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Assessment of the effects of sulfated polysaccharides extracted from the red seaweed Irish moss *Chondrus crispus* on the immune-stimulant activity in mussels *Mytilus spp.*

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1 Abstract

2 Seaweeds contain a number of health enhancing and antimicrobial bioactive compounds
3 including sulfated polysaccharides (SP). In the present study, SP extracted from a European
4 red seaweed Irish moss *Chondrus crispus* was chemically analyzed, SP content extracted and
5 the immune-response effect on wild Irish mussels *Mytilus* spp. investigated for the first time.
6 A high percent yield of SP was extracted from *C. crispus* and the immune-stimulant activity
7 of SP was assessed in a laboratory trial with mussels exposed to three different treatments of
8 low ($10 \mu\text{g mL}^{-1}$), medium ($20 \mu\text{g mL}^{-1}$) and high ($50 \mu\text{g mL}^{-1}$) SP dose concentrations and a
9 control mussel group with no exposure to SP. An initial mussel sample was processed prior to
10 the trial commencing and mussels were subsequently sampled on Days 1, 2, 3, 4, 7, and 10
11 post SP exposure. Both cell, humoral and immune related gene responses including
12 haemocyte cell viability, haemocyte counts, lysozyme activity and expression of immune
13 related genes (defensin, mytimycin and lysozyme mRNA) were assessed. No mussel
14 mortalities were observed in either the treated or non-treated groups. Mussels exposed with
15 SP showed an increase in haemocyte cell viability and the total number of haemocytes
16 compared to control mussels. Lysozyme activity was also higher in treated mussels.
17 Additionally, up-regulated expression of defensin, mytimycin and lysozyme mRNA was
18 observed in SP treated mussels shortly after exposure (on Days 1, 2, and 3) to SP. These
19 results indicate that a high quality yield of SP can be readily extracted from *C. crispus* and
20 more importantly based on the animal model used in this study, SP extracted from *C. crispus*
21 can rapidly induce health enhancing activities in *Mytilus* spp. at a cellular, humoral and
22 molecular level and with a prolonged effect up to ten days post treatment.

23 **Keyword:** *Chondrus crispus*; immune responses; *Mytilus spp*; sulfated polysaccharides

24

25 1. Introduction

26 Marine natural products have received much attention in recent years due to their
27 health benefits and bioactivity. More than 16,000 natural products have been isolated from
28 marine organisms including proteins, amino acids, lipid, fibers, pigments and
29 polysaccharides, all of which have the potential to modulate the immune system [1,2]. In
30 particular, seaweeds are recognized as a rich source of diverse bioactive compounds such as
31 sulfated polysaccharides (SP). Marine seaweeds contain many different polysaccharides
32 related to their taxonomic classifications [3]. It is acknowledged that SP from seaweed groups

33 include carrageenan and agaran from red seaweeds, ulvan from green seaweeds and fucoidan
34 from brown seaweeds [4,5]. SP present in the extracellular matrix of seaweeds [6] have been
35 increasingly reported to induce many health enhancing biological activities and are
36 considered to be antioxidant, anticancer, anti-inflammatory, antiviral, and
37 immunomodulatory [7]. The immunostimulatory effect induced by SP derived from marine
38 seaweeds has been widely studied in vertebrate and other invertebrate animal models [4,8]. In
39 other invertebrate animals, laminaran and β -glucan from the brown macroalgae *Laminaria*
40 *hyperborea* showed immunomodulatory effects on Atlantic salmon *Salmo salar* macrophages
41 [9]. Alginate extracted from seaweed increased the proportion of neutrophils, phagocytic
42 activity, respiratory burst and expression of interleukins in rainbow trout *Oncorhynchus*
43 *mykiss* post injection [10]. Administration of fucoidans from the brown macroalgae
44 *Sargassum wightii* by immersion in Pacific white shrimp *Litopenaeus vannamei* culture was
45 demonstrated to increase haemocyte proliferation and the mitotic index of haematopoietic
46 tissue [11]. Oral administration of carrageenan, a red seaweed polysaccharide from *Gigartina*
47 *sp.*, supplemented diets has been reported to increase the immune-related expression in *L.*
48 *vannamei* [12]. Additionally, treatment of alginate and fucoidans from the brown seaweed,
49 *Macrocystis pyrifera* and *Fucus vesiculosus* enhanced the immune response activity in head
50 kidney leukocytes of cod, *Gadus morhua* [13].

51 *Chondrus crispus*, a red seaweed from the family *Gigartinaceae*, is found abundantly
52 on the Atlantic coasts of Europe and North America [14]. It has been recognized that *C.*
53 *crispus* is a source of SP since the 1960s [15] and besides having a relatively high SP content,
54 *C. crispus* is also rich in proteins, amino acids, lipids and pigments [16]. The biochemical
55 composition of SP from seaweeds is dependent on the species, anatomical regions, growing
56 conditions, extraction procedures and analytical methods [4]. Protein content in seaweeds has
57 not been studied as well as polysaccharide content but it is acknowledged that seaweed
58 protein content can be species dependent with protein content in green and red seaweeds
59 being higher (up to 35-44% of dry mass) compared to brown seaweeds (less than 5%)
60 [16,17]. In addition to protein and polysaccharide content, polyphenols such as phenolic
61 compounds (2.5%) and flavonoids (0.1%), are also produced by seaweeds [18]. In a recent
62 study, water-soluble polysaccharides from *C. crispus* were observed to enhance the immune
63 response in the free-living nematode *Caenorhabditis elegans* and to suppress the expression
64 of quorum sensing and the virulence factors of the gram-negative bacteria *Pseudomonas*
65 *aeruginosa* [19]. In that study, it was suggested that the water-soluble polysaccharides
66 derived from *C. crispus* may play a health-promoting role in animals and humans. A growing

67 interest in the biomedical prospects of seaweed-derived SP in human health care has
68 emphasized the need for strategies to maximize SP extraction, bioavailability and investigate
69 precisely the therapeutic mechanisms of SP.

