

***Enterocytozoon hepatopenaei* Proliferate in *Procambarus clarkii*: A Warning for Crayfish and Shrimp Aquaculture**

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Abbreviation: Pc, *Procambarus clarkii*; Lv, *Litopenaeus vannamei*; EHP, *Enterocytozoon hepatopenaei*; AHPND, acute hepatopancreatic necrosis disease; WSD, White spot syndrome; HP, hepatopancreas; WF, white faces; IT: intestine; PM, peritrophic membrane; EL, epithelial layer; CT, connective tissue layer; ML, muscle layer; S, EHP mature spores; MP, merogonic plasmodia; Ex, exospore; En, endospore; Pm, plasm membrane or sporoplasm membrane; P, polaroplast; PF, polar filament; PT, polar tube; N, nucleus; AD, anchoring disc; EDD, electron-dense disk; *EHPptp2*, EHP polar tube protein 2; *cox*, mitochondrial cytochrome oxidase *c* subunit I; D-loop, mitochondrial control region (Displacement loop region); gDNA, genomic DNA; IFA, indirect-immunofluorescence assay; TEM, transmission electron microscopy; HPM, hepatopancreas microsporidiosis; SPF, specific pathogen free; PC, positive control; NC, negative control; NTC, no template control.

18 Abstract

19 The pacific whiteleg shrimp (*Litopenaeus vannamei*) and the crayfish
20 (*Procambarus clarkii*) are the most productive aquatic animals in the world. The
21 prevalence of the microsporidium *Enterocytozoon hepatopenaei* (EHP), an
22 intracellular spore-forming unicellular parasite which leading retarded growth of *L.*
23 *vannamei*, has caused severe economic losses in most shrimp farming country. In this
24 study, we found that the wild *P. clarkii*, living in shrimp ponds with EHP outbreak,
25 excreted white faeces after temporary laboratory culture. The hepatopancreas (HP) of
26 symptomatic crayfish exhibited a lighter color and severely atrophied. H&E-stain
27 showed tissue lesions in both hepatopancreas and intestine, and clustered
28 microsporidian spores were filled in the cytoplasm of the cells. PCR using *EHPptp2*
29 and two microsporidian-universal primers sets demonstrated the existence of EHP in
30 the hepatopancreas, intestine and the white feces of *P. clarkii*. The EHP loads of
31 10^3 - 10^4 copies of *EHPptp2*/50 ng HPgDNA were detected by qPCR. Developing
32 stages and mature spores of EHP were observed in hepatopancreas of *P. clarkii*
33 through indirect-immunofluorescence assay (IFA) and transmission electron
34 microscopy (TEM). Considering the large-scale cultivation of *L. vannamei* and *P.*
35 *clarkii*, overlap farming areas between these two species as well as the ability of
36 crayfish to crawl on land and water, our finding indicates the potential role of *P.*
37 *clarkii* in the transmission of EHP, and it is an early warning for crayfish and shrimp
38 farming.

39 **Keywords:** *Procambarus clarkii*, *Litopenaeus vannamei*, *Enterocytozoon*

40 *hepatopenaei*, Hepatopancreatic microsporidiosis, Transmission

41 **1. Introduction**

42 Pacific whiteleg shrimp (*Litopenaeus vannamei*) is one of the most productive
 43 aquatic animal in the world, reaching 5.8 million tons in 2020 (FAO, 2022). With the
 44 rapid expansion, disease problems plague the healthy development of shrimp farming.
 45 *Enterocytozoon hepatopenaei*, the pathogen of hepatopancreas microsporidiosis
 46 (HPM), is the microsporidium that can infect shrimp, resulting growth retardation and
 47 susceptibility to multiple infections with bacteria and viruses (Chayaburakul et al.,
 48 2004; Flegel, 2012). Since the first report of EHP in black tiger shrimp (*Penaeus*
 49 *monodon*) with stunted growth in Thailand (Tourtip et al., 2009), the pathogen has
 50 had a wider impact on *L. vannamei*, leading huge economic losses in most shrimp
 51 farming country and region (Ha et al., 2010; Wang et al., 2013; Tang et al., 2015; Tang
 52 et al., 2016; Biju et al., 2016; Shen et al., 2017; Tang et al., 2017; Kim et al., 2021).
 53 Aquaculture farmer lost their investment of manpower, resources, financial and time
 54 due to the chronic infection caused by EHP. For instance, Thailand's national
 55 economy lost has rose to US\$180 million from 2010 to 2016, and to US\$232 million
 56 in 2018 (Shinn et al., 2018; Patil et al., 2021). India lost US\$567.64 million due to
 57 EHP in 2018-2019 (Patil et al., 2021). Moreover, lost is even immeasurable in China
 58 (Wang et al., 2013).

