1 Ibuprofen As A Water Pollutant On The Defensive Behaviour And

2 Microbiome Of Grass Shrimp *Palaemonetes* Spp.

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4

5 ABSTRACT

6 Over-the-counter use of painkillers dates to the 1900s, and has become 7 increasingly normalized in modern society. The first-pass effect coupled with 8 insufficient sewage treatment mechanisms have led to significant levels of 9 pharmaceutical pollution in the environment. Scientists are only recently delving into 10 the implications of chronic use on the human gut microbiome, and similar 11 ramifications may hold for marine life that ingest pharmaceutical waste. As one of the 12 first research efforts to understand the effects of non-steroidal anti-inflammatory drugs (NSAIDs) on the gut microbiome of common marine species, this study points 13 14 to potential microbiome alterations that may be correlated to behavioural complications from ibuprofen pollution. 15

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17 INTRODUCTION

With increasing environmental conservation efforts, the body of research 18 surrounding the impact of personal care and pharmaceutical products (PCPPs) 19 20 pollution in natural waters on humans and other organisms has grown significantly in 21 recent years. 19% of U.S adults are estimated to be chronic users of NSAIDs(1), of 22 which ibuprofen is one. While this behaviour has long been linked to stomach ulcers, 23 intestinal bleeding and perforation(2), recent research has shown effects on human 24 gut microbiome profiles, specifically an increased abundance of 25 Pseudomonadaceae, Puniceicoccaceae families(3). Altered microbial expression 26 has yet to be definitively linked, however, to gastrointestinal effects in humans. 27 Seawater harbours up to 3 µg/L of ibuprofen(4), making it one of top eight PCPP 28 contaminants (5). This study aims to understand the effect of different ibuprofen

29 concentrations on *Palaemonetes* spp., a common grass shrimp.

30 Current literature on the effects of PCPPs on marine life focuses on direct consequences for the organism itself, such as decreased motor skills(6), metabolic 31 32 disorders(7), endocrine disruption, and oxidative stress(8), with limited investigation of effects to the host's microbiome. Thus, my hypothesis is two-fold: firstly, the 33 34 shrimp will exhibit lethargic and delayed defensive mechanisms with increasing 35 ibuprofen concentration. Secondly, taxonomic abundance of *Pseudomonadaceae*, Puniceicoccaceae families in the shrimps' microbiome will increase with increasing 36 37 ibuprofen concentrations.

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39 METHODOLOGY

40 Sample Collection

Twenty-four shrimp were collected off the dock at the Marine Resources Center at the Marine Biological Laboratory (MBL), in Woods Hole, Massachusetts. Six shrimp each were put into four three-liter ambient seawater tanks dosed with ibuprofen at final concentrations of 0 (Control-IB), 3 (Low-IB), 21 (Medium-IB), and 80 (High-IB) mg/L respectively. Ibuprofen was added using powdered 200 mg dosage Advil tablets.

To sample the gut microbiome, I extracted the digestive tract through dissection at the time of death or after five days, whichever occurred earlier. Three samples were collected per treatment, and stored at -80°C.

50 Behavioural Analysis

In response to perceived threats, shrimp flick their tails and jump a horizontal distance (9). Throughout the experiment, the shrimp were tested regularly on their defence mechanisms by inserting a glass rod into the water and moving it closer to individual shrimp. Two metrics were recorded: perceived threat – the distance between the rod and the shrimp right before it jumps; the defence distance – defined as the distance jumped following the disturbance.

57 DNA isolation, 16S rRNA amplification, Illumina MiSeq sequencing

58 DNA was extracted following manufacturer protocol for the QIAGEN DNeasy 59 PowerSoil kit (10), except the entire shrimp gut was used in place of 0.25 g of

- 60 sample to avoid contamination. The V4 region of 16S rRNA gene was amplified with
- 61 primers 515F and 816R, following 16S Illumina Amplicon protocol in the Earth
- 62 Microbiome Project (11). DNA concentration was confirmed with Nanodrop (12),
- 63 PicoGreen assay (13) and agarose gel electrophoresis (14). PCR products were
- 64 sequenced on the Illumina MiSeq platform in the Bay Paul Center at the MBL (15).

65 Sequencing Data and Statistical Analysis

66 Sequence data was demultiplexed using the "demux emp-paired" command (16). 67 One Low-IB sample was lost due to low yield. The DADA2 plugin in QIIME2-2018.2 (17) was used to group related sequences into exact sequence variants (ESVs). All 68 69 sequences with Phred scores below 25 were excluded, and taxonomy was assigned 70 using the Greengenes database v13.8 (18). Sequences unidentifiable beyond 71 kingdom level were discarded. Rarefaction curves using the Shannon H index (19) 72 were plotted to ensure the filtered data accurately represented samples' microbial 73 diversity.

I used UniFrac distances (20) to examine inter-sample taxonomic differences,
alongside principal coordinate analysis (PCoA) (21) and permutational multivariate
analysis of variance (PERMANOVA) (22). Power calculations were conducted using
HMP R package (23) and G*Power ANOVA (24).

