic
125 Pty. Ltd. and finally, *Bacillus subtilis* (PM3) and *Bacillus* sp. (PM4) were selected from
126 another commercial probiotic product supplied by Enviroplus Pty Ltd., Perth Australia.: All
127 probiotics were identified by the Bacteriology Laboratory, Animal Health Laboratories,
128 Department of Agriculture and Food, Western Australia. These selected probiotics have not
129 been reported as known pathogens of marron [38]. A basal diet without any probiotic
130 supplementation was used as a control diet.

131 The probiotics were supplemented to the basal diet using a described procedure [28],
132 with some modifications. The isolated probiotics from stock culture were re-grown onto a
133 new blood agar plate. After overnight incubation at 25°C, an appropriate inoculum of each
134 probiotic species was diluted into 20 mL of sterilized normal saline. Before being sprayed
135 onto the basal diet, all feeds were coated with fish oil blend (Bait mate®, Western Australia)
136 at 20 mL per kg basal diet. A concentration of 10⁸ cfu/g of feed was selected following the
137 previous studies [39-42]. The probiotic species were sprayed onto 1 kg of basal diet (10⁸
138 cfu/g feed) and then immediately covered with aluminium foil and stored in a refrigerator at
139 4°C to avoid bacterial growth. The concentration in cfu/mL, of each probiotic bacterium
140 sprayed onto the feed was determined using an established method [43] where optical
141 density (Spectrophotometer, BOECO S-20, Hamburg, Germany) correlates to the bacterial
142 concentration. The concentration sprayed onto the feed was confirmed by performing a total
143 bacterial count using blood agar plates and an overnight incubation at 25°C.

144

145 2.4. Feeding the marron with different probiotics supplemented diets – phase 1

146

147 Eighteen 250 L cylindrical plastic tanks were used to culture marron in a laboratory
148 scale experiment. Each tank was stocked with 7 healthy juvenile marron which were cultured

149 for 70 days, a time considered optimal for studying the marron growth and effects of
150 probiotics. The marron were fed with a basal (control) diet and probiotic-supplemented diets
151 at a rate of 1.5% / body weight every alternate day during the experimental period. Each
152 treatment was set up in triplicate. The effect of feeding treatments were measured in terms
153 of growth, survival, intermoult period, physiological response (organosomatic indices),
154 immune responses (total haemocyte count, differential haemocyte count and bacteraemia),
155 bacteria load in the intestine and water quality parameters.

156

157 2.5. Challenge test with *Vibrio mimicus* – phase 2

158 The marron from phase 1 were used in the challenge test (phase 2). In six culture
159 tanks, each tank was stocked with 3 marron and were fed at a rate of 1.5% / body weight
160 every day during the challenge test. Uneaten food and faeces were removed before feeding.

161 Pathogen bacteria *Vibrio mimicus* (isolated from blisters of dead yabbies, *Cherax*
162 *albidus*) was obtained from the Department of Agriculture and Food, Western Australia. A
163 stock solution of 2.04×10^8 cfu/mL was prepared for the injection. All marron in all tanks
164 were injected through the base of the fifth thoracic leg with 50 μ l of *V. mimicus*. Three
165 marron were injected with 50 μ l of artificial saline water as a mock challenge test (control).
166 The injected marron were kept in separate tanks before being put back into their original
167 tanks to avoid repeating sampling. The infected marron were monitored for survival, total
168 haemocyte count, differential haemocyte count and bacteraemia at 24, 48 and 96 h post-
169 injection.

170

171 2.6. Data collection

172 2.6.1. Feeding the marron with different probiotics supplemented diets – phase 1

173 - Survival rate of marron was determined by the formula:

174 Survival rate (%) = $(N_t - N_o) \times 100$,

175 where N_t and N_o are the number of marron at the end (t) of the experiment and at the
176 commencement (o), respectively.

177 - Marron biomass increment (BI) was measured by the following formula:

178 BI (%) = $[(\text{final biomass} - \text{initial biomass}) / \text{initial biomass}] \times 100$

179 - Specific Growth Rate (SGR) was measured by the following formula:

180 SGR (%)/g/day = $100 \times (\ln W_t - \ln W_o) / t$, where SGR is the specific growth rate in weight (%
181 g/day), and W_t and W_o are the weight of marron at current time (t) and at the
182 commencement of the experiment (o), respectively

183 Marron survival rate, specific growth rate (SGR) and biomass increment were
184 determined after every 14 days.

185 The presence of moults in tanks were checked daily and removed immediately.
186 intermoult period was determined as per established equation [44, 45].

187 $T_{im}(\text{day}) = T_{n+1} - T_n$

188 Where T_n = date of n moult; and T_{n+1} = date of $n+1$ moult

189

190 The wet hepatosomatic index (H_{iw}), tail muscle to wet body ratio (T/B_w), the
191 percentage moisture of the hepatopancreas (HM%) and tail muscle (TM%), wet
192 hepatopancreas index (H_{iw}), dry hepatopancreas index (H_{id}), wet tail muscle to wet body
193 weight ratios (T/B_w) and dry tail muscle to wet body weight ratios (T/B_d) were calculated as
194 per established equations [46, 47]. Moisture content and organosomatic indices
195 assessments were conducted at day 0, 35 and 70.