59 The spores of EHP are oval profile, and $0.7 \times 1.1 \mu\text{m}$ in size (Tourtip et al., 2009).
 60 EHP has spore wall enveloping sporoplasm, in which it contains a nucleus, 5-6 coils

61 of the polar filament, a posterior vacuole, polaroplast, an anchoring disk attached to
62 the polar filament and organelles such as ribosome, endoplasmic reticulum, Golgi
63 apparatus, and mitochondrial remnants named mitosome (Tourtip et al., 2009; Weiss
64 and Becnel, 2014). The life cycle of EHP can be divide into proliferative stage,
65 sporogonic stage, and infective stage (mature spores) (Tourtip et al., 2009; Vavra and
66 Lukes, 2013; Chaijarasphong et al., 2021). At proliferative stage, cells divide firstly
67 by binary fission, followed by multiple nuclear divisions without cell division (Ning,
68 2020). At sporogonic stage (sporogonic plasmodium), the electron-dense disk (EDD)
69 forms the polar filament, and then dense secretions thickens the plasma membrane,
70 accompanying with the sporoblasts beginning to form. Spore wall of the sporoblasts
71 gradually thicken and spores develop into mature ones (Ning, 2020). EHP can be
72 transmitted horizontally through water (Salachan et al., 2017), feces (Singh and Singh,
73 2018) and carcasses (Tangprasittipap et al., 2013; Jang et al., 2022). Besides *L.*
74 *vannamei* and *P. monodon*, EHP was found proliferate in blue shrimp (*Penaeus*
75 *stylirostris*) (Tang et al., 2015), giant river prawn (*Macrobrachium rosenbergi*) (Wang
76 et al., 2022) and dragonfly (*Anax parthenope*, *Pantala flavescens*, and *Ischnura*
77 *senegalensis*) (Kumar Dewangan et al., 2023). EHP can also be transmitted between
78 live foods and other animals living in the same waters, such as polychaetes (Desrina
79 et al., 2020), brine shrimp (*Artemia salina*) (Karthikeyan and Sudhakaran, 2020), false
80 mussels (*Mytilopsis leucophaeata*) (Munkongwongsiri et al., 2022), marine crabs
81 (Mani et al., 2022) and some macrofauna (Wan Sajiri et al., 2023). Although a variety
82 of methods including specific pathogen free (SPF) shrimp larvae breeding has been

83 taken, the outbreak of EHP remained frequently happening, suggesting a complex
84 biological transmission pathway of EHP (Chayaburakul et al., 2004; Jang et al.,
85 2022).

86 The crayfish (*Procambarus clarkii*), native to North America, is farmed all over the
87 world and forms large wild populations now. It was introduced into China in the
88 1930s, then its production has soared yearly and reached 2.6 million tons in 2021
89 (Westman, 2002; Guo et al., 2017; Fisheries administration of the Ministry of
90 Agriculture and Rural Affairs of the People's Republic of China, 2022). Its main
91 domestic breeding areas in China are concentrated in the middle and lower reaches of
92 the Yangtze River at present (Guo et al., 2017; Fisheries administration of the
93 Ministry of Agriculture and Rural Affairs of the People's Republic of China,
94 2022). The crayfish and shrimp farming areas are usually overlapped in China, even
95 plenty of shrimp ponds are converted from crayfish ponds. The crayfish is capable of
96 crawling, digging and breeding, and can survive cold winters in burrows. They have
97 the habit of climbing ashore, cross dikes and enter other bodies of water (Xie et al.,
98 2008). These characteristics allow crayfish to transmit pathogens easily between
99 different water bodies.

100 In this study, we found the wild crayfish *P. clarkii* which inhabited in the farming
101 ponds of *L. vannamei* with slow growth syndrome were infected with EHP. The *P.*
102 *clarkii* shed white faeces during temporary laboratory culture. Molecular detection,
103 histopathological observation and ultrastructure analysis were used to demonstrate the

104 present and proliferation of EHP in the hepatopancreas of *P. clarkii*.

105 2. Material and methods

106 2.1 Sample collection

107 The wild crayfish ranging from 9.5 to 10.5 cm were collected from a *L.*
108 *vannamei* shrimp pond which suffered an outbreak of EHP in Chongqing, China.
109 These crayfish were temporary reared in a 100 L ton within 40 L water at 26°C. The
110 excrement and food residue were cleaned up, and half of the water was replaced by
111 fresh ones every day. White feces were collected after one day of laboratory culture.
112 Segments of white feces were fixed in 4% paraformaldehyde, while the remained
113 were stored at -20°C. The hepatopancreas and intestines of healthy and diseased
114 crayfish were sampled for EHP detection and histological analysis.

115 2.2 DNA extraction

116 Sampled hepatopancreas, intestine and white feces of the infected crayfish were
117 grounded thoroughly. Genomic DNA was extracted by 2% CTAB (1.40 M NaCl, 0.55
118 M CTAB, 125 mM EDTA, 40 mM Tris-HCl, and 0.2%-1% β-Mercaptoethanol,
119 pH=8.0) and 1 mg/ml protease K, and incubated at 65 °C for 2 h. After incubation, the
120 equivalent value of phenol/chloroform/isoamylalcohol extraction was added to the
121 digests. The extracted DNA in aqueous phase was successively precipitated by
122 isopropanol and washed by gradient ethanol (90% and 70%). The purified DNA was
123 dissolved in sterilized ddH₂O, diluted to 50 ng/μL, and stored at -20°C.

