

1 **Black sea cucumber (*Holothuria atra* Jaeger, 1833) protects *Caenorhabditis elegans***
2 **against *Pseudomonas aeruginosa* infection**

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20 **Authors and respective contribution**

- 21 • **Wan-Ting Lee: Conducted killing assays, anti-virulence and gene expression**
22 **studies, draft of manuscript**
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- 24 • **Su-Anne Eng: Data analysis, manuscript preparation**
- 25 • **Gan Chee Yuen: Natural product extraction and fractionation**
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30 **supervision of WT Lee, BK Tan, S Eng, draft and finalization of manuscript**

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32

33 **Abstract:**

34 **Ethnopharmacological relevance:** Sea cucumbers are known to have beneficial effects
35 towards many ailments. Among traditional documented use of edible sea cucumbers are
36 as anti-infective tonic food. Metabolites with immunomodulatory properties have also
37 been reported from temperate sea cucumber species. The black sea cucumber
38 (*Holothuria atra*) is a widely distributed tropical marine sea cucumber species traditionally
39 consumed by Asian populations for its medicinal values.

40 **Aim of the study:** A strategy to circumvent the problem of multidrug resistant
41 *Pseudomonas aeruginosa* is discovery of novel anti-infectives targeting bacterial virulence
42 or host immunity. This study aims to decipher the anti-infective properties of *H. atra*.

43 **Materials and methods:** Here, we utilized the *Caenorhabditis elegans*-*P. aeruginosa*
44 infection model to assess the anti-infective properties of *H. atra*. We conducted the *C.*
45 *elegans*-*P. aeruginosa* slow killing assay to isolate a fraction with capable of conferring
46 survival to pathogen-infected worms, followed by a series of tests to assess the
47 anti-virulence and immune-related properties of the sea cucumber bioactive fraction.

48 **Results:** The *H. atra* methanol extract improved survival of infected worms without
49 affecting pathogen viability. Further extraction and assays sequestered a fraction with
50 significant anti-infective activities and was able to attenuate the secretion of virulence
51 factors and biofilm formation. Treatment of *P. aeruginosa*-infected transgenic *lys-7::GFP*
52 worms with this fraction also restored the repressed expression of host *lys-7*, a defense
53 enzyme. QTOF-LCMS analysis of this fraction revealed the presence of
54 aspidospermatidine, an indole alkaloid and inosine.

55 **Conclusion:** Collectively, results shows that *H. atra* confers survival advantage against *P.*
56 *aeruginosa* infection through inhibition of pathogen virulence and the restitution of host
57 *lys-7* expression.

58 **Keywords:** *Pseudomonas aeruginosa*; *Caenorhabditis elegans*; *Holothuria atra*;
59 anti-virulence; immunomodulator

60 **1. Introduction**

61 *Pseudomonas aeruginosa* is an opportunistic pathogen involved in urinary tract and
62 bloodstream infection, nosocomial pneumonia and contamination of surgical sites (Driscoll
63 et al., 2007; Taylor et al., 2014). Increasing number of antibiotic resistance *P. aeruginosa*
64 strains are being reported globally (Yayan et al., 2015). Attempts to overcome bacteria
65 resistance through modification of existing drugs or drug discovery are often ineffectual
66 (Parisien et al., 2008). Collectively, these issues underscore the need for alternative
67 therapies with different modes of targeting mechanism involving host immune response or

68 pathogen virulence. In comparison to direct killing of pathogens, these paths are projected to
69 invoke milder evolutionary pressure responsible for resistance development (Allen et al.,
70 2014).

71 The nematode *Caenorhabditis elegans* is a proven model for deciphering host–pathogen
72 interactions due to its readiness for infection by numerous human pathogens (Moy et al.,
73 2006). As a model organism, it is also supported by an array of tools useful for deciphering
74 immunity related pathways (Aballay and Ausubel, 2002). The *C. elegans* host-pathogen
75 interaction is a proven powerful platform for discovery of anti-infectives, and is frequently
76 utilized to screen for novel anti-infectives against a myriad of pathogens, (Conery et al.,
77 2014; Kim et al., 2017; Maria et al., 2017; Utari and Quax, 2013; Zhu et al., 2015). An
78 advantage in having both host and pathogen in a screening assay is early detection of
79 undesirable toxic properties in hit candidates (Squiban and Kurz, 2011).

80 The diverse marine environment impose the selection of an array of metabolites
81 produced by an immeasurable source of organisms (Molinski et al., 2009). Secondary
82 metabolites with anti-infective properties against *P. aeruginosa* have been identified from
83 numerous marine organisms including marine bacteria and seaweeds (Fatin et al., 2017;
84 Kandasamy et al., 2012; Kulshreshtha et al., 2016; Liu et al., 2013; Naik et al., 2013; Papa et
85 al., 2015). . To date, the only published report of an anti-infective from marine invertebrates
86 using the *C. elegans* platform is from a tropical mollusk (Kong et al., 2014).

87 The consumption of sea cucumber, a marine echinoderm as medicine can be traced back
88 to medical records documented during the Ming Dynasty (Chen 2003). Sea cucumbers
89 prepared for medicinal purposes are known as ‘gamat’ in Malaysia, with different forms of
90 gamat being utilized for various types of ailments (Fredalina *et al.* 1999). From the
91 estimated 1000 varieties of sea cucumbers worldwide, around 20 species have known
92 ethnopharmacological values (Bordbar et al., 2011). Asians typically regard sea cucumbers
93 as effective remedy of asthma, hypertension, joint pain, sprains intestinal and urinary
94 problems (Chen, 2004; Dakrory et al., 2015). Correspondingly, studies on sea cucumber
95 metabolites have reported multiple activities including anticancer, anticoagulant,
96 anti-hypertension, anti-inflammatory, antimicrobial, antioxidant, antithrombotic,
97 antihyperglycemic, anti-ageing and wound healing (Aminin et al., 2015; Bordbar et al., 2011;
98 Janakiram et al., 2015; Jattujan et al., 2018; Mourão et al., 1998; Song, M. et al., 2013). These
99 bioactivities are linked to a wide range of metabolites, such as triterpene glycosides, peptides,
100 collagen, mucopolysaccharides, carotenoids, polyunsaturated fatty acids and phospholipids

101 (Aminin et al., 2015; Bahrami and Franco, 2016; Ibrahim, 2012; Saito et al., 2002; Shi et al.,
102 2016).