196 The immune parameters of marron, such as total haemocyte count (THC) and
197 differential haemocyte counts (DHC) were determined following the established protocol [46,
198 48]. Bacterial loads in the haemolymph (bacteraemia) were assessed by using the
199 established method [28]. All of the immune parameters were determined at commencement
200 of the experiment (0 day), in the middle of the experiment (35th day) and at the end of the
201 experiment (70th day).

202 The bacterial load in the haemolymph (bacteraemia) was assessed using total bacterial
203 counts on blood agar plates. Individual drops of the haemolymph aliquot were placed onto

204 blood agar plates and the lawn inoculated. For each marron, 3 drops of haemolymph were
205 tested for bacteraemia rank. The plates were then incubated for 24 hours at 25°C and CFUs
206 were determined for each drop. Total cfu/mL haemolymph was calculated on the basis of a
207 total volume of 50 µL for each drop. The bacterial loads in the haemolymph were ranked
208 from 1 (0 – 19 cfu/mL) to 10 (180-199 cfu/mL) and the rank 11 was used for “too numerous
209 for an accurate count”.

210

211 The bacterial load in the intestine (IBL) was determined following the protocol [4], with
212 some modification, on the initial (0 day), middle (35th day) and at the end of the experiment
213 (70th day). The marron samples were selected and rinsed in distilled water, quickly washed
214 with 70% alcohol and then rinsed again in sterilised distilled water to remove the external
215 bacteria. The intestinal tract of each marron was removed and weighed. A sterilised mortar
216 was used to homogenise the intestine, then diluted serially with sterilised normal saline
217 water, and was lawn inoculated to blood agar plates. The plates were incubated for 24 hours
218 at 25 °C. The bacterial loads were ranked from 1 (1 – 250 cfu/mL) to 10 (2501-3000 cfu/mL)
219 and the rank 11 was used for “too numerous for an accurate count”.

220 To maintain good water quality in every tank, water exchange at a rate of 10-15 % of the
221 total water volume was performed twice a week, after siphoning out the faeces and uneaten
222 feed. Water quality parameters were measured weekly which included: total ammonia and
223 nitrite which were measured using Calorimeter PR 1890, USA; temperature and pH using a
224 digital pH/mV/°C meter, Cyberscan pH300, Eutech instruments Singapore; and dissolved
225 oxygen was measured using a digital DO meter SM600, Milwaukee, Romania.

226

227 2.6.2. Challenge test with *Vibrio mimicus* – phase 2

228

229 Total haemocyte count (THC) and differential haemocyte count (DHC) of the marron
230 were determined before the *V. mimicus* injection and at 24 hours, 48 hours and 96 hours
231 after the injection. Bacteria in marron haemolymph (bacteraemia) were counted before the
232 *V. mimicus* injection and at 24 hours, 48 hours and 96 hours after the injection. The blood
233 agar plates were used for culture of haemolymph bacteria. The bacterial loads were ranked
234 from 1 (0 – 49 cfu/mL) to 10 (450 - 499 cfu/mL) and the rank 11 was used for “too numerous
235 for an accurate count. The number of dead marron was recorded daily after the injection to
236 determine accumulative mortality rate.

237

238 2.7. Data analysis

239 All collected data were analysed with SPSS statistical package version 18.0 (PASW
240 version 18.0) for Windows and Microsoft Excel. The results were presented as means ± SE
241 (standard error) and significant differences among treatment means were determined using
242 one way ANOVA (Analysis of variance) and Independent-Samples T test. The Tukey post
243 hoc test was used for multiple mean comparisons when the *P* value showed significance.
244 For ranking data (bacteraemia) and where the data were not normally distributed, the
245 Kruskal-Wallis test and Mann Whitney U test were used. All significant tests were performed
246 at *P*<0.05 level.

247

248 3. Results

249 3.1. Feeding the marron with different probiotics supplemented diets – phase 1

250 3.1.1. Organosomatic indices

251 There were no significant differences (*P* > 0.05) in HM%, TM%, H_{iw}, H_{id}, T_{iw} and T_{id}
252 among any marron fed probiotic supplemented diets over any sampling periods (Table1).
253 However, marron fed AQ2 showed a significantly higher (*P*<0.05) in T_{iw} and T_{id} at day 35
254 than day 0.

255

256 3.1.2. Immune responses

257

258 The mean THC, the mean proportion of granular cell (GC) and the mean haemolymph
259 bacteria ranks (bacteraemia) of the marron fed different probiotic diets were not significantly
260 different ($P > 0.05$) until day 35 of feeding (Table 2). However, at the end of phase-1, the
261 mean THC in all marron fed probiotic diets was significantly higher ($P < 0.05$) than the control
262 diet. The highest mean THC was for the A10 diet and the lowest was for the marron fed the
263 control diet, while other treatments had similar levels of THC. At the end of the day 70, the
264 mean GC was significantly higher ($P < 0.05$) in the marron fed AQ2, A10 and PM4
265 supplemented diets. The highest proportion of GC was associated with the PM4
266 supplemented diet (35.54 ± 4.79 %) and the lowest one was with the control diet
267 (10.64 ± 0.60). At the end of phase-1, the mean bacteraemia was significantly lower ($P < 0.05$)
268 in the marron fed probiotic supplemented, than in the marron fed the control diet. The
269 marron fed PM3 and PM4 diets had the lowest bacteraemia, followed by marron fed A10 and
270 AQ2 respectively. The bacteraemia of PM3-treated marron decreased gradually over the
271 culture period, while other parameters fluctuated during phase 1 of the trial.