124 2.3 EHP detection and the crayfish identification

125 The primer sets targeting universal for microsporidia and EHP polar tube protein

126 2 (*EHPptp2*) were used to identify the infection of EHP (Zhu et al., 1993; Baker et al.,
127 1995; Wang et al., 2021; Wu et al., 2022). Two pairs of primers amplifying
128 mitochondrial genes were employed to identify the crayfish (Folmer et al., 1994; Li et
129 al., 2015). The details of primers used in this study were listed in Table 1. The PCR
130 products were cloned into pESI-Blunt vector and sequenced (Sangon, China).
131 Sequences of *SSU rRNA* (GenBank: MNPJ01000021.1), *EHPptp2* (GenBank:
132 MT249228.1), mitochondrial cytochrome oxidase *c* subunit I (*coxI*) (GenBank:
133 JN000903.1) and mitochondrial control region (Displacement loop region) (D-loop)
134 (GeneBank: KC556829.1) were download from NCBI. Sequence analysis was carried
135 out using MEGA (11.0.13) (Tamura et al., 2021) and GeneDoc (Nicholas and
136 Nicholas, 1997). qPCR was performed using Hieff qPCR SYBR® Green Master Mix
137 (Yeasen, China) in LightCycler® 96 real-time PCR instrument (Roche, China),
138 followed the instructions.
139

140 **2.4 Fluorescence microscopy and load assay of EHP**

141 Tissue samples were homogenized and appropriate amount were used to detect
142 the EHP spores through fluorescence microscopy. Samples were incubated with
143 Fluorescent Brightener 28 (Sigma Aldrich, USA) and Propidium iodide (Thermo
144 Fisher Scientific, USA) for 5 min, and then observed by fluorescence microscope
145 (Olympus, Japan) using differential interference contrast and ultra violet filter (Zhao
146 et al., 2020; Chen et al., 2021).

147 **2.5 Histological analysis by H&E-staining**

148 An appropriate tissue block was immediately cut from the live crayfish and fixed
149 in Davidson's AFA fixative. The tissues were embedded in paraffin and sliced to 5 μ m
150 thick. The paraffin sections were stained by the hematoxylin eosin staining kit
151 (Beyotime, China), then stained with Fluorescent Brightener 28 (Sigma Aldrich, USA)
152 for 10 minutes, and observed with microscope (Olympus, Japan) using differential
153 interference contrast and ultra violet filter.

154 **2.6 Indirect-immunofluorescence assay (IFA)**

155 IFA was performed to specifically recognize different phases of EHP in paraffin
156 sections prepared previously. Dewaxed paraffin sections were dealt with Citrate
157 antigen repair solution (Beyotime, China) for 30 minutes at 98 $^{\circ}$ C. After blocking,
158 sections were incubated with anti-EHP antisera or unimmunized antiserum (negative
159 serum) (diluent in 1:200), which is prepared by total proteins of EHP mature spores,
160 at room temperature for 2 hours. Then the sections were incubated with Alexa Flour®
161 488 conjugate goat anti-mouse IgG (Thermo Fisher Scientific, USA) following

162 washing steps. Fluorescent Brightener 28 (FB 28) and Propidium iodide (PI) were
163 used to stain chitin layer of spores and nuclei of cells respectively for 15 min. All
164 samples were sealed with ProLong® Gold antifade reagents (Thermo Fisher Scientific,
165 USA). Images were carried out by Olympus FV1200 laser scanning confocal
166 microscope (Olympus, Japan).

167 **2.7 Transmission Electron Microscopy (TEM)**

168 TEM was performed as described by Meng et al. (Meng et al., 2018) with slight
169 modifications. Hepatopancreas and feces were cut into small pieces ($\leq 1 \text{ mm}^3$) and
170 fixed in 5% glutaraldehyde over 24 hours. After washed with 0.1 M PBS buffer
171 (pH=7.4) three times (10 minutes each), samples were postfixated in 1% osmium
172 tetroxide for 2 h. The fixed tissues were embedded in epoxy resin Epon-812 and
173 polymerized at 37°C for 24 h and at 60°C for 48 h. Ultrathin sections (60-90 nm) were
174 collected on copper grids and stained with uranyl acetate and lead citrate.
175 Micrographs were taken using the transmission electron microscope (HITACHI,
176 Japan).

177 **3. Results**

178 **3.1 *P. clarkii* in *L. vannamei* farming ponds suffered EHP outbreak was positive** 179 **for EHP and exhibited white feces**

180 The wild crayfish, collected from a *L. vannamei* farming ponds suffered an EHP
181 outbreak, excreted white feces in the first five days under laboratory culture. Species
182 identification, using primers set LCO1490 and HCO2198 (Folmer et al., 1994), CRF
183 and CRR (Li et al., 2015) indicated the crayfish is *Procambarus clarkii* (Fig. S1 & Fig.

184 S2). The hepatopancreas of the symptomatic crayfish was discolored and atrophied.
 185 Besides lightened color, part of the intestine exhibited empty (Fig. 1 A-C).
 186 Fluorescence microscopy of the tissues smear showed numerous microsporidian
 187 spores and germinated spore coats presented in hepatopancreas, intestine and white
 188 feces (Fig. 2). Genomic DNA isolated from these samples was subjected to PCR
 189 amplification using primers targeting *SSU rRNA* of microsporidia and *EHPptp2* (Fig.
 190 3 A). These sequenced fragments were matched with the *SSU rRNA* and the
 191 conference *EHPptp2* respectively (Wiredu Boakye et al., 2017) (Fig. 3 C & D),
 192 demonstrating that the microsporidium found in wild *P. clarkii* is coincident with the
 193 conference EHP strain TH1. *EHPptp2* qPCR results showed 1.62×10^3 , 1.93×10^3 and
 194 5.61×10^4 copies/50 ng gDNA loads in intestine, hepatopancreas and white feces of
 195 EHP infected *P. clarkii* respectively (Fig. 3 B).