103 *Holothuria atra* Jaeger 1833 (Holothuridae), colloquially known as black sea cucumber
104 or lollyfish, is one of the most commonly found sea cucumber species in the Indo-Pacific
105 region, dwelling in seagrass beds and rocky reefs (Conand, 2004). In Fiji, this species is
106 traditionally cooked or marinated before consumption (Vuki and Viala, 1990). Among the
107 many medicinal properties linked with sea cucumbers, sea cucumbers are also traditionally
108 consumed for its immune boosting properties (Chen, 2004). In tandem, metabolites with *in*
109 *vitro* or *in vivo* immunomodulatory activities have been isolated from several sea cucumber
110 species (Aminin et al., 2016; Aminin et al., 2008; Aminin et al., 2006; Song, Y. et al., 2013).
111 Therefore, sea cucumber is an opportune organism for discovery of novel and potent
112 anti-infective entities. Using the *C. elegans* host-pathogen platform, we investigated the
113 anti-infective properties of *H. atra*. Overall, we demonstrate the capacity of an extract from
114 this sea cucumber in rescuing worms from a lethal *P. aeruginosa* infection. The capacity of
115 *H. atra* metabolites in impeding the production of well-characterized *P. aeruginosa*
116 virulence factors and boosting an immune-related gene were also studied.

117 **2. Materials and Methods**

118 *2.1. Bacterial and Nematode Strains*

119 *P. aeruginosa* strain PA14 (Rahme et al., 1995), which is rifampicin-resistant, was
120 grown in King's B broth supplemented with rifampicin (100 µg/mL) while the
121 streptomycin-resistant *E. coli* strain OP50 (Brenner, 1974) was cultured in Luria Bertani
122 (LB) broth with streptomycin (100 µg/mL). Cultures were grown overnight with aeration at
123 37°C. The wild type *C. elegans* Bristol N2 (N2) and transgenic *lys-7::GFP* worm strain
124 SAL105 were obtained from the *Caenorhabditis* Genetics Center (CGC), USA, with growth
125 and handling as described elsewhere (Alper et al., 2007; Brenner, 1974).

126

127 *2.2. Preparation of Holothuria atra extract*

128 *Holothuria atra* was collected from Pulau Songsong Kedah situated at the northern
129 region of Peninsular Malaysia (N5°48'33.72" E100°17'32.63") and authenticated by the
130 Centre for Marine and Coastal Studies, Universiti Sains Malaysia. Samples were stored at
131 -20°C until further processing. Frozen samples were cleaned with distilled water, sliced into

132 smaller pieces and freeze-dried for about 5-7 days. Dried samples were extracted with either
133 methanol or acetone, followed by 15 minutes of sonication at room temperature and lastly,
134 filtration with Whatman filter paper No. 1. Extraction procedure was repeated thrice.
135 Filtrates were then concentrated using a rotary evaporator at 39°C. All dried extracts were
136 kept at -20°C before dilution in DMSO at 20mg/mL, followed by storage at 4°C.
137 Depending on assays, the extract stock solution is further diluted to achieve the desired
138 concentrations.

139 2.3. *C. elegans* Survival Assay

140 The slow killing assay was performed according to the protocol described elsewhere
141 (Dharmalingam et al., 2012). PA14 cultured overnight on King's B broth was evenly spread
142 on Pseudomonas Infection Agar (PIA) assay plates supplemented with natural products,
143 followed by incubation at 37°C for 24 hours to allow growth of PA14. Triplicate plates
144 containing 50-60 young adult worms each were used. At least three independent repeats were
145 performed.

146 Percentage survival of the infected worm population was determined after 48 hours of
147 incubation at 25°C. DMSO (0.5%) and curcumin (100 µg/mL), a known anti-infective agent,
148 was used as negative and positive control, respectively. (Rudrappa and Bais, 2008).

149 2.4. Antimicrobial Test

150 The minimum inhibitory concentration (MIC) broth microdilution antimicrobial test was
151 carried out to determine bactericidal activities of natural product. Stock solution of extracts
152 or fractions were further diluted in DMSO to achieve concentration ranging from 25 µg/mL
153 to 1000 µg/mL. Overnight culture of *P. aeruginosa* PA14 was adjusted to 0.5 McFarland
154 standard turbidity and diluted to 1:20. A total of 10 µL of inoculum was added into the wells,
155 followed by incubation at 37°C for 24 hours. Streptomycin (100 µg/mL) and DMSO (0.5%)
156 were also tested as positive and negative controls, respectively.

157 2.5. Bacterial Growth Assay

158 The assay was conducted with similar preparations as the MIC broth microdilution test.
159 Upon inoculum addition, the well plate was incubated at 37°C with aeration and bacterial cell
160 density was measured at OD₆₀₀ every 2 hours.

161 2.6. Liquid-liquid Partitioning of Methanolic *H. atra* Extract

162 The methanol extract was subjected to liquid-liquid chromatography, to separate
163 compounds according to polarity. The extract was suspended in distilled water and
164 subsequently partitioned with butanol. The partition was collected and dried with a rotatory
165 evaporator, followed by storage at -20°C. Stock solution for the partition was prepared in
166 DMSO and stored at 4°C.

167 2.7. Fractionation of *H. atra* Butanol Partition

168 The butanol partition was column chromatographed over silica gel 60 (70-230 mesh) and
169 eluted stepwise from 100:0:1 to 0:100:1 with the chloroform:methanol:water solvent system
170 at ratio of 80:20:1. The process was repeated until sufficient fractions were obtained for
171 further analysis.