70 Mussels *Mytilus spp.* belonging to the Mytillidae family are important aquatic animals
71 that are harvested for human consumption worldwide [20] and also play an important
72 ecological role in the marine environment, forming biogenic reefs [21]. The blue mussel
73 *Mytilus edulis* are boreo-temperate in their distribution on both coasts of the Atlantic Ocean
74 in Europe and North America and are found in abundance, intertidally and subtidally, in both
75 sheltered and exposed sites, attached to hard substrates [22,23]. The Mediterranean mussel
76 *Mytilus galloprovincialis* is endemic to the Mediterranean, Black Sea and Adriatic Sea and
77 has expanded its range to the British Isles [24]. Evidence of hybridisation and hybrid zones of
78 *M. edulis* and *M. galloprovincialis* in the south west of England and Ireland were first
79 recorded in the 1970s and subsequent studies have further documented this phenomenon
80 [25,26,27]. The immune response of *Mytilus spp.* like other invertebrates includes a cellular
81 response and soluble haemolymph factors including lysozyme, an antibacterial enzyme, and
82 other antimicrobial peptides (AMPs) including defensins and mytimycins [28,29].
83 Haemocytes play an important role in invertebrate cellular immune response carrying out
84 phagocytosis, melanization, encapsulation and cell-to-cell communication and indirect
85 humoral immune response [30]. Several studies have shown that lysozymes are able to kill
86 Gram-negative bacteria, which has been demonstrated in bivalves [31,32].

87 The objectives of the present study were (a) to extract, quantify and chemically
88 analyze the SP derived from *C. crispus* and (b) assess the immune effects of exposure to the
89 extracted SP using mussels *Mytilus spp.* as the animal model for the first time. Findings from
90 this study will contribute to a better understanding of naturally derived biotherapeutics and
91 their contribution not only to animal health and well-being, in particular in species involved
92 in the aquaculture industry, but possibly to vertebrate health including humans.

93

94 **2. Materials and methods**

95

96 *2.1. Cold water extract of sulfated polysaccharides from Chondrus crispus*

97

98 Red seaweed Irish moss *C. crispus* (50 g) was collected in November 2015 from Inch
99 Strand, Midleton Co. Cork, Ireland (51.79508, -8.180008). Seaweed samples were freshly

100 harvested, returned to the laboratory and sand and epiphytes were removed. The *C. crispus*
101 was dried in an oven (35-40 °C) for 20-24 h. SP was extracted as previously described by
102 Wongprasert et al. (2014) [33]. Briefly, the dry seaweed was ground and de-pigmented with
103 benzene (24 h) and acetone (24 h). Five grams of de-pigmented seaweed powder was stirred
104 at 35-40 °C in 500 mL distilled water (DW) for 4 h. The extract was diluted with 500 mL of
105 hot water (100 °C) and centrifuged at $6000 \times g$ for 5 min. The seaweed residue was re-
106 extracted again by adding 800 mL of DW and stirring for 4 h and its supernatant was filtered
107 through a white cloth (35-48 mesh). The filtrate was allowed to cool to room temperature
108 (RT) and was subsequently kept at -20 °C overnight. The frozen supernatant was thawed and
109 centrifuged at $6000 \times g$ for 5 min to separate gel and non-gel fractions. The gel fraction was
110 discarded and the non-gel fraction SP was collected, freeze-dried and stored at -20 °C for
111 further study.

112

113 2.2. Chemical analysis

114

115 2.2.1. Sulfate content analysis

116

117 The sulfate content of SP was measured using a K_2SO_4 solution as a standard. Briefly,
118 20 mg of SP was hydrolyzed for 2 h at 100 °C in 0.5 mL of HCl (2N). The SP solution was
119 then transferred and made to volume in a 10 mL volumetric flask. Humic substance was
120 removed from the SP solution by centrifugation ($3000 \times g$, 10 min). The 2 mL of the
121 supernatant was diluted with 18 mL of DW and 2 mL of HCl (0.5 N). Then 1 mL of $BaCl_2$ -
122 gelatin reagent was added, swirled and retained for 30 min at RT. The mixture solution was
123 measured at 550 nm using a spectrophotometer and the percentage of sulfate was calculated
124 with reference to the standard curve given by a K_2SO_4 solution of known concentrations in
125 the ranges of 10-50 μg sulfate mL^{-1} .

126

127 2.2.2. Total polysaccharide analysis

128

129 Total polysaccharide content of SP was estimated by phenol-sulfuric acid method
130 using a galactose solution as a standard. One mL of SP solution (1 mg mL^{-1}) was mixed with
131 1 mL of 5% phenol in DW and 5 mL of sulfuric acid. The mixture was vortexed and allowed
132 to stand for 10 min at RT. The mixture solution was then cooled in an ice bath (15-20 °C) for
133 15 min and its absorbance was measured at 490 nm using a spectrophotometer. The

134 percentage of polysaccharide in SP was calculated with reference to the standard curve given
135 by a galactose solution of known concentrations in the ranges of 10-2000 $\mu\text{g mL}^{-1}$.

136

137 *2.2.3. Total protein analysis*

138

139 Total protein content of SP was measured by BCA protein assay [34]. An aliquot (25
140 μL) of SP or standard BSA solution (final concentrations at 0-2000 $\mu\text{g mL}^{-1}$) was added to
141 200 μL by BCA solution (BCATM protein assay kit, Thermo scientific, Inc., USA), mixed and
142 incubated at 37 °C for 30 min. After incubation, the solution was measured the absorbance at
143 562 nm using a spectrophotometer. The total protein content of SP was calculated with
144 reference to the standard curve given by a BSA solution of known concentrations.