196 **3.2 Histological examination showed tissue damage in hepatopancreas and** 197 **intestine of EHP infected *P. clarkii***

198 H&E stained paraffin section showed a various degree of tissue lesion in the
 199 EHP-positive *P. clarkii* (Fig. 4). The hepatopancreas showed severely necrotic
 200 pathologic-changes that the structure of hepatopancreatic tubules was damaged and
 201 infiltrated by a large amount of hemolymph accompanied by a severe hemocyte
 202 aggregation (Fig. 4 A1). The basal layers of hepatopancreatic tubules were ruptured
 203 along with epithelial cells disruption and shed, and EHP spores distributed among the
 204 cytoplasm of necrotic epithelial cells and lumen of necrotic tubules (Fig. 4 A1-A3 &
 205 C1-C3). While in healthy tissues, the hepatopancreatic tubules maintain intact

206 morphological structure with epithelial cells tightly bound to the basal membrane, and
207 the hemocyte was blocked from hepatopancreatic tubule epithelial cells (Fig. 4 B &
208 D).

209 The intestine exhibited abnormal characteristics that epithelial cells necrosis,
210 cytoplasmic staining was not obvious, the interstices in the tissue increased, the edges
211 of the pleated ridge were sharper, and the muscle layer atrophied slightly (Fig. 4
212 E1-E3, G1-G3). In EHP-free crayfish, the peritrophic membrane, epithelial layer,
213 connective tissue layer and muscle layer were tightly ordered (Fig. 4 F & H). There
214 were some inclusions in the lumen (Fig. 4 F).

215 **3.3 Developing stages of EHP were found in the hepatopancreatic cells of *P.***

216 *clarkii*

217 In order to specifically indicate different stages of EHP, the polyclonal antiserum
218 against EHP proteins extracts was prepared. Anti-EHP serum can recognize
219 developing stages and mature spores of EHP. Combined with specific anti-EHP serum
220 and Fluorescent brightener 28 labeling chitin layer of spores, proliferative stage,
221 sporogonic stage and mature spores can be distinguished (Chen et al., 2017;
222 Senderskiy et al., 2021). Early merogonic plasmodia, characterized by what anti-EHP
223 serum can recognize without obvious chitin layer signal, were found in the
224 hepatopancreas of EHP infected *P. clarkii* (Fig. 5 A1-C5). Sporogonic stages and
225 mature spores were abundant in the hepatopancreas and intestine (Fig. 5 & Fig. S3).
226 No hybridization signal was observed in EHP infected *P. clarkii* slices with
227 unimmunized antiserum (Fig. S3) and in EHP-free *P. clarkii* slices with anti-EHP

228 serum (Fig. S4).

229 TEM further demonstrated the proliferation of EHP in crayfish hepatopancreatic
230 cells. Numerous mature spores and sporogonic stages of EHP clustered in the
231 cytoplasm (Fig. 6 A & B). Some sporogonic spores distributed near the host lipid
232 droplet, and the membrane of lipid droplet contacted closely by the spores was
233 deformed (Fig. 6 A & C). The mature spores were clearly characterized by exospore,
234 endospore, plasm membrane, polaroplast, anchoring disc, single nucleus and 5-6 coils
235 of polar filament (Fig. 6 E & F). EHP in sporogonial phase (during the sporogonic
236 stage) was distinguished by thinner chitin layer and more unordered polar filaments
237 (Fig. 6 E). Some spores were surrounded by a membranous structure individually or
238 in pairs (Fig. 6 B & D). The germination behavior in hepatopancreas was verified by a
239 germinated polar tube connected with EHP spore (Fig. 6 F).

240 4. Discussion

241 *L. vannamei* of and *P. clarkii* cultivation are the two most productive aquatic
242 animals. The world production of their exceeds 8 million tons, accounting for more
243 than 70% of the farmed crustaceans (FAO, 2022). With the rapid expansion of
244 breeding scale, bacteria, viruses, parasites and other diseases have brought great
245 trouble for aquaculture. It is noteworthy that the effects of pathogens vary from
246 periods and species. For instance, the impact of White spot syndrome virus (WSSV)
247 on crayfish was not obvious at first (Chou et al., 1995), but now it becomes one of the
248 most threatening disease agents (Baumgartner et al., 2009; Dragicevic et al., 2021).
249 EHP was initially neglected due to the prevalence of acute hepatopancreatic necrosis

disease (AHPND) and White spot syndrome (WSD), while it is one of the most influential pathogen in shrimp farming now (Tang et al., 2015). In 2022, EHP is classified as a pathogen of Class II animal diseases in China. The pathogen found to proliferate in *P. clarkii* is noteworthy and alarming.

Microsporidia are important ecological regulators in nature. When the population of a species is massive increase, microsporidia will then proliferate and play a role in population size regulation. Compared with viral and bacterial diseases, EHP is characterized by relatively low mortality and huge economic loss (Patil et al., 2021). Therefore, it is particularly important to prevent intra- and interspecies infection of EHP to reduce the risk at the source.