172 2.8. Virulence Factors and Biofilm Production Assays

173 Protease and elastase activity assays were performed as described elsewhere, with minor
174 modifications (Prithiviraj et al., 2005). Briefly, *P. aeruginosa* PA14 was grown in the
175 presence of bioactive fraction at 200 µg/mL or DMSO (0.5%) for 24 hours at 37°C. The
176 supernatants were collected and filtered with a 0.22 µm nylon filter. Reaction mixture
177 consisting of 0.8% azocasein in 500 µL of 50mM K₂HPO₄ (pH7) and 100 µL of the purified
178 supernatant were incubated at 25°C for 3 hours. A total of 0.5 mL of 1.5M HCl was added
179 into the reaction, followed by placement on ice for 30 minutes and centrifugation. After the
180 addition of 0.5 mL 1M NaOH, the absorbance was measured at 400nm (SpectraMax M5).

181 As for elastase activity, 100 µL of the purified supernatant described above was added
182 into tubes pre-prepared with 1 mL of 10mM Na₂HPO₄ and 20 mg of elastin-Congo red. The
183 mixtures were then incubated at 37°C for 4 hours with agitation. After the incubation, the
184 mixtures were centrifuged at 14 000 rpm for 10 minutes and absorbance determined at
185 495nm.

186 The pyocyanin quantification assay was conducted as described (Essar et al., 1990). *P.*
187 *aeruginosa* PA14 was grown in the presence of bioactive fraction for up to 24 hours at 37°C.
188 Cultures were collected at 6, 12 and 24 hours after treatments and centrifuged. A total of 3
189 mL chloroform was then added into the resulting supernatant, vortexed briefly and
190 centrifuged for 10 minutes at 10 000 rpm. The bottom blue layer was isolated, and the
191 extraction processes were repeated using 1.5 mL of 1M HCl. The absorbance at 520 nm of

192 the pink layer was measured. Pyocyanin concentration was determined by multiplying the
193 absorbance value with 17.072.

194 A microtiter biofilm formation assay was used to determine the effect of *H. atra* fraction
195 on *P. aeruginosa* PA14 biofilm formation (O' Toole, 2011). *P. aeruginosa* PA14 was grown
196 overnight followed by dilution to 1:100. A total of 100 μ L of the inoculum was added into 96
197 wells plate preseeded with *H. atra* fraction or 0.5% DMSO. The plate was then incubated for
198 24 hours at 37°C. Liquid and cells were discarded, followed by rinsing of plate with distilled
199 water. For visualisation, 125 μ L of 0.1% crystal violet solution was added into each well,
200 followed by incubation for 15 minutes at room temperature. Excessive stain was washed off
201 with water and wells left air-dried for 2 hours. Stained biofilm was treated with 125 μ L of
202 30% acetic acid followed by 10-15 min incubation. A total of 125 μ L of solubilized crystal
203 violet was transferred into a new microtiter for measurement at 550 nm .

204 2.9. Visualization of Transgenic *lys-7::GFP* *C. elegans*

205 Transgenic *C. elegans lys-7::GFP* strain SAL105 were exposed *P. aeruginosa* PA14 in
206 the presence or absence of *H. atra* fractions as described in the slow killing assay. The
207 fluorescence intensities of young adult worms at 12 hpi for all the experimental conditions
208 were visualized using Leica Microsystem M205 FA microscope and digitally captured with
209 Leica DFC310 FX camera (60x magnification)

210 2.10. Real-time PCR based quantification of *lys-7* expression in *C. elegans*

211 Primers were designed accordingly for the amplification of *C. elegans lys-7* and
212 pan-actin as housekeeping gene. Total RNA was treated with RNase-free DNase (Promega,
213 USA) for removal of genomic DNA. PCR optimization and amplification was performed
214 under condition described previously (Dharmalingam et al., 2012). Melt curve analysis was
215 performed to ensure the specificity of the PCR products. Quantitative expression of *lys-7*
216 was calculated using CFX Manager software (BioRad, USA) by normalizing its level to
217 housekeeping gene. Three independent replicates were performed for each analysis.

218 2.11. Chemical Profiling of Fractions

219 Compounds identification of both the *H. atra* active fractions were performed using the
220 Agilent 6520 accurate-mass quadrupole time-of-flight liquid chromatography mass
221 spectroscopy (QTOF-LCMS). Zorbax SB-C18 column (1.5 μ m particle size, 0.5x150mm)
222 was used and separation was achieved with a 22.5 minutes gradient of 3-90% acetonitrile,

223 followed by 3% acetonitrile for 10 minutes at a flow rate of 20 μ L/min. Solvents contained
224 0.1% formic acid. The compound prediction was achieved by comparing the obtained
225 individual MS/MS fingerprinting spectrum with METLIN, MassBank and Chemspider
226 databases.

227 2.12. Statistical Analysis

228 Data were analyzed using Student's *t*-test whereby in all comparisons, $p < 0.05$, $p < 0.01$
229 and $p < 0.001$ were accepted as statistically significant.

230

231 3. Results

232 3.1. *H. Atra* Promotes Survival of *P. aeruginosa* PA14 Infected *C. elegans* Without 233 Displaying Any Direct Bactericidal Effect

234 The slow-killing assay showed methanol extract of *H. atra* causing a significantly higher
235 percentage of worm survival ($60.21 \pm 6.7\%$) at 48 hours after treatment, which is comparable
236 to value obtained with curcumin ($60.68 \pm 8.6\%$), a known anti-infective against PA14
237 infection (Figure 1). A lower percentage of survival ($56 \pm 7.6\%$) was obtained with acetone
238 extract of *H. atra*.

239 Next, we determined the bactericidal effect of the methanol extract on *P. aeruginosa*
240 PA14. Extract concentrations ranging from 25 μ g/mL to 1000 μ g/mL did not reduce *P.*
241 *aeruginosa* PA14 growth, as compared to streptomycin at 100 μ g/mL (Supplementary 1).
242 This extract was also non-detrimental towards the growth kinetic of *P. aeruginosa* PA14, as
243 no significant difference was observed between *H. atra* and DMSO exposed bacteria
244 population (Figure 2).