145

146 *2.2.4. Total phenolic analysis*

147

148 The total phenolic content of SP was determined by using the Folin-Ciocalteu assay [35].
149 An aliquot (25 μL) of SP or standard solution of gallic acid (final concentrations at 0-100 mg
150 L^{-1}) was added to 250 μL of DW. A reagent blank was DW. Then, 25 μL of Folin-Ciocalteu's
151 phenol reagent was added to the mixture and shaken. After 5 min, 250 μL of 7 % Na_2CO_3
152 solution was added to the mixture. The solution was diluted to 625 μL with DW and mixed by
153 vortex. After incubation for 90 min at RT, the absorbance was measured at 750 nm using a
154 spectrophotometer. Total phenolic content of SP was expressed as mg gallic acid equivalents
155 (GAE) per 100 g of SP.

156

157 *2.2.5. Total flavonoid analysis*

158

159 Total flavonoid content of SP was measured by the aluminum chloride colorimetric assay
160 [36] and modified as described by Marinova et al. (2005) [37]. An aliquot (65 μL) of SP or
161 standard solution of catechin (final concentrations at 0-100 mg L^{-1}) was added to 325 μL of
162 DW and mixed. Then, 19.5 μL of 5% NaNO_2 was added. After 5 min, 19.5 μL of 10% AlCl_3
163 was added. At 6 min, 130 μL of 1 M NaOH was added and the total volume was made up to
164 650 μL with DW. The solution was mixed and shaken and the absorbance measured at
165 510 nm using a spectrophotometer. Total flavonoid content of SP was expressed as mg
166 catechin equivalents (CE) per 100 g of SP.

167

168 2.3. *Immune-stimulant activity of SP derived from C. crispus in the mussel Mytilus spp.*

169

170 2.3.1. *Experimental design*

171

172 Wild mussels *Mytilus spp.* (n=148) were randomly sampled from rocks at the
173 intertidal at Ringaskiddy, Cork Harbour, Co. Cork (51°50'N, 8°19'W), a known hybrid zone
174 for *M. edulis*, *M. galloprovincialis* and hybrids of both parent species, and a Class C water
175 quality site influenced greatly by anthropogenic effects such as agricultural and industrial
176 run-off, leachate from landfills and untreated waste water and sewage [27]. A total of 136
177 mussels were arbitrarily divided into 4 groups, each with duplicates of 17 mussels in stand-
178 alone rectangular plastic aquaria containing 8 L of artificial seawater (ASW) at a salinity of
179 34 and at 12 °C (8 tanks in total) and aerated using an air stone. The photoperiod was 12/12 h
180 of day/night cycles. The four groups consisted of Group 1, control mussels, not exposed to SP
181 and Groups 2, 3 and 4 consisting of mussels exposed to SP at final concentrations of 10, 20
182 and 50 µg mL⁻¹, respectively. Water changes were not carried out for the trial duration to
183 ensure that the SP dose amount added to the water at the beginning of the trial was not altered
184 and/or removed from each system. The trial was carried out in a constant temperature room
185 and with a constant photoperiod. Holding conditions were optimal for the mussels as no
186 mortalities were observed in either the control or treated mussels. Mussels were fed with 2
187 mL of Shellfish Diet 1800 (Reed Mariculture) consisting of a heteromorphic mix of
188 phytoplankton at a total concentration of 2 x 10⁹ cells mL⁻¹ containing *Isochrysis sp*, *Pavlova*
189 *sp*, *Thalassiosira weissflogii*, and *Tetraselmis sp* at day 4 of the experiment. At days 1, 2, 3,
190 4, 7 and 10 of the trial, 4 mussels were arbitrarily selected from each treatment (n =2/
191 replicate/ treatment) for immunological analyses including haemocyte viability, total
192 haemocyte counts, lysozyme activity assay and immune related mRNA expression. All
193 assays were performed in triplicate.

194

195 2.3.2. *Cell viability in haemolymph of mussels Mytilus spp.*

196

197 The neutral red retention time (NRRT) assay used to determine the haemocyte
198 viability in mussels as previously described [38] was carried out. Briefly, haemolymph (750
199 µL) was collected from individual mussels using a 20-gauge sterile needle (Microlance) fitted
200 on a 1 mL syringe containing 250 µL of tris-buffered saline solution (TBS). Haemolymph
201 samples were constantly vortexed to resuspend haemolymph and prevent aggregation. One-

202 hundred microliters of haemolymph sample was then transferred into microplate well, 20 μL
203 of 2% neutral red solution (Sigma Aldrich, USA) was added and then sample was held in a
204 dark humid chamber for 30 min. The supernatant was subsequently drained by tilting the
205 plates at an angle, facilitating drainage, followed by a gentle rinse with TBS, to remove any
206 excess neutral red dye. The neutral red was extracted by displacing the haemocytes using 100
207 μL of extraction solution (1% acetic acid in 50% ethanol) for 30 min. The plates were
208 measured at 450 nm and 570 nm using a UV max spectrophotometer (ELx808IU, Mason
209 Technology, USA).

210

211 2.3.3. Total haemocyte counts

212

213 Haemolymph sample (200 μL) was individually collected and fixed with 6% formalin
214 in Alsever's solution (200 μL) to prevent cell aggregation. After 10 min, 20 μL portions of
215 the fixed haemocyte suspension were mixed with 20 μL Rose Bengal solution (1.2% Rose
216 Bengal in 50% ethanol) and incubated at RT for 20 min before being used to determine the
217 total haemocyte counts using a hemocytometer under light microscopy at 20 x magnification
218 (Eclipse 80i microscope, Nikon Instruments Inc., USA). Haemocytometer (improved
219 Newbauer bright line) counts were made for 5/25 squares (volume of 1 square = $0.2 \times 0.2 \times$
220 0.1 mm^3) to calculate total haemocyte count mL^{-1} of hemolymph ($5 \times \text{count} \times 10^4 \times \text{dilution}$
221 factor) [39].