The color of hepatopancreas and intestine turned white, which is a signal of EHP infection. This phenomenon may be related to lipid reduction in the crab and shrimp (Ding, 2021; Wu et al., 2022), and the presence of a large number of spores in the *P. clarkii* (Fig. 1 B & C; Fig. 2). EHP may supply its own development by absorbing and converting lipids from host. It is clear that EHP in the proliferative stage are closely bound to the lipid droplet of the host (Fig. 6 A & C). But there's no way to see white hepatopancreas and intestine visually in *P. clarkii*. Because of the opaque exoskeleton, and EHP infected *P. clarkii* have no lesions on their appearance, it is hard to find crayfish infected. Flexible crawling ability also allows *P. clarkii* infected with EHP spread pathogens between different water bodies.

It is interesting that the crayfish in the same batch of samples (n=10) were negative to EHP by real-time qPCR after 2 weeks. However, the intestines of the same

batch of crayfish were similar to that after starvation, with atrophy of the intestinal villi and enlargement of the intestinal lumen (Fig. 4 G). We suspect that probably due to regeneration of hepatopancreas epithelial cells in crayfish. The *P. clarkii* infected with EHP in the beginning, with daily water change, may somehow drained EHP out and recovered finally, since the regeneration of the complete hepatopancreas epithelium of crayfish seems to last less than 2 weeks (Davis and Burnett, 1964; Strus et al., 2019; Vogt, 2019). The short-term expulsion of large number of active EHP spores is beneficial for the transmission of EHP, but harmful to the farmed animals it parasitizes. However, this strategy in crayfish has implications for treatment of EHP infected shrimps.

In general, this article is a warning to the shrimp and crayfish aquaculture. Our results provide evidence that EHP can infect and proliferate in the hepatopancreas of crayfish *P. clarkii*. The crayfish prefer to swim upstream, resting in lakes, rivers, reservoirs, swampy ponds and ditches, and sometimes in the fields. Combining these features, *P. clarkii* was able to approach shrimp ponds and spread EHP they carried, or healthy crayfish entered EHP infected shrimp ponds, become infected accidentally and carry EHP to other healthy ponds or their habitats. All these remind farmers to take certain measures to avoid direct contact between *L. vannamei* and *P. clarkii* in culture processes. For example, it is forbidden to exchange ponds between the two species, to prevent wild *P. clarkii* from entering shrimp ponds.

5. Conclusion

The study showed EHP can proliferate in the hepatopancreatic cells of *P. clarkii*,

294 and cause damage of hepatopaneas and intestine. Considering the large-scale
295 cultivation of *L. vannamei* and *P. clarkii*, overlap farming areas between these two
296 species as well as the ability of crayfish to crawl on land and water, our finding
297 indicates the potential role of *P. clarkii* in the transmission of EHP, and this alerts us
298 to the potentially greater risk of EHP in crayfish and shrimp aquaculture.

299 **Funding**

300 This research was funded by the Natural Science Foundation of Chongqing:
301 2022YSZX-JCX0010CSTB and cstc2021jcyj-cxttX0005.

302 **Authors' contribution**

303 Bingxin Ling: Writing - original draft, Methodology, Investigation. Yujiao Wu:
304 Writing - original draft, Methodology, and Validation. Qing Yu, Chunxia Wang and
305 Mengjiao Hu: Investigation. Xianzhi Meng: Resources and Investigation. Mengxian
306 Long: Conceptualization and Visualization. Guoqing Pan: Conceptualization, Writing
307 - review & editing and Funding acquisition. Zhonghuai Xiang: Conceptualization,
308 Funding acquisition. Zeyang Zhou: Supervision and Writing - review & editing. Jie
309 Chen: Supervision, Writing - review & editing and Funding acquisition.

310 **Declaration of Competing Interest**

311 The authors declare that they have no conflict of interest.

312 **Acknowledgments**

313 The authors are very grateful to Qing Lv for the help of staining the TEM
314 samples, and thank to Yuan Wang for revision of the manuscript.