245 The lethality of *P. aeruginosa* PA14 infection in the slow killing assay involves the
246 colonization of *C. elegans* intestinal tract (Tan et al., 1999). Therefore, we ascertain if the
247 improved survival of worms treated with extract is an outcome of reduced worm feeding
248 activities. We observed the worm pharyngeal pumping rate, a reliable indicator of
249 bacterial feeding in this species (Kong et al., 2014). Results show no significant difference
250 in feeding activities between OP50 fed worms and untreated PA14 fed worms, 0.5%
251 DMSO treated PA14 fed worms or *H. atra* methanol treated PA14 fed worms, respectively

252 (Figure 3). Therefore, worms exposed *H. atra* methanol extract still display normal
253 feeding activities, leading to colonization of gut by PA14.

254 Column chromatography fractionation of the *H. atra* butanol partition resulted in 9
255 fractions (F1-F9). Based on the slow-killing assay, fractions F2 ($55.37 \pm 8.4\%$), F3 ($64.05 \pm$
256 8.8%), F4 ($52.75 \pm 6.3\%$) and F9 ($52.00 \pm 8.0\%$) resulted in significantly higher worm
257 survival as compared to 0.5% DMSO treatment (Figure 4a). Fraction F3 also did not appear
258 to perturb *P. aeruginosa* PA14 growth (Figure 4b).

259 Dose dependent assay for fraction F3 showed that 200 $\mu\text{g/mL}$ was the minimal
260 concentration resulting in significantly highest percentage of worm survival, comparable to
261 value obtained with curcumin (Figure 5). At higher concentrations, a decreasing trend of
262 survival rates was observed.

263 3.2. *H. atra* fraction Inhibits the Production of *P. aeruginosa* PA14 Elastase, Protease, 264 Pyocyanin and Formation of Biofilm

265 In comparison with 0.5% DMSO, treatment of *P. aeruginosa* PA14 with F3 significantly
266 suppressed the production of elastase and protease at 24 hr after treatment (Figure 6a).
267 Formation of biofilm in microtiter plate was also attenuated with F3. As for pyocyanin,
268 significant reduction in pyocyanin production was obtained at 6 and 12 hr after incubation,
269 respectively, but not 24 hr. (Figure 6b).

270

271 3.3. *H. atra* Induces the Expression of *C. elegans* Immune Effector *lys-7* in PA14 Infected 272 Worms

273 At 12 hr after infection, GFP intensity of *P. aeruginosa* PA14 infected worms was lower
274 than uninfected worms, indicating diminished *lys-7* expression (Figure 7a,b). Treatment
275 with *H. atra* restored *lys-7* expression to a level comparable with uninfected worms (Figure
276 7c). Correspondingly, qPCR analysis of *lys-7* in worms also showed higher expression
277 levels in worms exposed to F3 at 12 or 24 hours after treatment (Figure 7d). *H. atra* fraction
278 also resulted in higher *lys-7* expression as compared to curcumin.

279 2.4. Tentative identification of major compound in *H. atra* F3 and F4 Fractions

280 QTOF-LCMS separation detected two major peaks in F3 at Retention Times (R_T) 17.764
281 min and 18.395. The compounds were identified using an HR MS (accurate mass in

282 negative mode) and HR MS/MS spectra (fragmentation) that matched with METLIN,
283 MassBank and Chemspider databases. From the accurate mass analysis (Table 1), the
284 molecular formula was calculated as $C_{18}H_{22}N_2$ and $C_{10}H_{12}N_4O_5$, respectively. The
285 compounds were putatively identified as aspidospermatidine and inosine.

286 **4. Discussion**

287 An ongoing strategy to overcome the problem of drug-resistant pathogens is discovery
288 of anti-infective activity, which either disrupt virulence-related pathways (Clatworthy et al.,
289 2007) or boost host immunity (Hamill et al., 2008). Marine natural products with
290 anti-virulence (Naik et al., 2013; Papa et al., 2015) or immunomodulatory properties
291 (Kandasamy et al., 2012) have been reported. Among edible marine organisms, sea
292 cucumber is regarded as a potent source of novel entities useful for both nutraceutical and
293 pharmaceutical industries. There is an increasing reports of bioactivities from *H. atra*, a
294 widely distributed tropical sea cucumber species. This includes antibacterial, antifungal,
295 anti-proliferative, anti-apoptotic and antioxidant properties (Dakrory et al., 2015;
296 Dhinakaran and Lipton, 2014; Dhinakaran and Lipton, 2015; Esmat et al., 2013; Nishikawa
297 et al., 2015; Ridzwan et al., 1995; Saad et al., 2016).

298 Our results identified a sea cucumber *H. atra* methanol extract with capacity to confer
299 survival in *P. aeruginosa* PA14-infected *C. elegans*. This extract does not possess direct
300 bactericidal activity nor impedes growth of *P. aeruginosa* PA14, which rules out bacteria
301 mortality as reason for improved worm survival. It was previously reported that a methanol
302 extract of *H. atra* did not exert any bactericidal effect on several pathogenic bacteria species,
303 including *P. aeruginosa* (Lawrence et al., 2010). Taken together with other investigations, *H.*
304 *atra* bioactivities is therefore dependent on extraction solvent and target pathogen species
305 (Abraham et al., 2002; Ibrahim, 2012). Further separation of the *H. atra* methanol extract
306 yielded a fraction with similar bioactive properties. The range of worm survival obtained
307 with this fraction is comparable to levels using a similar slow-killing assay where infected
308 worms were treated with a terrestrial plant seed extract (Dharmalingam et al., 2012) or
309 curcumin (Rudrappa and Bais, 2008). These levels however, are lower than a fraction
310 isolated from a tropical marine actinomycete (Fatin et al., 2017).