272

273 3.1.3. Intestinal bacteria load (IBL)

274 The mean intestinal bacteria load (IBL) of marron fed different probiotic test diets were
275 the same ($P > 0.05$) at the commencement of phase-1 (Fig 1). At day 35, the IBL was
276 significantly different among the treatments ($P < 0.05$): the highest IBL was in the marron fed
277 A10, while the lowest were in the control and PM4 dietary treatments. At the end of the trial,
278 means IBL were also significantly different ($P < 0.05$) among marron fed different diets but,
279 the marron fed A10 had the highest IBL and the marron fed the basal diet had the lowest
280 rank.

281

282 3.1.4. Survival rate, growth and intermoult period

283

284 There was no significant difference ($P > 0.05$) in the mean survival rates, mean
285 biomass increment (BI %) and specific growth rate (SGR) in marron fed any diets. Intermoult
286 period (days) ranged between 25.25 ± 8.50 to 43.00 ± 1.00 and there was no significant
287 difference ($P > 0.05$) between marron fed different diet.

288

289 3.2. Challenge test with *Vibrio mimicus* – phase 2

290 3.2.1. Total haemocyte count (THC)

291

292 The mean THC of marron were not significantly different among any dietary treatments
293 before the *V. mimicus* challenge (Table 3). After *V. mimicus* injection, there were significant
294 decreases ($P < 0.05$) in the mean THC in all treatments at 24 hours, 48 hours and 96 hours.
295 At 24 hours post injection, the THC of marron fed probiotic-supplemented diets were
296 significantly higher ($P < 0.05$) than those fed the control diet. The marron fed the diet
297 supplemented with PM3 showed the highest THC ($1.53 \pm 0.35 \times 10^6$ cells/mL), whereas the
298 marron fed the control diet had the lowest THC ($0.36 \pm 0.07 \times 10^6$ cells/mL). After 48 hours
299 into the challenge, the THC of marron fed probiotic-supplemented diets were significantly
300 higher ($P < 0.05$) in marron fed A10, A12 and PM4, whereas the other dietary treatments
301 showed similar values as the control. After 96 hours of *V. mimicus* injection, A10, PM3 and
302 PM4 treatments showed significant differences ($P < 0.05$) than the control. Whereas, other
303 probiotic fed marron had similar mean THC to those of the control.

304

305 3.2.2. Differential Haemocyte Count (DHC)

306 Before the *V. mimicus* was injected, the mean proportions of granular cells (GC) of
307 marron fed the probiotic-supplemented diets were significantly different ($P < 0.05$) than the
308 control. The PM4 treatment produced the highest GC proportion, followed by the A10
309 treatment (Table 3). Other treatments had a similar proportion of GC as the control. After *V.*
310 *mimicus* injection, marron exhibited a declining trend ($P < 0.05$) in the mean GC proportions
311 in all treatments. At 24 hours post-injection, there was a significant difference ($P < 0.05$)
312 among treatments, with the PM3 producing the highest GC proportion and the control

313 showing the lowest. At 48 hours into the challenge, there was a significantly higher ($P <$
314 0.05) GC proportion for A12 and PM4 supplemented diets than others. At 96 hours, the GC
315 proportion of marron fed probiotic-supplemented diets were significantly higher ($P < 0.05$)
316 than the marron fed control diet.

317

318 3.2.3. Bacteraemia

319 The mean bacteraemia ranks of the marron fed probiotic-supplemented diets were
320 significantly lower than the marron fed control diet. At 24 h post-injection with *V. mimicus*,
321 there were significant increases ($P < 0.05$) of bacteraemia ranks in all treatments. However,
322 there was no significant difference ($P > 0.05$) of bacteraemia 24 h post *V. mimicus* injection.
323 These bacteraemia values remained similar to 48 h post-injection.

324 After 96 h, there was a significant decrease ($P < 0.05$) of the bacteraemia ranks of
325 marron fed AQ2, A10 and PM4. There was also a significant difference ($P < 0.05$) in
326 bacteraemia ranks among treatments, of which the lowest bacteraemia was observed in
327 marron fed A10 and PM4 supplemented diets. However, AQ2 and PM3 diets produced
328 similar ranks as the control diet.

329

330 3.2.4. Survival Rate

331 At 96 h post-injection, all marron injected with *V. mimicus* and those injected only with
332 normal saline water remained alive and started to respond to the feed given on day 4. The
333 normal saline injected marron, however, remained actively responsive to the feed given
334 earlier than bacteria injected marron.