222

223 2.3.4. Lysozyme activity

224

225 Lysozyme activity was measured in serum as previously described by Prado-Alvarez
226 et al. (2015) [40]. Haemolymph sample (200 μL) was individually collected from mussels
227 and mixed with EDTA (200 μL) to prevent coagulation. Samples were centrifuged at $1200 \times$
228 g for 10 min to separate cells from the serum. Serum samples and serial dilution of standard
229 hen egg white lysozyme suspensions (Sigma Aldrich, USA) in 0.1 M phosphate buffer (final
230 concentrations at $0\text{--}5 \mu\text{g mL}^{-1}$) (30 μL) were dispensed in triplicate in 96 well plates before
231 adding a *Micrococcus luteus* (*M. lysodeikticus*) suspension (170 μL) at 0.2 mg mL^{-1} in 0.2 M
232 monobasic sodium phosphate and 0.2 M dibasic sodium phosphate (Sigma Aldrich, USA). A
233 set of five measurements of the optical density at 620 nm was recorded every minute using a
234 UV max spectrophotometer (ELx808IU, Mason Technology, USA).

235

2.3.5. Expression of immune-related mRNA in mussels

The haemocytes from mussels (4 mussels/ group) was RNA extracted in 200 μL TRI reagents according to the manufacturer's protocol (Sigma Aldrich, USA). The concentration and quality of RNA was determined by measuring the absorbance at 260/280 nm using a NanoDrop 2000 spectrophotometer (Thermo Scientific, USA). cDNA was transcribed from RNA (1 μg) using the Thermo Scientific RevertAid First Strand cDNA Synthesis Kit (Thermo Scientific, USA) containing RevertAid reverse transcriptase (200 U μL^{-1}), RiboLock RNase inhibitor (20 U μL^{-1}), Oligo (dT)18 primer (100 μM), dNTP mix (10 mM), and 5X reaction buffer (250 mM tris-HCl (pH 8.3), 250 mM KCl, 20 mM MgCl_2 , 50 mM DTT) at 42 $^\circ\text{C}$ for 1 h. The immune related mRNA was amplified by PCR using 1 μL (100 ng) of cDNA with specific primer sets and conditions (Table 1). 18S specific primer was also amplified as an internal control. Twenty-five microliters of PCR reactions contained Phusion DNA polymerase (2 U μL^{-1}), 10 mM dNTP mix, 10 μM of each forward and reverse primers and 5x Phusion HF buffer containing 7.5 mM MgCl_2 (Thermo Scientific, USA). The hot start PCR program used for immune related mRNA was performed with 98 $^\circ\text{C}$ for 30 s, followed by 40 cycles of 98 $^\circ\text{C}$ for 10 s, annealing temperature for each particular mRNA as shown in table 1, extension at 72 $^\circ\text{C}$ for 30 s followed by final extension at 72 $^\circ\text{C}$ for 5 min. The RT-PCR product was analyzed by 1.5% agarose gel electrophoresis, stained with 3% ethidium bromide, and visualized under ultraviolet light and documented using the EpiChemi3 darkroom (UVP, Inc., Upland, CA). Expression was semi-quantitatively determined from the ratio of band intensity to the internal control (18S) using ImageJ analysis program (from NIH website by Scion Corporation, Frederick, MD). Each assay was carried out in triplicate.

2.4. Statistical analysis

All assays were performed in triplicate. The data were presented as mean \pm SD and analyzed by one-way ANOVA followed by Tukey's multiple comparison and statistically significant difference was required at p-value less than 0.05.

3. Results

3.1. Extraction and chemical analyses of SP from *C. crispus*

270 SP extracted from the red seaweed *C. crispus* by cold water extraction was 15.36%
271 yield of dry weight. The chemical analysis showed that SP contained a sulfate content of 9.78
272 $\pm 0.42\%$ w/w, a polysaccharide content of $73.94 \pm 20.61\%$ w/w, a total protein content of
273 $7.08 \pm 0.45\%$ w/w, phenolic content of $2.55 \pm 0.24\%$ w/w and flavonoid content of $0.1002 \pm$
274 0.003% w/w (Table 2).

275

276 3.2. *C. crispus* SP extract effect on immune parameters in mussel *Mytilus* spp.

277

278 3.2.1. Effect of SP on haemocyte cell viability in *Mytilus* spp. using the NRRT assay

279

280 The results of haemocyte cell viability, analysed using the NRRT assay, are shown in
281 Fig. 1. Haemocyte cell viability of SP treated groups (final concentrations of 10, 20 and 50
282 $\mu\text{g mL}^{-1}$) were higher than the control group on Days 1, 2, 3, 4 and 10. While on day 7, the
283 haemocyte cell viability was not different.

284

285 3.2.2. Effect of SP on total haemocyte counts (THCs) in *Mytilus* spp.

286

287 The results of THCs from SP treated groups (final concentration of 10, 20 and 50 μg
288 mL^{-1}) and control group are shown in Fig. 2. THCs were higher in the SP exposed groups
289 than control mussels on Days 1, 2, 3, 4 and 10, with the exception of day 7. Moreover, the
290 results also show a positive correlation between THCs and SP dose concentration, with an
291 increase in THCs with increasing dose concentration. For example, on Day 3, mean THCs of
292 control and SP treated groups at the concentrations of 10, 20 and 50 $\mu\text{g mL}^{-1}$ were $0.993 \pm$
293 0.24×10^6 , $1.262 \pm 0.36 \times 10^6$, $1.915 \pm 0.15 \times 10^6$ and $1.724 \pm 0.31 \times 10^6$ cells mL^{-1} ,
294 respectively. On Day 10, THCs of control and SP treated groups at the concentrations of 10,
295 20 and 50 $\mu\text{g mL}^{-1}$ were $1.593 \pm 0.47 \times 10^6$, $1.599 \pm 0.90 \times 10^6$, $1.864 \pm 0.32 \times 10^6$ and 2.263
296 $\pm 0.67 \times 10^6$ cells mL^{-1} , respectively.