311 In order to further understand the mechanisms responsible for the survival of infected
312 worms, we assessed the effect of *H. atra* towards several *P. aeruginosa* PA14 virulence

313 factors. Although the complete spectrum of the pathogenesis of *C. elegans*-*P. aeruginosa*
314 slow killing assay is not fully understood, the involvement of multiple virulence factors such
315 as pyocanin, pyoverdine, various proteases, elastase, phospholipase and exotoxin is
316 canonical (Kipnis et al., 2006; Rahme et al., 1995; Tan et al., 1999). Our results indicate *H.*
317 *atra* lowered proteolytic and elastolytic activities, diminished the production of pyocyanin
318 and the formation of biofilm, collectively signifying attenuation of *P. aeruginosa* PA14
319 virulence. In addition, the collective mitigation of these virulence factors suggests a possible
320 interference of PA14 quorum sensing (QS) by *H. atra* metabolites. QS have been
321 corroborated as one of the crucial factor responsible for host mortality in the *C. elegans* slow
322 killing assay (Papaioannou et al., 2009). Elsewhere, a marine seaweed extract was reported to
323 repress *P. aeruginosa* QS genes, leading to diminished production of virulence factors and
324 survival of infected worms (Kulshreshtha et al., 2016). Future work should ascertain the
325 capacities of *H. atra* metabolites in interfering with pivotal elements of the *P. aeruginosa*
326 QS.

327 The bioactive fraction was also able to restore the expression of *lys-7*, which encodes an
328 immune-specific lysozyme which facilitates the clearance of bacterial infection through
329 disruption of bacterial cell wall peptidoglycans (Evans et al., 2008; Mallo et al., 2002). Two
330 plausible reasons could explain the restitution of *lys-7*. A known characteristic of *P.*
331 *aeruginosa* infection in *C. elegans* is the suppression of host *lys-7* through the subversion of
332 the DAF-2/DAF-16 insulin-like signaling pathway, leading to the downstream
333 transcriptional suppression of several antimicrobial factors, including *lys-7* (Evans et al.,
334 2008; Garsin et al., 2003; Murphy et al., 2003). Since the *H. atra* F3 fraction could mutually
335 reduce the production of virulence factors and restore expression of *lys-7* in infected worms,
336 it is possible that the latter observation is a consequence of halting the *P. aeruginosa*
337 virulence activities upstream of the DAF2-DAF16 pathway. This would mimic the effects
338 exerted by a marine brown seaweed extract in overcoming the pathogenicity of PA14
339 towards *C. elegans* (Kandasamy et al., 2012). Alternately, *H. atra* could also contain
340 compounds capable of directly inducing *lys-7* activities, as reported elsewhere (Fatin et al.,
341 2017; Li et al., 2014).

342 Compounds with immunomodulatory activities have been isolated from edible sea
343 cucumber species such as *Cucumaria japonica*, *C. frondosa* and *Stichopus japonicus*
344 (Aminin et al., 2016; Aminin et al., 2008; Aminin et al., 2006; Song, Y. et al., 2013).
345 Among these anti-infective molecules, polysaccharides are the most studied group (Kale et

346 al., 2013; Song, Y. et al., 2013; Wang et al., 2017). In addition, some triterpene glycosides,
347 which are prominent secondary metabolites in sea cucumbers, also possessed
348 immune-boosting effects (Bahrami and Franco, 2016). Two examples, cucumariosides and
349 frondoside, were also reported to promote cellular immunity through stimulation of
350 phagocytosis and lysosomal activities (Aminin et al., 2016; Aminin et al., 2001; Aminin et
351 al., 2008). In our case, the main constituents detected in the bioactive *H. atra* fraction are
352 aspidospermatidine and inosine. Aspidospermatidine is an indole alkaloid, a large and
353 growing group of potentially useful marine-based metabolites (Netz and Opatz, 2015).
354 Elsewhere, alkaloids have been identified in *H. atra* extract (Dhinakaran and Lipton, 2014).
355 Indole and its derivatives have been demonstrated to repress the *P. aeruginosa* QS system,
356 leading to diminished production of protease, pyocyanin and biofilm (Kim and Park, 2013;
357 Lee et al., 2009; Lee et al., 2012). Inosine, a deaminated product of adenosine, was initially
358 speculated as not having useful biological effects (Hasko et al., 2004). However, various
359 studies have highlighted immunomodulatory and neuroprotective properties in inosine
360 (Benowitz et al., 2002; Chen et al., 2002; Jin et al., 1997).

361 **5.0 Conclusion**

362 In conclusion, we propose that *H. atra* has anti-infective properties against *P.*
363 *aeruginosa*. A bioactive fraction of this sea cucumber species inhibits the production of
364 pathogen virulence factors. Concomitantly, *H. atra* also restored the expression level of host
365 *lys-7*. High levels of an indole alkaloid and inosine were detected in this bioactive fraction.

366

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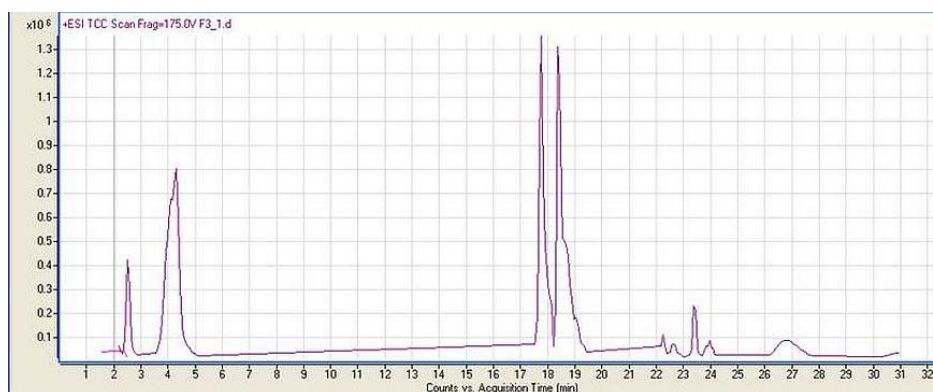
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1 **Table 1.** Aspidospermatidine and inosine, the main compounds identified in F3
 2 fraction of *Holothuria atra*

ESI-MS RT (min)	Detected ion (<i>m/z</i>) [M-H] ⁻	Fragments	Molecular Formula	Identification
17.764	266.1789	266, 144, 136, 130 268, 215,	C ₁₈ H ₂₂ N ₂	Aspidospermatidine
18.395	268.194	185, 136, 133, 109, 93, 82, 54	C ₁₀ H ₁₂ N ₄ O ₅	Inosine

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9 **Supplementary 1**

10 The antimicrobial effect of *Holothuria atra* on *P. aeruginosa* PA14 as determined by Minimum
 11 Inhibitory Concentration (MIC) test*.