335

336 4. Discussion

337 Hepatopancreas and tail muscle indices have been widely used as tools to monitor the
338 effects of different culture environments and diets on penaeid [49-53] and non-penaeid
339 shrimps [54, 55]. The present study shows that probiotic supplementation of *Bacillus sp* on
340 marron diet did improve the physiological condition of marron, as shown by the changes in
341 dry tail index (T_{id}). The significantly higher T_{id} observed in marron fed a diet with *Bacillus sp*.
342 indicates that this probiotic bacteria helped the marron to absorb nutrients more efficiently
343 and thus, they could store more energy in their tail muscle [56]. Similarly, probiotic bacteria
344 in the formulated shrimp diet, act as a facilitator to digest all of the protein components,
345 which in turn synthesises required enzymes responsible for increasing the shrimp's protease
346 activity and food digestibility [57].

347 Haemocytes perform a key role in the host-defence mechanism of crustaceans by
348 destroying micro-organic invaders [58, 59]. Environmental conditions, such as the presence
349 of stressors and disease, have a relationship with the number of circulating haemocytes in
350 crayfish [60]. A poor health condition in crayfish can be reflected by a low number of total
351 haemocyte counts in their haemolymph [55]. The results from this study showed that marron
352 fed probiotic diets had higher THC counts than those fed the control diet; this indicates that
353 marron are healthier when they are fed a probiotic supplemented diet. The THC of the
354 western king prawn (*Penaeus latisulcatus*) also increased gradually when they were treated
355 with two selected probiotics (*Pseudomonas synxantha* and *Pseudomonas aeruginosa*) [61].
356 Administration of *Bacillus licheniformis* in the *Litopenaeus vannamei* diet enabled an
357 increase in the THC [62]. In the present study, *Bacillus mycoides* produced the highest THC,
358 demonstrating its higher effectiveness than other probiotics.

359 After being injected with *V. mimicus*, the THC decreased sharply. Similar results has
360 been reported in Chinese shrimp (*Fenneropenaeus chinensis*) injected with *Vibrio*
361 *anguillarum* [63]. Another study also found the THC of *V. mimicus* injected marron declined
362 [64]. Environmental stressors and diseases can influence the number of circulating
363 haemocytes [55, 60, 65]. The decrease in THC after *V. mimicus* injection, could be
364 associated with an inflammatory response of the haemocytes leaving the circulation and
365 migrating to the site of the injection [66, 67]. However, the THC of marron fed probiotic diets
366 remained higher than those fed the probiotic-free diet during the challenge test. Similar
367 results has been reported in white shrimp (*Litopenaeus vannamei*) when challenged with *V.*

368 *harveyi*, after administration of probiotic bacteria (*Lactobacillus plantarum*) [68]. The higher
369 THC of the marron fed a probiotic-supplemented diet in the present study indicates the
370 greater ability of the marron haemocytes in the degranulation process to fight against the
371 foreign substance. Among the probiotics diet, PM4 (*Bacillus* sp.) resulted in the highest
372 THC. *Bacillus* sp. is known as an effective probiotic bacteria in shrimp aquaculture and has
373 been studied for its effectiveness by many researchers, under both laboratory and
374 commercial trials [6, 12, 24, 69, 70].

375 In the present study, the marron diets with probiotic supplementation of AQ2 (*Bacillus*
376 sp.), A10 (*Bacillus mycooides*) and PM4 (*Bacillus* sp.) produced significantly higher
377 proportions of granular cell in marron haemolymph than the control diet (Table 2). Therefore,
378 indicating a positive effect of the probiotic, *Bacillus spp* in enhancing the health condition of
379 these marron [55, 71]. After *V. mimicus* injection, the proportions of granular cells in marron
380 were decreased in all treatments. Similar results were also reported where an injection of 20
381 μL of 0.53 ± 10^6 cfu/mL *V. mimicus* reduced the proportion of GC [64]. In other species the
382 reduction of GC proportion have been reported in black tiger shrimp (*Penaeus monodon*)
383 [67] and Chinese shrimp (*Fenneropenaeus chinensis*) [63] when they were injected with
384 *Vibrio anguillarum*. A decreasing GC proportion suggests that GC degranulates at first,
385 followed by lysis of the lysosomes [63]. Nevertheless, the proportion of GC of marron fed
386 with probiotic diets remained higher than GC of marron fed probiotic-free diet (control) during
387 the challenge test. Similar results have been reported for western king prawn (*Penaeus*
388 *latisulcatus*) when challenged with *Vibrio harveyi* at 10^3 , 10^5 , and 10^7 CFU/mL [72]. The
389 marron fed the diets supplemented with genus *Bacillus* (AQ2, PM3 and PM4) and then
390 challenged with high dosage of *V. mimicus* had higher GC proportions than other treatments.
391 Similar to shrimps, *Bacillus* surface antigens, or their metabolites, act as immunogens by
392 stimulating phagocytic activity of granulocytes [73].