297

298 3.2.3. Effect of SP on lysozyme activity in *Mytilus* spp.

299

300 Lysozyme activity of mussels treated with SP from *C. crispus* was evaluated by
301 measuring the lysozyme activity in serum. Compared to control, lysozyme activity in the
302 serum of mussels treated with SP significantly increased on Days 1 and 2 (Fig. 3). On Day 3,
303 lysozyme activity in the serum of mussels treated with SP declined to control levels. On Days

304 4 and 7, lysozyme activity increased in treated and control groups from Day 3 but this
305 increase was not significant in both the control and treated mussels. On Day 10, lysozyme
306 activity in each groups had decreased from Days 4 and 7.

307

308 *3.3. The effects of SP stimulation on immune related genes expression in Mytilus spp.*

309

310 Post treatment with SP (final concentrations of 10, 20 and 50 $\mu\text{g mL}^{-1}$), the relative
311 expression of defensin, mytimycin and lysozyme with 18S mRNA was evaluated in treated
312 and control mussels on Days 1, 2, 3, 4, 7 and 10. The levels of defensin mRNA expression in
313 mussels from the SP treated groups was significantly higher than that of mussels in the
314 control group as shown in Fig. 4A and B. On Days 7 and 10, the levels of defensin mRNA
315 expression showed higher than that of the control group but it was not significant. For
316 mytimycin mRNA expression, on Days 1 and 2, the levels of mytimycin mRNA expression
317 in mussels in the SP treated groups were significantly higher than that of mussels in the
318 control group. On Day 3, the levels of mytimycin mRNA expression of SP treated at 50 μg
319 mL^{-1} remained significantly high from the control group. Whereas, on Days 4, 7 and 10, the
320 levels of mytimycin mRNA in mussels treated with the three SP concentrations showed no
321 difference to the mussels from the control group (Fig. 4A and C). For lysozyme mRNA
322 expression, on Days 1 and 2, the levels of lysozyme mRNA expression in mussels, from the
323 treated groups, was significantly higher than that of mussels in the control group. On Day 3,
324 the levels of lysozyme mRNA in mussels treated with the three SP concentrations showed no
325 difference to the mussels from the control group. An increase of lysozyme mRNA expression
326 in SP treated groups was observed again on Day 4. A decrease in lysozyme mRNA
327 expression was observed from Day 7 onwards and this was relative to SP dose concentration.
328 For example, on Day 7, the levels of lysozyme/18S mRNA expression in SP treated mussels
329 at concentrations of 10, 20 and 50 $\mu\text{g mL}^{-1}$ were 0.67, 1.1 and 2.0 folds of control,
330 respectively and on Day 10, it was 1.6, 3.1 and 3.1 folds of control, respectively (Fig. 4A and
331 D).

332

333 **4. Discussion and Conclusion**

334

335 In the present study, a high quality yield of SP from the red seaweed *C. crispus* was
336 extracted successfully by cold-water extraction following the methodology previously
337 reported for the red seaweed *G. fisheri* in Wongprasert et al. (2014) [33]. The yield of SP

338 from *C. crispus* (15.36%) was high and similar to other seaweeds, which have yielded a range
339 of 14.2-19.7%, using cold water extraction of the red seaweed *G. corticata* (2.8-19.7%) [41]
340 and the green seaweed *U. pertusa* (14.2%) [42]. Chemical analysis in this study indicated that
341 the SP in *C. crispus* contained high levels of carbohydrate including sulfate ester and low
342 levels of protein, which is comparable with other red seaweeds (*Porphyra*, *Palmaria*,
343 *Gracilaria*) [16,41]. It has been demonstrated that different thermal extractions obtain
344 different levels of polysaccharide and sulfate contents [41], with the cold-water extraction
345 method yielding higher levels of sulfate ester compared to the hot water extraction method, as
346 was also observed in this study.

347 The immune response in *Mytilus spp.* was observed in the present study to be
348 enhanced at a cellular, humoral and molecular level shortly after exposure to *C. crispus* SP
349 (on day 1). Under optimal conditions, bivalves such as *Mytilus spp.* will filter seawater at a
350 maximum rate [43] to facilitate feeding and respiration. Valve opening and continuous
351 filtering activity was observed in the mussels in this study thus readily exposing the treated
352 mussels to the SP dissolved in the tank seawater. Haemocyte cell viability and total
353 haemocyte counts were increased with SP treated mussel groups compared to the nontreated
354 mussel group. The immune system of bivalves lacks immune specificity and memory,
355 however, innate defense mechanisms that can identify and protect against non-self-material
356 have evolved, with invertebrate immune response centered largely on the multifunctional
357 haemocytes [44]. Early effects of physiological alterations are often seen as changes in
358 haemocyte counts with elevated cell counts a common response to environmental stress [45]
359 or as observed in this study exposure to an immunostimulant biocompound. The rapid SP
360 induced increase in haemocyte cell count and cell viability, observed in the treated mussels in
361 this study, highlights the effective delivery mechanism of SP to the mussels via their filter
362 feeding activity.