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Samples	Concentration (µg/mL)				
	25	50	100	500	1000
<i>Holothuria atra</i> -methanol	-	-	-	-	-
<i>Holothuria atra</i> -fraction F3	-	-	-	-	-
Streptomycin**			+		

13 * + denotes inhibition; - denotes no inhibition of PA14 growth

14 ** 100 µg/ml streptomycin was used as control

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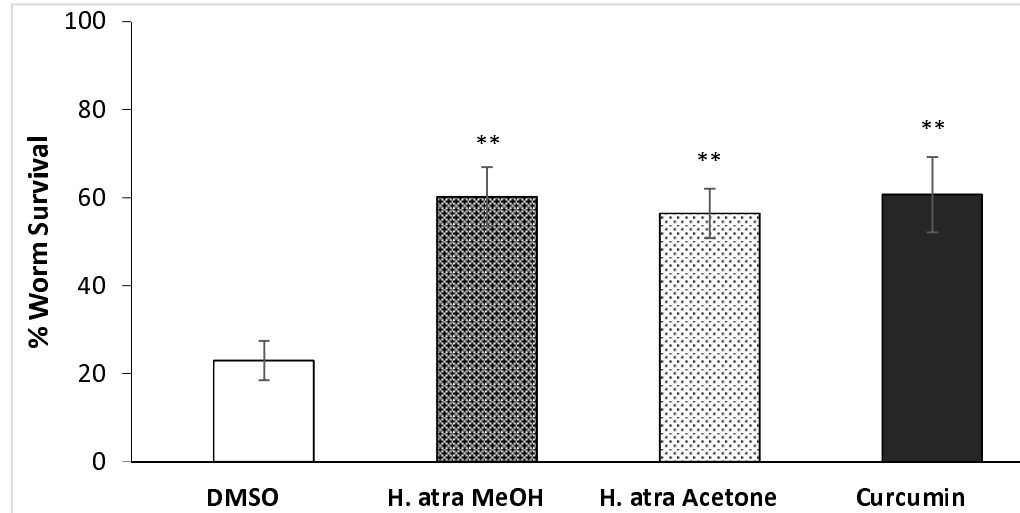
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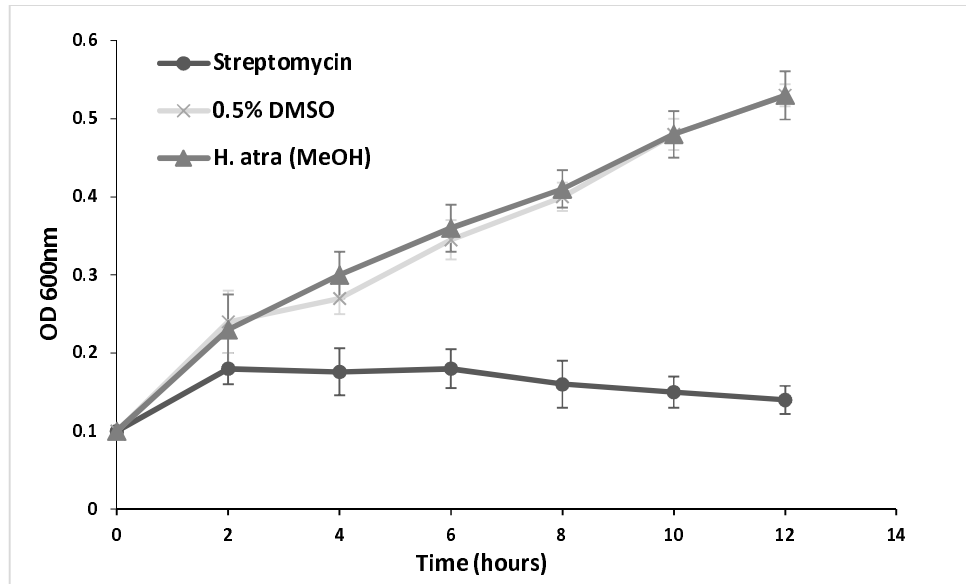
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Figure 1. Improved survival of *P. aeruginosa* PA14-infected *C. elegans* at 48 hours after treatment with *Holothuria atra* extracts (200 μ g/mL). The methanol extract showed the highest percentage of survival. Data were analyzed using Student's *t*-test where ** denotes significance at $p < 0.01$ when compared to 0.5% DMSO treatment. Curcumin was used as positive control.



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21 **Figure 2.** Growth rate of *P. aeruginosa* PA14 in the presence of 200 $\mu\text{g/mL}$
22 *Holothuria atra* methanolic extract was comparable to 0.5% DMSO treatment while
23 streptomycin (100 $\mu\text{g/mL}$) reduced bacterial proliferation.