393 The bacteraemia of marron fed probiotic supplementation in their diets was lower than
394 the marron fed control diet. The probiotics act as immunostimulants to stimulate the non-
395 specific immune system against bacterial infection [6, 13, 74]. PM3 (*Bacillus subtilis*) and
396 PM4 (*Bacillus* sp) supplementations had the lowest haemolymph bacteraemia. Probiotic
397 bacteria *Bacillus*, not only competes for nutrients and thus inhibits other bacteria from
398 growing rapidly [5], they naturally are also able to produce many different antibiotic
399 compounds [21]. At 24 and 48 h post injection with *V. mimicus*, the bacteraemia values
400 increased sharply in all treatments. Similar results has been reported wherein bacteraemia
401 of marron fed the control diets (without immunostimulant Bio-Mos^T) increased significantly 24
402 and 48 h post *V. mimicus* injection [64]. However, the bacteraemia of marron fed probiotic-
403 supplemented diets had lower levels of bacteraemia than the control diet after 96 hours.
404 This indicates that the probiotics were effective in reducing the bacterial load in the marron
405 haemolymph. In western king prawn, administration of two combined probiotics,
406 *Pseudomonas synxantha* and *Pseudomonas aeruginosa* (10^5 cfu/mL), at 20 mL/kg feed,
407 reduced the number of bacteria in the haemolymph [28, 61]. The increase bacteraemia
408 levels in haemolymph indicates that the immune capacity of the animal has declined and
409 thus, possibly can result in increased susceptibility to infections [71].

410 The average survival rate of the marron in the feeding trial (phase 1) was 46.03% of
411 the initial stocking biomass of approximately 103.82 g/m². There was no evidence to prove
412 that supplementation of the probiotics in marron diets improved marron survival rate.
413 However, this rate is still higher than marron cultured in the earthen ponds, where the
414 survival of marron ranges from 13.82% to 34.66% with a lower stocking density (4.01 ± 0.28
415 g/m²) [53, 75]. When beta 1,3 β glucan is used as an immunostimulant, more than 65% of
416 the initial stocking biomass (16.83 g/m²) of marron can be achieved [37]. In an intensive
417 battery culture system the survival rate can also be higher (71%) than the present trial, due
418 to absence of cannibalism [76]. In the present study, the average survival was relatively low
419 as marron mortality was mainly caused by cannibalism. Cannibalism frequently occurs when
420 a population of crayfish is at high density and it is often triggered by moulting individuals [77].

421 In shrimp aquaculture, the application of probiotics, either through mixing into the
422 water or supplementing with feed, has been reported to increase the survival rate [3, 6-8, 12,

423 28, 61, 70]. The injection of *V. mimicus* did not affect the survival of marron (59.04 ± 1.64 g
424 initial mean weight). However, it took a few days for marron to recover and resume its
425 response to the feed given. Different results have been reported where 20 μ L of 0.53×10^6
426 cfu/mL of *V. mimicus* stock solution injected into marron (10.44 ± 0.20 g initial mean weight)
427 resulted in 100% mortalities in marron [64]. The different results in the present experiment
428 could be attributed to the low dosage of injected *V. mimicus* per unit weight of the marron, as
429 the mean weight of the marron was five times higher than those used by other authors [64].

430 Injection of *V. alginolyticus* into white shrimp (*Litopenaeus vannamei*) at dosages of
431 10^5 and 10^6 cfu/shrimp could produce survival rate of 40 – 50% [78, 79]. *P. monodon* mean
432 weight of 18g survived for 6 days after challenged by bath exposure to 10^7 cfu/mL of *V.*
433 *harveyi* 1526 in aquarium water [80], whereas, when challenged with 40 – 200 $\times 10^6$ cfu of *V.*
434 *anguillarum* at body weight of 2 – 4 g, *P. monodon* survived for a week [67]. Furthermore,
435 the isolates of *V. mimicus* at a dosage of 10^5 cfu, can cause high mortalities in *Cherax*
436 *albidus* [81]. This comparison shows that the survival of injected animals is dependent on
437 species of the pathogenic bacteria, the mean weight of the host species and the virulent
438 effects of the pathogen, rather than numbers of bacteria. Hence, in considering the
439 resistance of marron against *V. mimicus*, the virulent effect and the size of marron need to
440 be considered to explain the effects of a probiotic on marron survival.

441 Marron fed probiotic-supplemented diet in the present study did not show any growth
442 improvement or biomass increment, although supplementation has been reported to
443 increase the growth in other cultured shrimps [7, 8, 82, 83]. The average SGR in the present
444 experiment ranged from 0.13 to 0.66, with moult interval ranging between 25.5 to 43 days.
445 The average SGR of marron was different in various culture conditions; pond culture studies
446 range from 0.5 – 2.1 and intensive crayfish culture system studies range from 0.4 – 1.1 [84].
447 The intermoult period in juvenile marron ranged between 15 to 45 days [55, 85]. The
448 presence of probiotic bacteria in shrimp intestine can increase the health status of the host
449 by providing competitive exclusion in the shrimp's gut [86, 87]. Moreover, probiotics may
450 synthesise the vitamins that can lead to increased digestive activity or improve enzymatic
451 activities; therefore, this could be responsible for weight increase, improved digestion and/or
452 nutrient absorption [9].

453 The present study showed that marron fed probiotics had higher intestinal bacteria
454 ranks than those fed the control diet, especially diet supplemented with *Bacillus mycoides*.
455 Though our studies could not confirm whether higher bacterial load in the intestine of marron
456 are due to increased densities of probiotic species, other studies have reported that the
457 administration of probiotic *Bacillus licheniformis* can improve the white shrimp's intestinal
458 micro-flora and its immunity by colonizing and replacing pathogen bacteria in their intestine
459 [12, 69].