363 Similar findings including haemocyte proliferation and an increase in the mitotic
364 index of haematopoietic tissue in response to SP exposure have been observed in crustaceans
365 and fish species [10,11]. However, SP showed a differentially distinctive increase in
366 percentage of total haemocyte counts, for instance *Mytilus spp.* exposed to SP from *C.*
367 *crispus* and rainbow trout injected with alginate from *L. digitata* showed a similar increase in
368 number of haemocyte. While shrimp immersion with fucodains from *S. wightii* showed
369 increased percentage of total haemocyte counts in a lesser degree than that of mussel and fish
370 at the same concentration and exposure time.

371 In the present study, lysozyme activity was measured in the serum of SP-treated
372 compared with SP-nontreated mussels, and it was observed that lysozyme activity increased
373 shortly (on Days 1 and 2) after exposure to SP and the level of lysozyme mRNA expression
374 was also up-regulated on Days 1 and 2 of the trial. Previous studies reported that marine
375 microalgae, *Chaetoceros calcitrans* (*C. calcitrans*) and *Tetraselmis suecica* (*T. suecica*),
376 presented in Shellfish Diet 1800, have been reported to evoke an immune response in
377 bivalves [46]. It was reported that *C. calcitrans* had a positive effect on total haemocyte
378 count, granulocyte percentage, phagocytic rate and oxidative activity of clam haemocytes.
379 Moreover, *T. suecica* had a positive effect on the phagocytic rate of oyster haemocytes.
380 Interestingly, our study demonstrated that on Day 4 when the SP exposed mussels were fed
381 the heteromorphic commercial Shellfish Diet 1800 (containing microalgae), lysozyme
382 activity and lysozyme mRNA expression increased more once again than the nontreated
383 control mussels, suggesting the SP possibly enhanced increased lysozyme activity after
384 Shellfish Diet 1800 feeding.

385 Antimicrobial peptides such as defensins, mytilins, myticins and mytimycin, have
386 been identified in mussel species (*M. galloprovincialis* and *Bathymodiolus azoricus*) and are
387 associated with a humoral immune response [28]. Consistent with lysozyme mRNA, defensin
388 and mytimycin mRNA expressions were up-regulated shortly (on Days 1, 2 and 3) after SP
389 treatment in this study. It was reported that polysaccharides, sulfated galactans from the red
390 seaweed *Gracilaria fisheri* stimulated immune AMPs-mRNA expressions in *L. vannamei*
391 shrimp haemocyte [47], however the response in *Mytilus spp.* is much more rapid following
392 exposure to SP.

393 In conclusion, the present study demonstrated that SP was efficiently extracted from
394 *C. crispus* using the cold-water extraction methodology and this extracted SP has a rapid
395 immune-stimulant effect in mussel *Mytilus spp.* This prompt immune response on a cellular,
396 humoral and molecular level, observed in *Mytilis spp.* and instigated by SP exposure, may be
397 beneficial to animals in assisting them in overcoming the challenges of parasites and disease.
398 More importantly the short-term energy output required for such a rapid immune response
399 may not be too costly to individuals whose health maybe already be compromised due to
400 stress and poor health.

401

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407 the study.

408

409 **Legends**

410

411 **Table 1** Specific primers and conditions used for the determination of immune-related
412 mRNA expression in mussels *Mytilus spp.*

413

414 **Table 2** Contents of sulfated polysaccharides (SP) from the red seaweed *C. crispus*. SP
415 showed sulfate, polysaccharide, protein, phenolic and flavonoid contents. Data are presented
416 as a mean of triplicate independent experiments.

417

418 **Fig. 1.** Haemocyte cell viability of mussels *Mytilus spp.* exposed with SP from the red
419 seaweed *C. crispus* at the concentrations of 10, 20 and 50 $\mu\text{g mL}^{-1}$ determined by neutral red
420 retention time (NRPT) assay. The bar graphs show the haemocyte cell viability presented as
421 the percent of control (mean \pm standard deviation of triplicate independent experiments). *
422 indicates value significantly different from the control ($P < 0.05$), $n = 4$.

423

424 **Fig. 2.** Total haemocyte counts of mussels *Mytilus spp.* exposed with SP from the red
425 seaweed *C. crispus* at the concentrations of 10, 20 and 50 $\mu\text{g mL}^{-1}$. Data is presented as a
426 mean \pm standard deviation of triplicate independent experiments. * indicates value
427 significantly different from the control ($P < 0.05$), $n = 4$.

428

429

430 **Fig. 3.** Lysozyme activity of mussels *Mytilus spp.* exposed with SP from the red seaweed *C.*
431 *crispus* at the concentrations of 10, 20 and 50 $\mu\text{g mL}^{-1}$. Data is presented as a mean \pm
432 standard deviation of triplicate independent experiments. * indicates value significantly
433 different from the control ($P < 0.05$), $n = 4$.

434

435 **Fig. 4.** Expression levels of immune related mRNA of mussels *Mytilus spp.* in the initial
436 samples, control and SP from the red seaweed *C. crispus* at the concentrations of 10, 20 and
437 50 $\mu\text{g mL}^{-1}$. (A) RT-PCR analysis of defensin, mytimycin and lysozyme mRNA expression
438 by 1.5% agarose gels. (B) Densitometry value of defensin mRNA relative to 18S in different

439 groups, (C) densitometry value of mytimycin mRNA relative to 18S in different groups and
440 (D) densitometry value of lysozyme mRNA relative to 18S in different groups. Bars indicate
441 mean \pm standard deviation. * indicates value significantly different from control group
442 ($P < 0.05$), $n = 4$.