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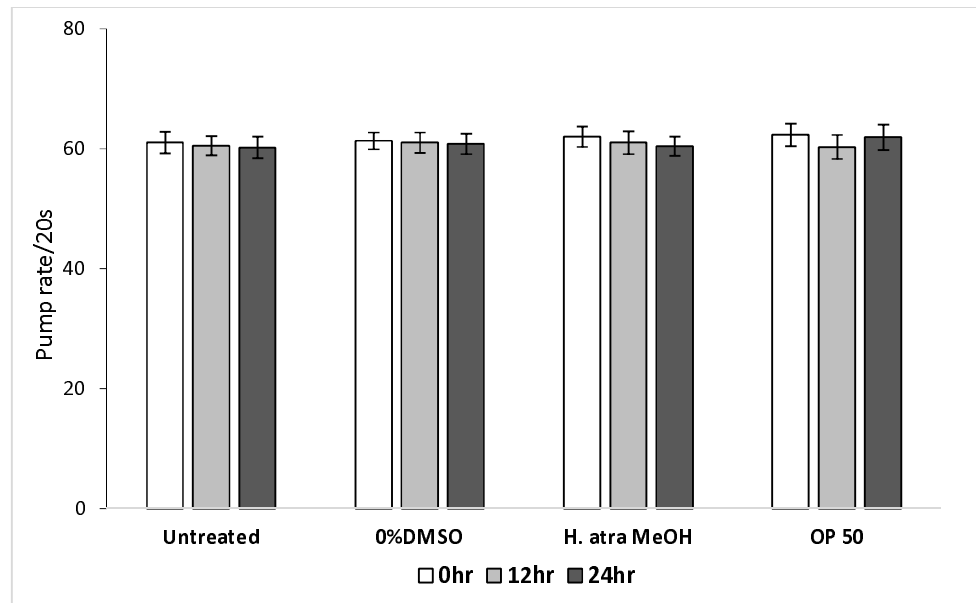
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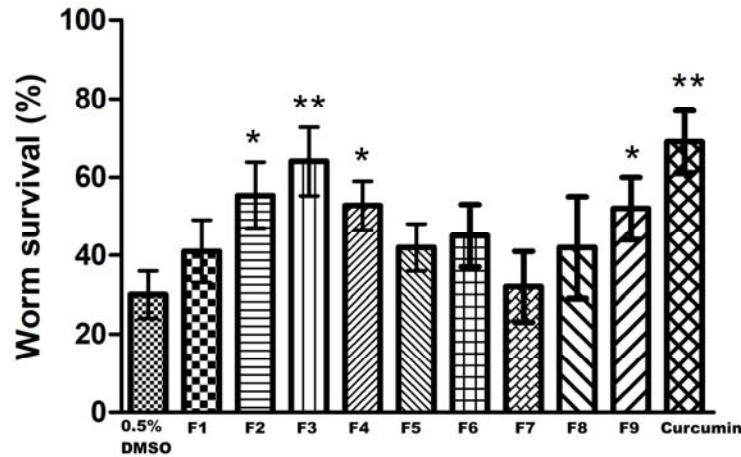


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Figure 3. Pharyngeal pumping rate of *C. elegans* fed OP-50, fed PA14 (untreated), fed PA14 (0.5% DMSO treated) and fed PA14 (*Holothuria atra* methanol extract at 200 $\mu\text{g}/\text{mL}$) at 0, 12 or 24hr after treatment.

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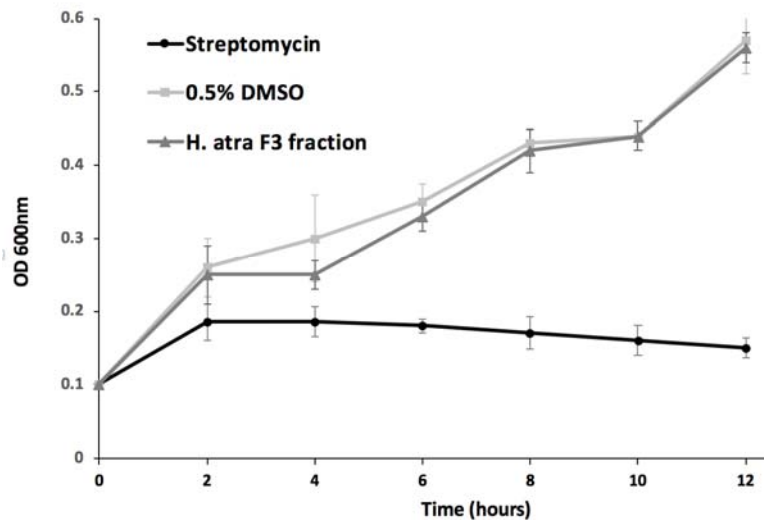
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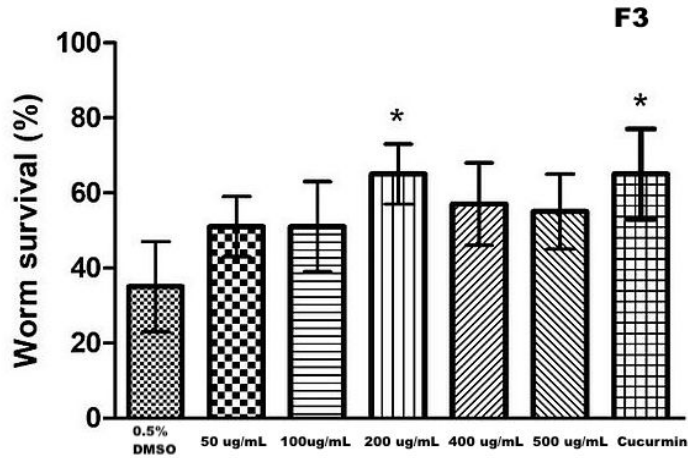
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Figure 4. a) Survival of *P. aeruginosa* PA14-infected *C. elegans* at 48 hours after treatment with respective *Holothuria atra* fractions. The F3 fraction showed the highest percentage of survival. Data were analyzed using Student's *t*-test where * and ** denotes significance at $p < 0.05$ and $p < 0.01$ respectively when compared to 0.5% DMSO treatment. Curcumin, used as positive control, also prolonged worm survival significantly. b) Growth rate of *P. aeruginosa* PA14 in the presence of 200 µg/mL *Holothuria atra* fraction F3 was comparable to 0.5% DMSO treatment (b). Streptomycin (100 µg/mL) reduced bacterial proliferation.



93 **Figure 5.** Dose dependent response of *P. aeruginosa* PA14-infected *C. elegans* at
94 48 hours after treatment with *Holothuria atra* fraction F3. Data were analyzed using
95 Student's t-test where * at $p < 0.05$ when compared to 0.5% DMSO treatment.
96 Curcumin, as positive control, also prolonged worm survival significantly.