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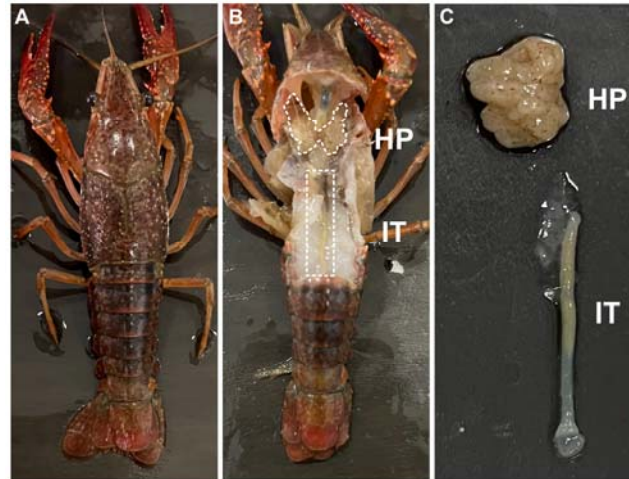
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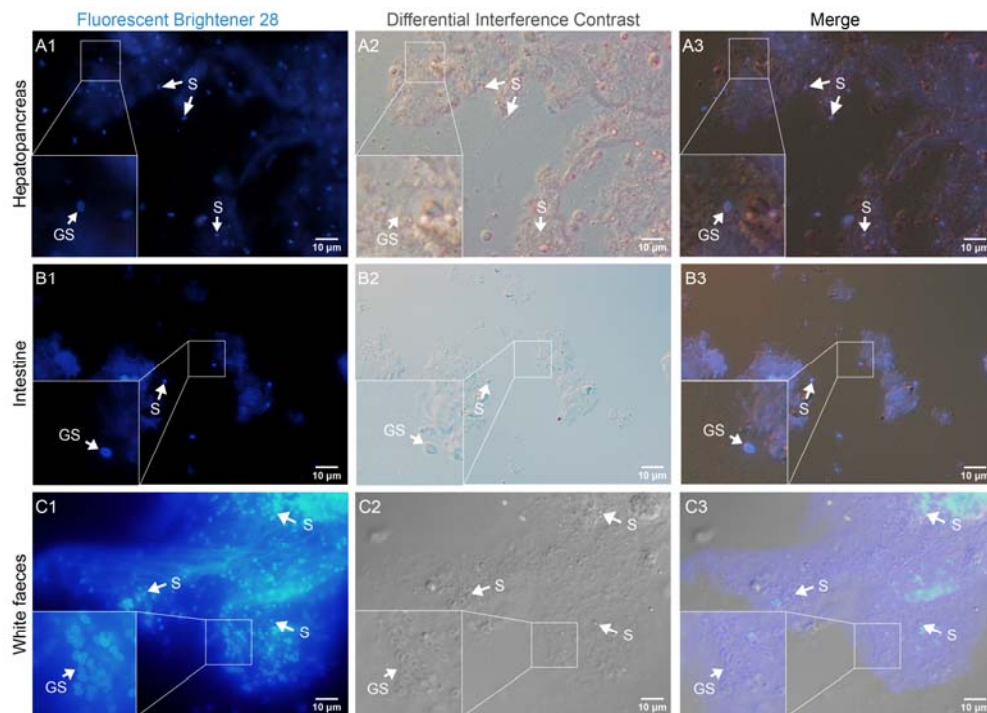
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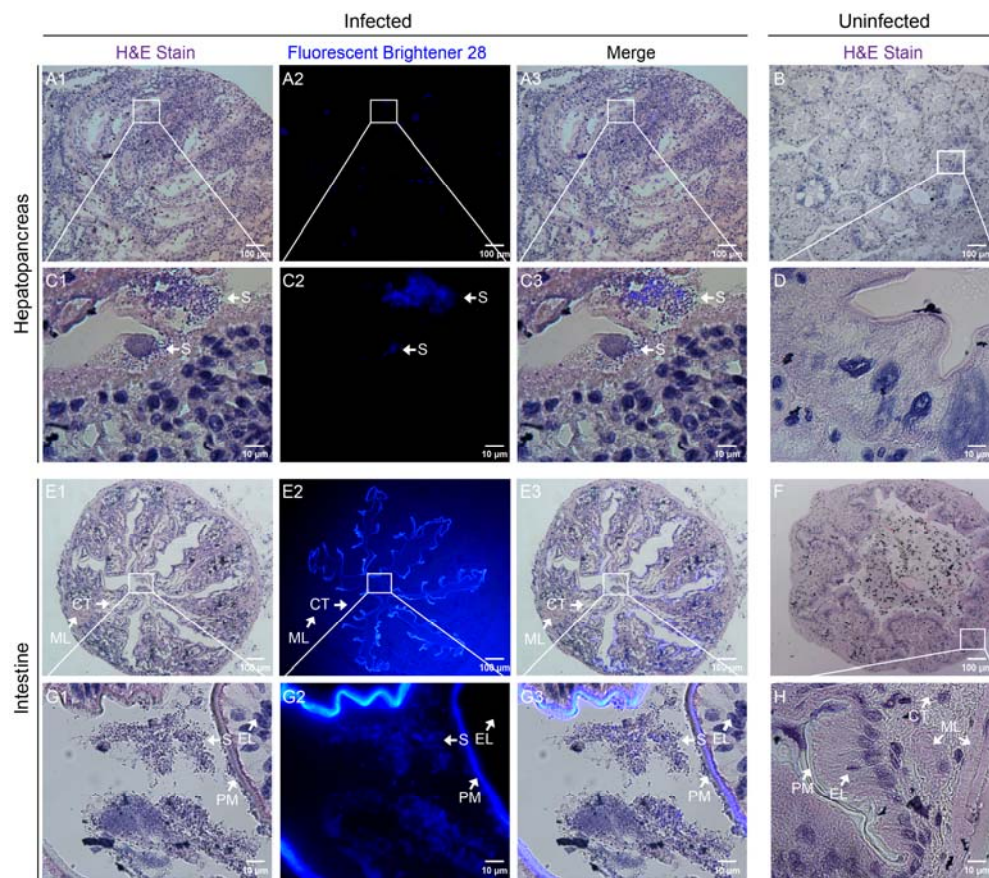
533 **Fig. 1 The anatomical morphology of digestive tract in infected *P. clarkii*.** The
534 external form of infected *P. clarkii* (A) had no difference with healthy ones, while the
535 anatomical digestive tract (B) was lighter in color. A large number of brown spots in
536 the hepatopancreas and white stuff in jejunum were observed (C). HP: hepatopancreas,
537 IT: intestine.