443

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Primers	Primer sequences (5' to 3')	References	Annealing temp
Lysozyme-F Lysozyme-R	ATGTGGAATCTGAAGGACTTGT CCAGTATCCAATGGTGTTAGGG	[48]	60 °C for 30 s
Defensin-F Defensin-R	GTGGCGTCTGCTGGGTTT GAATGGACTTACAATGTCGATGACA	[49]	58 °C for 30 s
Mytimycin-F Mytimycin-R	CAATCCATCACTGTTGAAT ATGGTAAATCGTGTTATGAACGTG	[50]	58 °C for 30 s
18S-F 18S-R	TTACGTCGGCGCAACTTCT CTGTTCCAAGGACTTTAATG	[50]	57.6 °C for 30 s

Table 1

Contents	% of SP extract (w/w)
Yield	15.36
Sulfate	9.78 ± 0.424
Polysaccharide	73.94 ± 20.61
Protein	7.08 ± 0.455
Phenolic	2.55 ± 0.236
Flavonoid	0.10 ± 0.003

Table 2

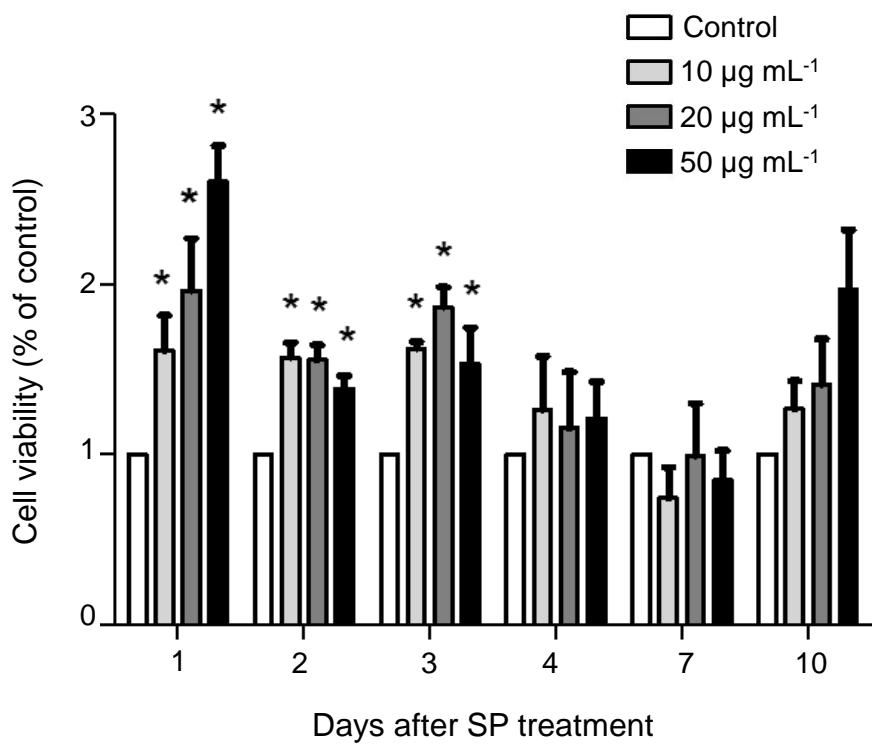


Figure 1

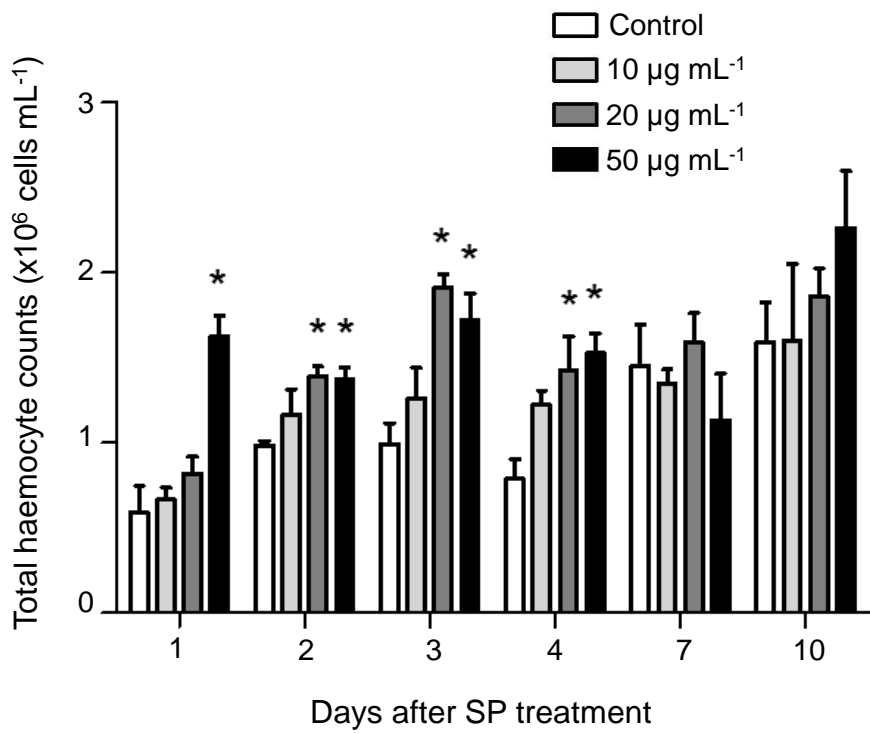


Figure 2

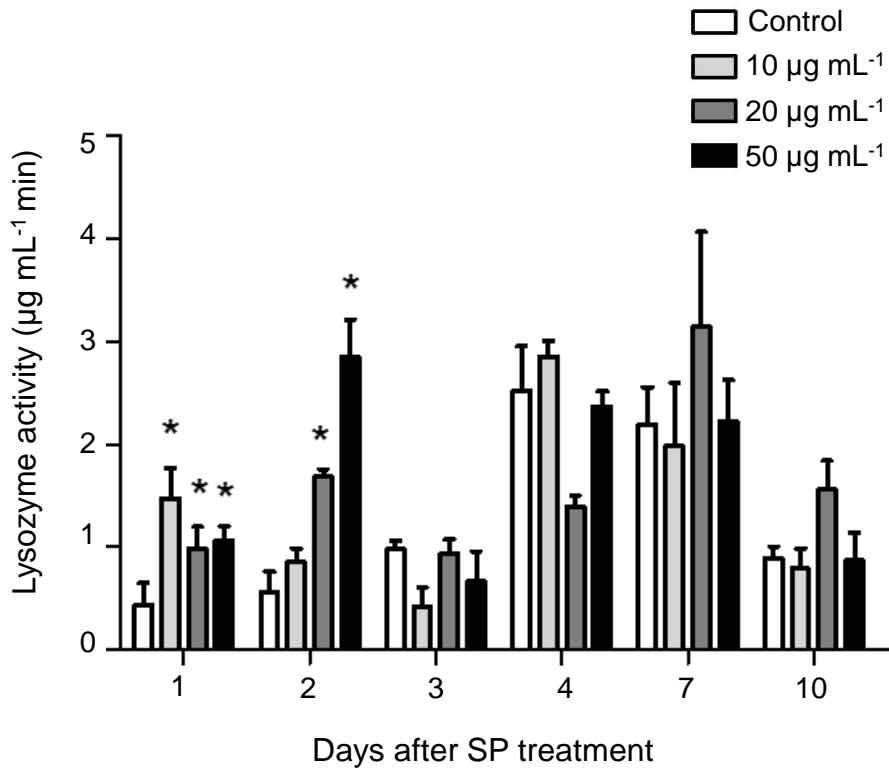


Figure 3

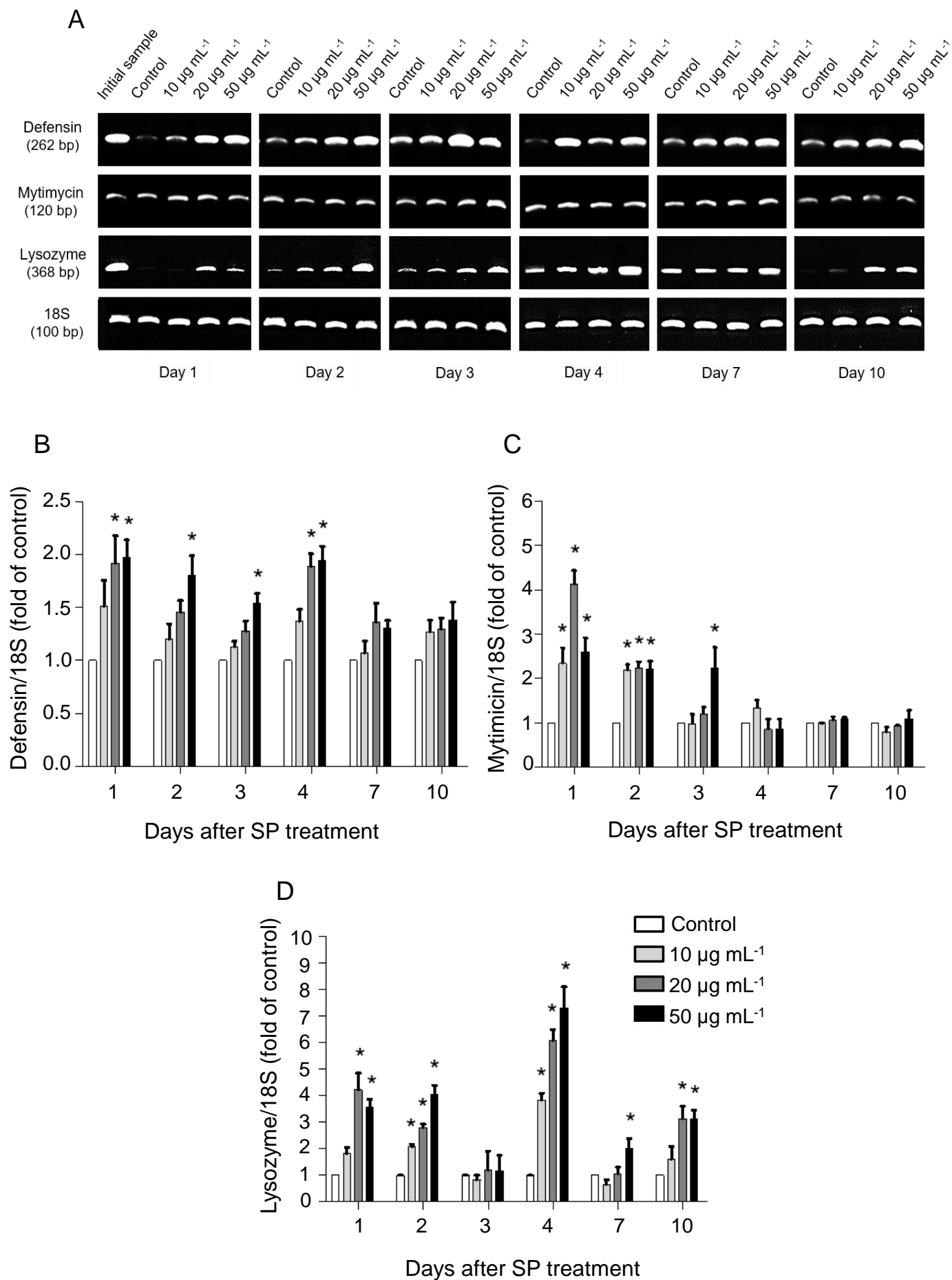


Figure 4

Highlights

- Sulfated polysaccharides (SP) were extracted from Irish moss *Chondrus crispus*
- SP enhanced immune parameters in mussels *Mytilus spp.*
- SP up-regulated expression of immune genes in mussels shortly after exposure