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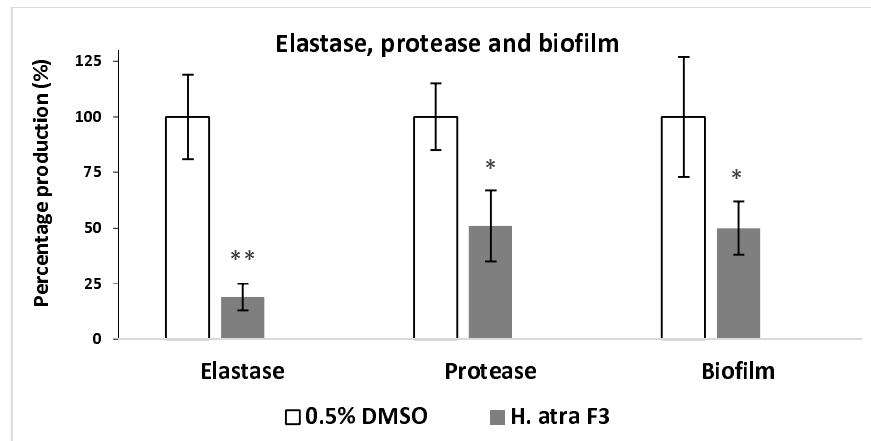
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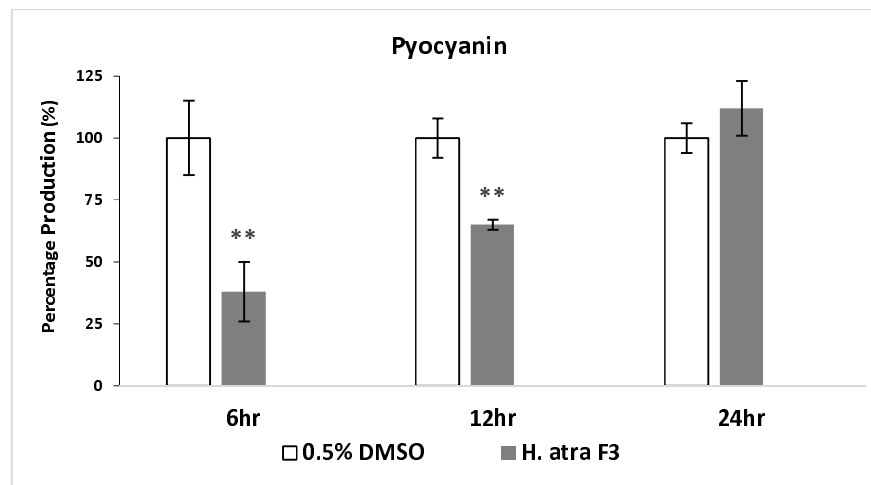
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111 **Figure 6** Effect of *Holothuria atra* F3 fraction on production of *P. aeruginosa*
112 PA14 virulence factors (a) Reduced production elastase, protease and biofilm at 24
113 hr after exposure to F3 (b) Reduced pyocyanin production at 6 and 12 hr after
114 exposure. Data were analyzed using Student's *t*-test where * and ** denotes
115 significance at $p<0.05$ and $p<0.01$ respectively when compared to 0.5% DMSO
116 treatment.

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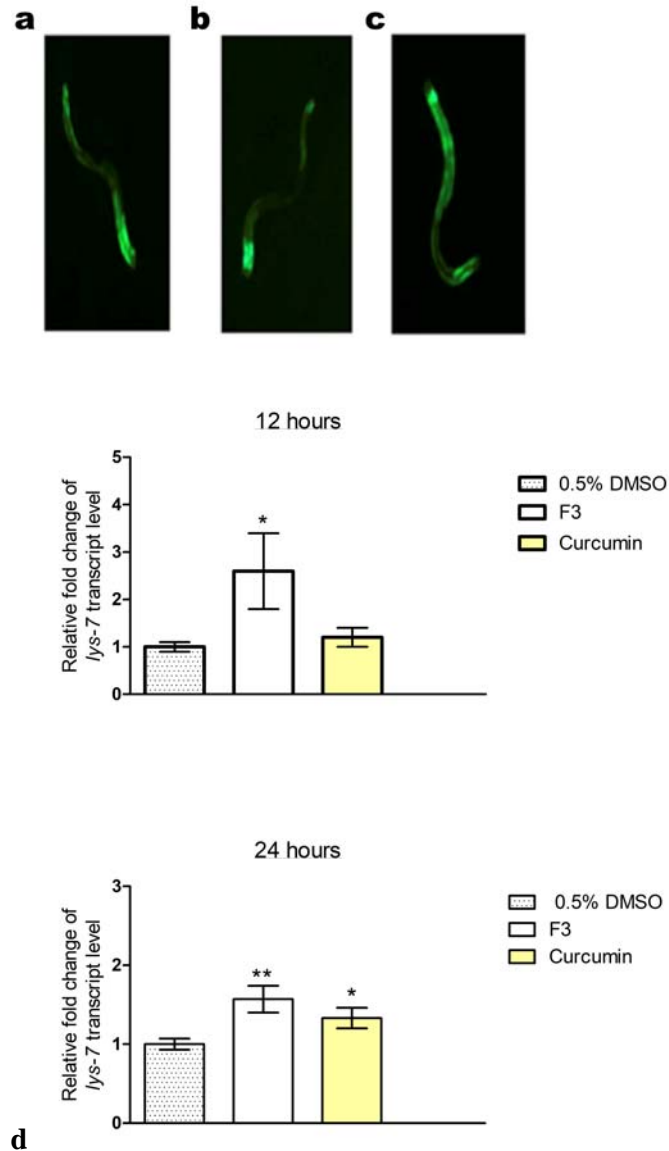
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125 **Figure 7.** *Holothuria atra* restore *lys-7* expression in *P. aeruginosa* PA14 infected
126 *C. elegans*. Representative fluorescence micrographs (60x magnification) of
127 transgenic *lys-7::GFP* worms (a) fed with *E. coli* OP50 (b) infected with *P.*
128 *aeruginosa* PA14 and treatment with 0.5% DMSO (c) infected with *P. aeruginosa*
129 PA14 and treatment with F3. Images were captured at 12 hr after treatment. (d)
130 qPCR of *C. elegans lys-7* in PA14 infected worms treated with 0.5% DMSO, *H. atra*
131 F3 fraction or curcumin at 12 or 24 hrs. Data were analyzed using Student's *t*-test
132 where * and ** denotes significance at $p < 0.05$ and $p < 0.01$ respectively when
133 normalized to 0.5% DMSO treatment.

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