460 In a shrimp's digestive tract, *Bacillus* enhance the specific activity of lipase, protease
461 and amylase [8]. In the current study, most species of probiotic bacteria for the trial came
462 from the genus *Bacillus*, as the enzymes of *Bacillus* are very efficiently in digesting a large
463 variety of carbohydrates, lipids and proteins into smaller units [5]. In shrimp culture within
464 ponds, *Bacillus* species also degrade organic materials [1, 12], thus improving the water
465 quality.

466 In conclusion, using the matrix (Table 4) results from phase 1 and 2 showed that A10
467 can achieved the highest score, followed by PM4, AQ2, PM3, and A12. Furthermore, A10
468 (*Bacillus mycoides*) and PM4 (*Bacillus sp.*) are most effective probiotic supplements in the
469 marron diets. In the present study, marron fed probiotic supplemented diets produced
470 beneficial outcomes in terms of physiological condition (organosomatic indices), immune
471 parameters (THC, DHC, GC proportion and bacteria load in haemolymph) and level of the
472 bacteria load in the intestine than control diet. However, further research is required to
473 investigate the effects of supplementing two or three different species in marron diets under
474 a commercial farming environment.

475
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720 Table 1. Hepatopancreas moisture content (HM%), tail muscle moisture content (TM%), wet
 721 hepatosomatic index (H_{iw}), dry hepatosomatic index (H_{id}), wet tail muscle indices (T_{iw}) and
 722 dry tail muscle indices (T_{id}) of marron with different species of probiotic bacteria
 723 supplementation
 724

Index	Day	Treatment					
		AQ2	A10	A12	PM3	PM4	Control
HM%	0	161.97±1.16	161.97±1.16	161.97±1.16	161.97±1.16	161.97±1.16	161.97±1.16
	35	162.53±1.46 ^a	162.05±5.38 ^a	165.33±6.48 ^a	162.76±5.61 ^a	168.42±5.54 ^a	175.16±7.07 ^a
	70	158.44±1.30 ^a	163.16±5.40 ^a	164.17±3.28 ^a	165.11±3.92 ^a	161.44±1.38 ^a	170.79±6.26 ^a
TM%	0	280.52±0.30	180.52±0.30	180.52±0.30	180.52±0.30	180.52±0.30	180.52±0.30
	35	179.13±0.26 ^a	182.00±2.193 ^a	179.55±0.88 ^a	179.47±0.90 ^a	180.57±0.727 ^a	180.82±0.41 ^a
	70	178.74±0.19 ^a	179.34±0.83 ^a	180.36±1.13 ^a	186.46±6.36 ^a	179.15±0.09 ^a	180.67±0.94 ^a
H_{iw}	0	16.06±1.10	16.06±1.10	16.06±1.10	16.06±1.10	16.06±1.10	126.06±1.10
	35	17.83±0.87 ^a	16.99±0.64 ^a	15.87±0.85 ^a	16.73±1.29 ^a	16.70±1.27 ^a	15.50±0.51 ^a
	70	15.86±0.68 ^a	16.30±0.66 ^a	16.17±0.65 ^a	16.72±0.50 ^a	16.44±0.81 ^a	28.15±0.91 ^a
H_{id}	0	12.33±0.49	12.33±0.49	12.33±0.49	12.33±0.49	12.33±0.49	12.33±0.49
	35	12.96±0.43 ^a	12.72±0.63 ^a	11.94±0.223 ^a	12.65±0.81 ^a	12.26±0.84 ^a	11.39±0.48 ^a
	70	12.76±0.17 ^a	12.34±0.46 ^a	12.19±0.19 ^a	12.32±0.13 ^a	12.50±0.29 ^a	12.29±0.88 ^a
T_{iw}	0	126.44±1.35	126.44±1.35	126.44±1.35	126.44±1.35	126.44±1.35	126.44±1.35
	35	229.69±0.91 ^a	135.32±6.51 ^a	128.60±0.77 ^a	128.04±0.54 ^a	127.21±1.51 ^a	127.43±1.21 ^a
	70	125.80±1.04 ^a	125.42±1.07 ^a	128.58±2.55 ^a	124.90±3.00 ^a	127.96±0.94 ^a	127.64±1.09 ^a
T_{id}	0	15.15±0.24 ^a	15.15±0.24 ^a	15.15±0.24 ^a	15.15±0.24 ^a	15.15±0.24 ^a	15.15±0.24 ^a
	35	26.20±0.26 ^a	16.07±0.26 ^a	15.84±0.17 ^a	15.75±0.23 ^a	15.27±0.24 ^a	15.27±0.32 ^a
	70	125.49±0.25 ^a	15.24±0.12 ^a	15.64±0.72 ^a	13.72±1.86 ^a	15.83±0.18 ^a	15.36±0.45 ^a

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 726 Data in the same column within an index having different subscript letters (_{1,2}) are significantly
 727 different at α level of 0.05. Data in the same row having the same superscript letter indicate a similar
 728 mean which is not significantly different at α level of 0.05.