538

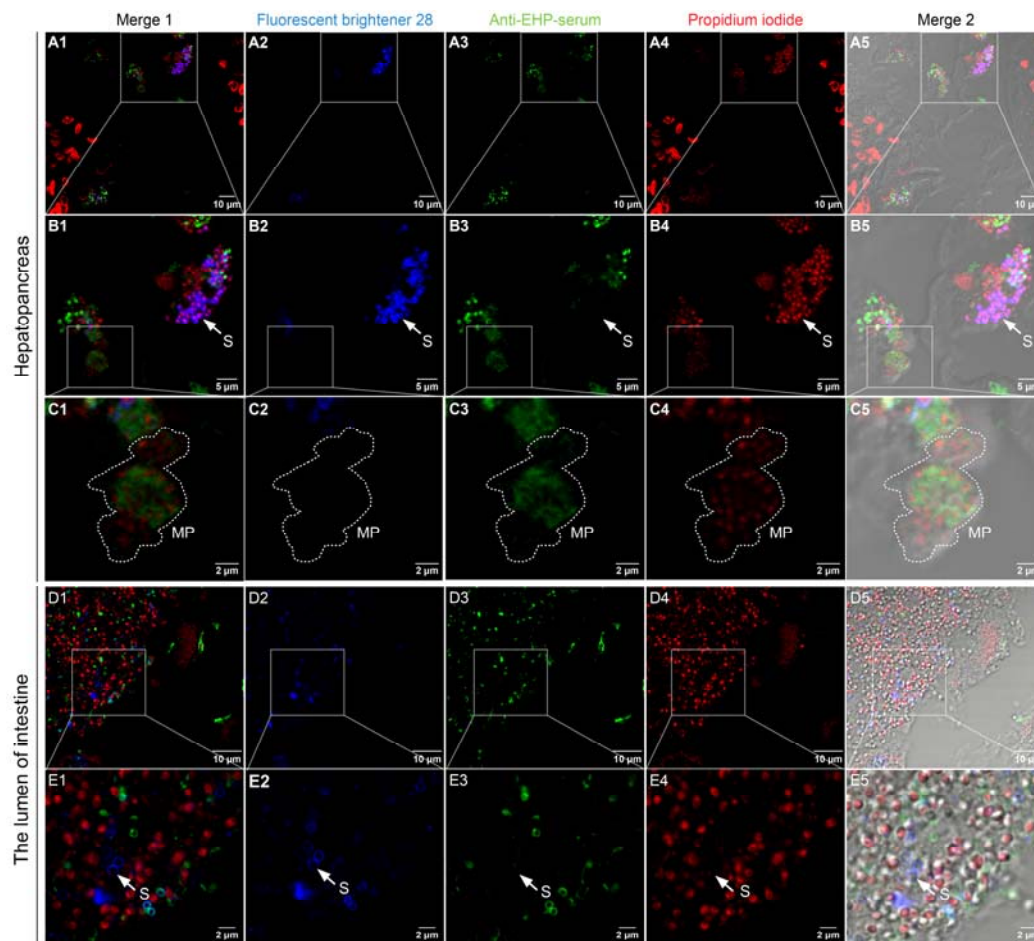
539 **Fig. 2 Fluorescence microscopic examination of hepatopancreas (A1-A3),**

546 *clarkii* by *EHPptp2* qPCR. Sequence alignments of *SSU rRNA* (C) and *EHPptp2* (D)
547 amplicons and reference. PC: positive control, PCR products amplified from EHP
548 infected *L. vannamei*; NC: negative control, no products were amplified from
549 EHP-free *P. clarkii*; NTC: no template control, ddH₂O was used as template;
550 SSU-rRNA: small subunit ribosomal RNA (GenBank: MNPJ01000021.1); ptp2: polar
551 tube protein 2 of EHP (GenBank: No. MT249228.1); Pc: *Procambarus clarkii*; Lv:
552 *Litopenaeus vannamei*; HP: hepatopancreas; IT: intestine; WF: white faces.



553
554 **Fig. 4 H&E-stain showed the tissue damage in the EHP infected *P. clarkii*.**
555 Abundant necrotic areas were seen in the infected hepatopancreas tissue and the
556 morphology of hepatopancreatic tubules were difficult to distinguish (A1-A3).

557 Clustered basophilic EHP spores were distributed among the necrotic areas (C1-C3)
 558 compared with those of EHP-free ones (B & D). The intestine of infected crayfish
 559 exhibited distinct epitheliolysis, enlarged interstices and sharpened pleated ridges
 560 (E1-E3 & G1-G3) compared with those of EHP-free ones (F & H). S: EHP mature
 561 spores; PM: peritrophic membrane; EL: epithelial layer; CT: connective tissue layer;
 562 ML: muscle layer.



563
 564 **Fig. 5 Developing stages of EHP in the hepatopancreas (A1-C5) and the lumen of**
 565 **intestine (D1-E5) of crayfish illustrated by indirect-immunofluorescent assay.**
 566 EHP was labeled with anti-EHP serum (green), chitin coats of spores were labeled
 567 using Fluorescent brightener 28 (blue) and nuclei was stained by Propidium iodide

(red). The regions selected by the rectangular boxes were magnified to the image below. Mature spores (B1-B5) and proliferative stage (C1-C5) of EHP can be seen in the hepatopancreas of crayfish. Different stages of EHP mix in the lumen of infected intestine (D1-E5). MP: merogonic plasmodia; S: EHP mature spores.