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79 **RESULTS**

80 Sequencing Analysis

81 Alpha and Beta Diversity

Shannon index analysis across the concentration gradient yielded no significant
differences between any treatment groups, with an average index of 6.68 at
insignificant p-value 0.082. Between dead and live samples however, there was a
significant H-index of 7.5 and p-value 0.006.

The weighted Unifrac metric (Figure 1) showed more meaningful clustering compared to the unweighted counterpart. Statistically significant clustering was not observed between treatments, but was observed between live and dead samples. Thus, both dead and live shrimp microbiome profiles show intra-group similarity in 90 phylogenetic distance and relative taxonomic abundance, but live profiles are more





Fig. 1: A) Plot of principal coordinate analysis (PCoA) from weighted Unifrac showing insignificant clustering of the gut microbiome profile difference between treatments, with pairwise PERMANOVA p-values above 0.05 threshold. B) Significant gut microbiome profile difference between live and dead samples observed, with significant p-value of 0.001.

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93 Differential taxonomic abundance

Fig.2: Taxonomic composition of samples by treatment. Asterisks denote samples that were dead upon collection. Only top 13 most abundant taxa are shown.

- Dead shrimp gut microbiota (Figure 2) has lower average taxonomic diversity, and is
 overwhelmingly populated by *Vibrionaceae* (96.2%). Average relative abundances
 are denoted in parentheses.
- 98 The gut microbiome of live shrimp is more diverse, with significant microbial
- 99 abundances of *Moraxellaceae* (30.1%), *Pseudomonadaceae* (16.1%),
- 100 Weeksellaceae (11.72%) and Comamonadaceae (10.5%). Vibrionaceae (3.02%)
- 101 was the sixth most abundant taxa, and notably contain a different, though
- 102 unidentified, strain compared to dead counterparts.
- ANCOM analysis (25) of the live shrimp (data not shown) did not identify any
 ESVs that differed significantly between treatments.

105 Behavioural Analysis

All shrimp in High-IB were dead by day 2, and 8 molts were deposited. On day 3,

5 shrimp in Medium-IB had died, and 4 molts were deposited. 1 molt in Low-IB and 0

108 molts in Control-IB were deposited by day 6.



Fig. 3: A) Perceived threat distances measured for Control-IB and Low-IB treatments on day 0, 1, 3, 6 of the experiment. Each record corresponds to one shrimp. Low-IB showed statistically significant decreasing relationship with time at 0.05 p-value threshold. **B)** Defence distances measured under the same conditions as (A).

- 110 Logistic regression of behavioral data (Figure 3) showed visually negative
- 111 correlations between ibuprofen concentration and defence capabilities, though only
- the trend for perceived threat was statistically significant.
- 113

114 **DISCUSSION**

115 Along the concentration gradient for live samples, I was unable to identify statistically significant differences in taxonomic abundances and diversity. However, 116 117 the microbiome profiles of live and dead samples did differ significantly, especially in the abundance of Vibrionaceae. While a form of predominant marine bacteria, 118 119 research has also shown that many strains are naturally virulent, and marine hosts 120 have evolved mechanisms to restrict pathogenesis (26). It is therefore possible that 121 higher fatality relative to concentration is linked to Vibrionaceae spp. by 122 compromising host immunity, aiding vibrio pathogenesis – by either colonization of external strains or once commensal strains. It is also possible that the predominant 123 124 strain observed in the dead samples is commonly found in decaying grass shrimp gut microbiome. 125

126 However, significant physical effects were observed. High mortality rates in High-127 IB and Medium-IB show that ibuprofen is lethal for grass shrimp at high 128 concentrations. Ibuprofen ingestion can compromise host survival in the face of 129 predators also, shown by significantly shorter perceived threat distances and visibly 130 lowered defence distances measured by Low-IB compared to Control-IB. Lastly, the 131 alarming count of shrimp molts in High-IB and Medium-IB suggests that ibuprofen 132 can indirectly alter the circadian rhythm of the molting cycle. Given that the regular 133 molting cycle ranges from fourteen to twenty-one days (27), the newly developed 134 shells may be lacking in protective capabilities and may compromise survival. Further research is needed to determine if weakened external shells are correlated 135 136 to or accelerate shrimp death.

I was underpowered to conclude significant differences in alpha and beta
diversity. Power analysis using G*Power ANOVA on the Shannon H index across the
concentration gradient for live samples only yielded 0.11 power, and the minimum
sample size for 0.8 power given my effect size is 36. Using Dirichlet-Multinomial

parameter test on the beta diversity across the concentration gradient yielded 0.50power. Thus, these results are best taken as a starting point for further research.

143 Aside from sample size, a few alterations should be made to the experimental 144 design in further research. Firstly, the duration of the experiment should be increased. Secondly, the experiment should be conducted with seawater diluted by 145 146 pure ibuprofen salt concentrations as well as powdered Advil solutions. While 147 ibuprofen is the active ingredient in Advil tablets, I cannot exclude the possibility that some trace ingredient in the tablets, undigested by the human body, is the true 148 culprit for host compromise and fatality in this experiment. Should this be true, this 149 could have implications for pharmaceutical companies as they revise peripheral 150 ingredients in pills and for waste management systems filtering for pharmaceutical 151 152 contaminants.

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