729 Note: AQ2 (*Bacillus* sp.); A10 (*Bacillus mycoides*); A12 (*Shewanella* sp.); PM3 (*Bacillus subtilis*); PM4
 730 (*Bacillus* sp)

731

732 Table 2. Mean \pm SE marron immune parameters
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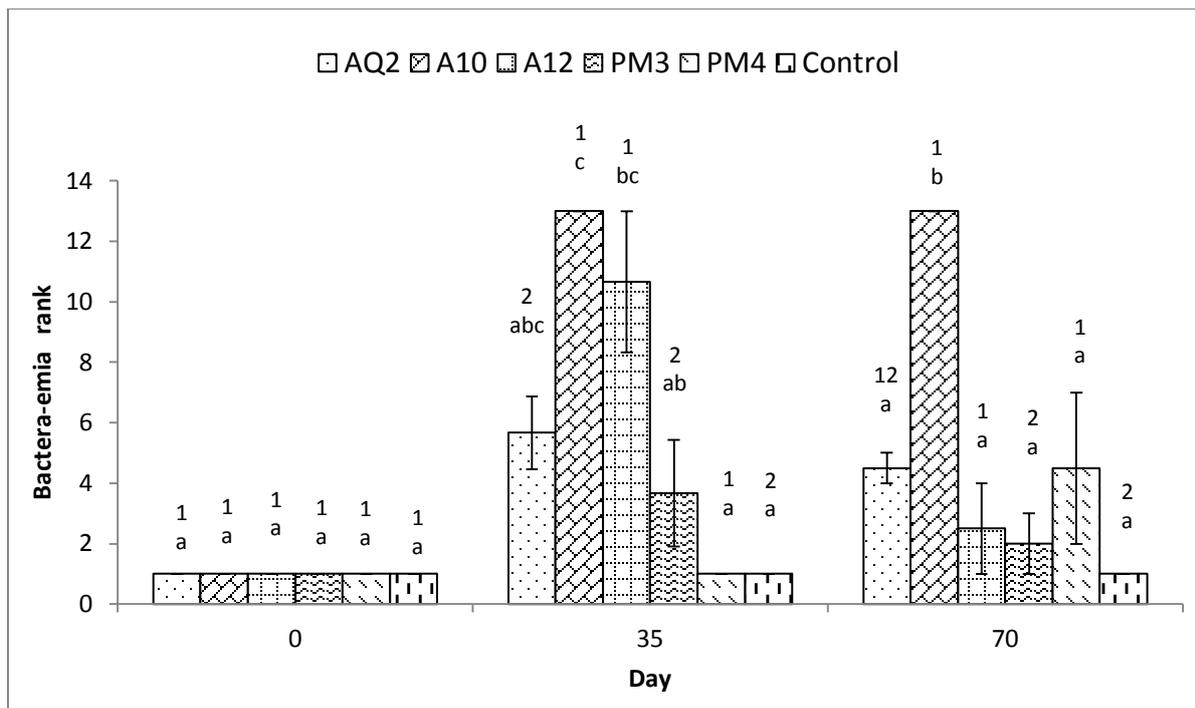
Immune parameter	Day	Treatment					
		AQ2	A10	A12	PM3	PM4	Control
Total Haemocyte Count (THC)	0	10.84 ± 0.00^a	10.84 ± 0.00^a	10.84 ± 0.00^a	10.84 ± 0.00^a	10.84 ± 0.00^a	10.84 ± 0.00^a
	35	210.17 ± 3.57^a	212.25 ± 2.46^a	211.79 ± 2.81^a	210.65 ± 1.45^a	210.13 ± 1.55^a	17.96 ± 0.88^a
	70	214.47 ± 0.46^{ab}	215.80 ± 1.30^b	29.81 ± 1.32^{ab}	211.43 ± 0.78^{ab}	211.06 ± 1.72^{ab}	17.73 ± 3.09^a
Granulocyte Cell (GC) (%)	0	349.96 ± 4.77^a	349.96 ± 4.77^a	349.96 ± 4.77^a	349.96 ± 4.77^a	349.96 ± 4.77^a	349.96 ± 4.77^a
	35	349.96 ± 4.77^a	1237.10 ± 8.60^a	1231.16 ± 7.02^a	231.44 ± 3.68^a	137.77 ± 2.28^a	223.31 ± 1.73^a
	70	120.14 ± 4.22^{ab}	$124.85.99 \pm 5.49^{ab}$	114.19 ± 5.25^a	113.39 ± 2.70^a	135.54 ± 4.79^b	110.64 ± 0.60^a
Haemolymph Bacterial Rank (BACTERAEemia)	0	25.00 ± 0.00^a	15.00 ± 0.00^a	15.00 ± 0.00^a	25.00 ± 0.00^a	25.00 ± 0.00^a	15.00 ± 0.00^a
	35	15.67 ± 1.20^a	113.00 ± 0.00^a	110.67 ± 2.33^a	13.67 ± 1.76^a	21.00 ± 0.00^a	11.00 ± 0.00^a
	70	14.50 ± 0.50^a	113.00 ± 0.00^a	2.50 ± 1.50^{ab}	2.00 ± 1.00^a	14.50 ± 2.50^a	11.00 ± 0.00^b

734
735 Data in the same column within an index having different subscript letters (1, 2) are significantly
736 different at α level of 0.05. Data in the same row having the same superscript letter indicate a similar
737 mean which is not significantly different at α level of 0.05.
738 Note: AQ2 (*Bacillus* sp.); A10 (*Bacillus mycoides*); A12 (*Shewanella* sp.); PM3 (*Bacillus subtilis*); PM4
739 (*Bacillus* sp)
740