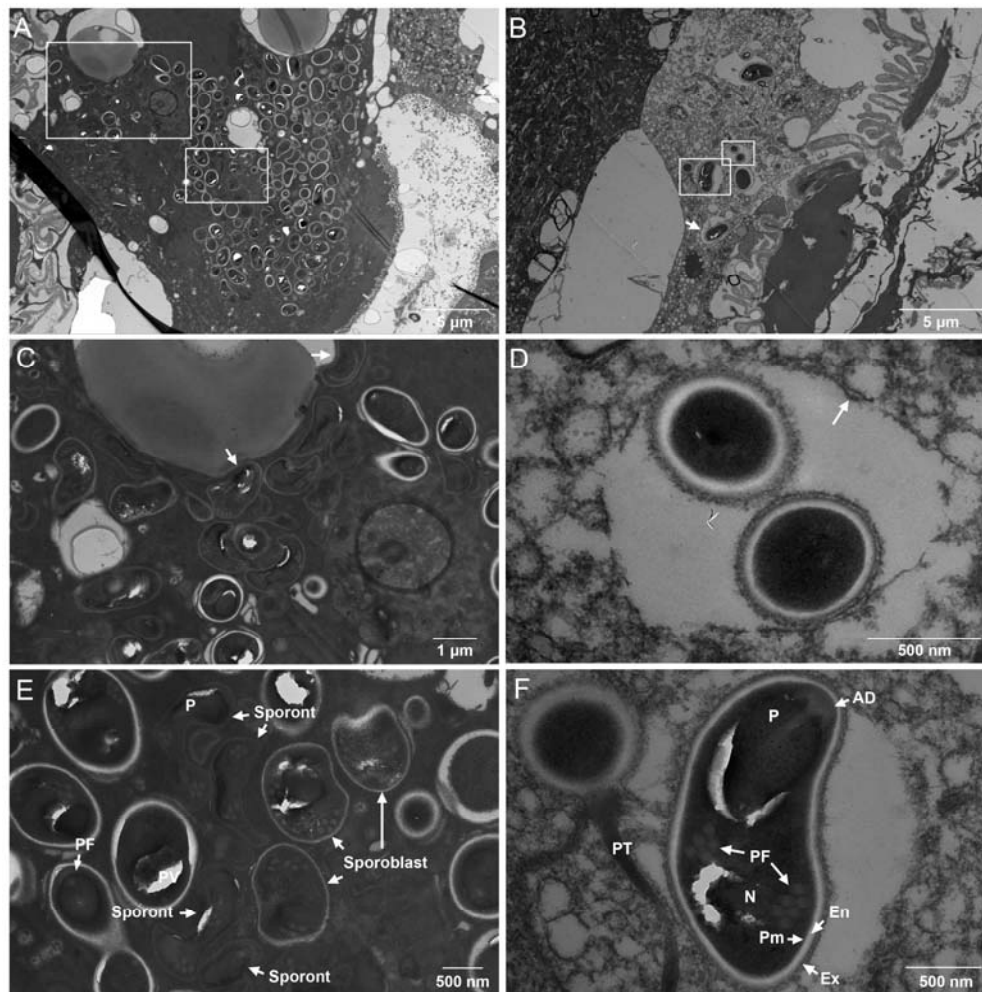


Fig. 6 Ultrastructure of sporogonic stages and mature spores of EHP in hepatopancreatic epithelia of crayfish. Abundant sporogonic and mature stage of EHP were distributed in the cytoplasm of hepatopancreatic epithelial cells. Sporogonic spores were found contacted closely to the membrane of host lipid droplet, and the membrane was deformed (A & C). Several spores were enveloped by a

membranous structure individually or in pairs (B & D). Mature spores were characterized by microsporidia ultrastructure with 5-6 coils of polar filament (F), while a thin spore wall was in the sporogonic phase of EHP (E). Image C and E, D and F were the magnification of the regions selected by the rectangular boxes in image A and B respectively. Ex: exospore; En: endospore; Pm: plasm membrane; P: polaroplast; PF: polar filament; PT: polar tube; N: nucleus; AD: anchoring disc.

TABLE 1 Primers used in this study.

Primer Name	Sequence (5`-3`)	Target Gene	References
EHP PTP2-F1	ATGAGTCTTTATAATGCACTG	<i>EHPptp2</i>	(Wang et al., 2020)
EHP PTP2-R1	TTATTCGTTGGATGTTAATG		
<i>EHPptp2</i> -192F	AATGGCTCAAGGGTTCA	<i>EHPptp2</i>	(Wu et al., 2022)
<i>EHPptp2</i> -192R	CTCCTGCCCTGTTTACG		
V1	CACCAGGTTGATTCTGCCTGAC	SSU rRNA of microsporidia	(Zhu et al., 1993; Baker et al., 1995)
530r	CCGCGGC(T/G)GCTGGCAC		
CRF	TCGCTGTAAAGTTGAAGAAGTT	Mitochondrial control region of <i>P. clarkii</i>	(Li et al., 2015)
CRR	TTAATCTCTTCATATCTTTAATTA		
LCO1490	GGTCAACAAATCATAAAGATATT	Mitochondrial cytochrome oxidase subunit I gene of metazoan invertebrates	(Folmer et al., 1994)
HCO2198	GG		
	TAAACTTCAGGGTGACCAAAAA		
	ATCA		