741 Table 3. Mean \pm SE marron immune parameters when challenged with high dosage of
742 *V. mimicus*
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Immune parameter	Hour	Treatment					
		AQ2	A10	A12	PM3	PM4	Control
Total Haemocyte Count (THC)	0	110.10 ± 3.62^a	111.10 ± 2.25^a	110.54 ± 2.46^a	110.31 ± 1.35^a	19.77 ± 1.49^a	18.04 ± 1.67^a
	24	21.25 ± 0.10^{ab}	20.87 ± 0.01^{ab}	21.09 ± 0.08^{ab}	21.53 ± 0.35^b	21.10 ± 0.08^{ab}	20.36 ± 0.07^a
	48	21.19 ± 0.35^{ab}	20.82 ± 0.18^{ab}	20.87 ± 0.13^{ab}	21.36 ± 0.40^b	21.00 ± 0.05^{ab}	20.22 ± 0.05^a
	96	21.19 ± 0.19^{ab}	20.61 ± 0.01^a	20.81 ± 0.16^{ab}	21.34 ± 0.15^b	21.49 ± 0.07^b	20.53 ± 0.07^a
Granulocyte Cell (GC) (%)	0	120.14 ± 4.22^{ab}	124.65 ± 5.58^{ab}	114.19 ± 5.24^a	1213.39 ± 4.67^a	235.54 ± 4.79^b	110.64 ± 0.60^a
	24	115.98 ± 4.91^a	118.35 ± 0.21^a	117.90 ± 3.31^a	226.10 ± 0.66^a	124.93 ± 6.53^a	19.21 ± 0.12^a
	48	117.98 ± 4.36^{ab}	118.37 ± 2.09^{ab}	119.52 ± 0.78^{ab}	115.65 ± 1.17^{ab}	1216.73 ± 3.10^b	113.06 ± 0.19^a
	96	110.95 ± 1.64^b	116.46 ± 0.12^{ab}	119.36 ± 1.25^b	118.98 ± 2.04^b	118.13 ± 0.57^{ab}	111.59 ± 0.43^a
Haemolymph Bacteria Rank (BACTERAEemia)	0	11.00 ± 0.00^a	11.33 ± 0.57^a	11.67 ± 0.33^a	11.00 ± 0.00^a	11.00 ± 0.00^a	13.67 ± 0.67^b
	24	310.00 ± 1.00^a	39.50 ± 1.50^a	211.00 ± 0.00^a	311.00 ± 0.00^a	16.50 ± 4.50^a	211.00 ± 0.00^a
	48	310.00 ± 1.00^a	236.00 ± 1.00^a	211.00 ± 0.00^a	28.00 ± 1.00^a	17.50 ± 1.50^a	29.50 ± 0.50^a
	96	26.00 ± 1.00^{ab}	122.50 ± 0.50^a	210.00 ± 1.00^b	27.50 ± 0.50^b	12.50 ± 0.50^a	210.00 ± 1.00^b

745
746 Data in the same column within an index having different subscript letters (1, 2) are significantly
747 different at α level of 0.05. Data in the same row having the same superscript letter indicate a similar
748 mean which is not significantly different at α level of 0.05.
749 Note: AQ2 (*Bacillus* sp.); A10 (*Bacillus mycoides*); A12 (*Shewanella* sp.); PM3 (*Bacillus subtilis*); PM4
750 (*Bacillus* sp)
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754 Fig. 1. Mean \pm SE of intestine bacteria rank (IBL) of marron fed on different probiotic supplementation
755 diets. Different letters (a, b, c) over bars indicate significantly different means for different treatments at
756 $P < 0.05$. Different numbers (1, 2) over bars indicating significantly different means at different times at
757 $P < 0.05$

758 Note: AQ2 (*Bacillus* sp.); A10 (*Bacillus mycoides*); A12 (*Shewanella* sp.); PM3 (*Bacillus subtilis*); PM4
759 (*Bacillus* sp)

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762 Table 4. Matrix of comparison between five probiotic diets using various parameters

PARAMETERS	AQ	A10	A12	PM3	PM4
Organosomatic index	+	-	-	-	-
Biomass increment (%)	-	-	-	-	-
Specific Growth Rate	-	-	-	-	-
Intermoult period (days)	-	-	-	-	-
Survival rate (%)	-	-	-	-	-
Immune parameter:					
a. Total haemocyte count	+	+	+	+	+
b. Granular cell proportion (%)	+	+	-	-	+
c. Haemolymph bacterial rank	+	+	-	+	+
Bacterial assessment:					
a. Intestinal bacteria rank	-	+	-	-	-
b. Total water bacteria	-	-	-	-	-
Challenge test:					
a. Total haemocyte count	-	+	-	+	+
b. Granular cell proportion (%)	+	+	+	+	+
c. Survival rate	-	-	-	-	-
d. Haemolymph bacterial rank	-	+	-	-	+
e. Water quality	-	+	-	-	-

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Note: AQ2 (*Bacillus* sp.); A10 (*Bacillus mycoides*); A12 (*Shewanella* sp.);
PM3 (*Bacillus subtilis*); PM4 (*Bacillus* sp)
(+) = Significant (1 point)
(-) = Not significant (0 point)