



Communication

# Response of Coral Reef Dinoflagellates to Nanoplastics under Experimental Conditions Suggests Downregulation of Cellular Metabolism

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Received: 25 September 2020; Accepted: 5 November 2020; Published: 9 November 2020



**Abstract:** Plastic products contribute heavily to anthropogenic pollution of the oceans. Small plastic particles in the microscale and nanoscale ranges have been found in all marine ecosystems, but little is known about their effects upon marine organisms. In this study, we examine changes in cell growth, aggregation, and gene expression of two symbiotic dinoflagellates of the family Symbiodiniaceae, *Symbiodinium tridacnidorum* (clade A3), and *Cladocopium* sp. (clade C) under exposure to 42-nm polystyrene beads. In laboratory experiments, the cell number and aggregation were reduced after 10 days of nanoplastic exposure at 0.01, 0.1, and 10 mg/L concentrations, but no clear correlation with plastic concentration was observed. Genes involved in dynein motor function were upregulated when compared to control conditions, while genes related to photosynthesis, mitosis, and intracellular degradation were downregulated. Overall, nanoplastic exposure led to more genes being downregulated than upregulated and the number of genes with altered expression was larger in *Cladocopium* sp. than in *S. tridacnidorum*, suggesting different sensitivity to nano-plastics between species. Our data show that nano-plastic inhibits growth and alters aggregation properties of microalgae, which may negatively affect the uptake of these indispensable symbionts by coral reef organisms.

**Keywords:** nanoplastics; dinoflagellate; coral reef; *Symbiodinium*; *Cladocopium*; gene expression

## 1. Introduction

Coral reefs provide a habitat for marine invertebrate and vertebrate species alike, sustaining the highest biodiversity among marine ecosystems [1]. Formed primarily by scleractinian corals and coralline algae, coral reefs are complex and vulnerable ecosystems. Structural complexity of coral reefs, and, by extension, the capability to sustain biodiversity often declines due to natural and human-related stressors [2,3].

One important stressor for coral reef ecosystems is plastic pollution. Small plastic particles (>1 mm) have been reported from coral islands at more than 1000 items/m<sup>2</sup> [4]. Further fragmentation of these particles leads to nano-plastics (<1 μm) [5]. Microplastic particles induce stress responses in scleractinian corals, and suppress their immune systems and capacity to cope with environmental toxins [6]. When ingested by corals [7–9], microplastics disrupt the anthozoan-algal symbiotic relationship [10]. They are also linked to potential adverse effects on calcification [11] with exposure resulting in attachment of microplastic particles to tentacles or mesenterial filaments, ingestion of microplastic particles, and increased mucus production [12]. Su et al. [13] exposed the coral

symbiont, *Cladocopium goreauii*, to 1- $\mu\text{m}$  polystyrene spheres, leading to diminished detoxification activity, nutrient uptake, and photosynthesis as well as increased oxidative stress, apoptosis levels, and ion transport. Plastic particles seem to negatively impact symbiotic relationships between corals and their microalgae, thereby degrading the entire coral reef ecosystem. However, this has not been systematically investigated.

Nano-plastics can originate by fragmentation of larger plastic objects through photochemical and mechanical degradation. There are also primary sources of nano-plastics. Medical and cosmetic products, nanofibers from clothes and carpets, 3D printing, and Styrofoam byproducts find their way into coral reef ecosystems via river drainages, sewage outfalls, and runoff after heavy rainfall, as well as via atmospheric input and ocean currents. Nano-plastics have recently been reported in ocean surface water samples [14]. Since the nanoplastic research is still in its infancy, many unanswered questions remain, starting with the environmental concentrations in various ecosystems [15,16]. Since detecting nano-plastics' concentrations directly is still not possible [17], a better understanding of the potential impacts is necessary to encompass a range of different concentrations. The miniature size of these particles leads to higher surface area to volume ratios and enhanced reactivity of smaller particles coupled with the ability to pass across biological barriers and enter cells [18] when compared to micro-plastics.

In this study, we focused on the microalgal symbionts of mollusks that inhabit fringing coral reefs of Okinawa. Knowledge of the effects of nano-plastics on the symbionts of Tridacninae (giant clams) and Fraginae (heart cockles) will benefit conservation and restocking efforts, since both are obligatory photo-symbionts and important contributors to coral reef ecosystems. Approximately 30 Symbiodiniaceae phylotypes are economically important for fisheries [19]. This study specifically investigated effects of nano-plastics (42-nm polystyrene spheres) on the growth rates, aggregations, and gene expression changes in *Symbiodinium tridacnidorum* (symbionts of the Tridacninae) and *Cladocopium* sp. (symbionts of the Fraginae).

## 2. Materials and Methods

### 2.1. Exposure to Nano-Plastics Using Roller Tanks

The majority of host animals obtain their indispensable symbiotic dinoflagellates from coral reef sand and the water column [20,21]. Roller tanks and tables were used to simulate the natural environment of the dinoflagellate vegetative cells in their free-living state [22,23]. Roller tanks have commonly been used to promote aggregation since Shanks and Edmondson [23,24]. Fifteen roller tanks of 13.4 cm in diameter and 7.5 cm in height with a capacity of 1057 mL were employed. In tanks, aggregation can occur [23], ensuring that microalgae are exposed to the polystyrene nano-plastics (nanoPS) in a way that mimics their natural habitat. Once rotation commenced, continuous aggregate formation and suspension were ensured [24] as well as continuous exposure to nanoPS. Roller tanks are closed for the entire duration of the experiment, so that exposure levels of the nanoPS remain constant throughout. Tanks were closed without bubbles so as not to disturb the aggregation process with turbulence. To compare differences between species, two dinoflagellates, *Symbiodinium tridacnidorum* (clade A3 strain, ID: NIES-4076) and *Cladocopium* sp. (clade C strain, ID: NIES-4077) were cultured in artificial seawater containing 0.2 $\times$  Guillard's (F/2) marine-water enrichment solution (Sigma-Aldrich) in roller tanks [25,26]. *S. tridacnidorum* and *Cladocopium* sp. were isolated from *Tridacna crocea* and *Fragum* sp. in Okinawa, Japan [25]. Using glass flasks, precultures for the stress experiment were established, as previously described [26].

Microplastics (>1 mm) from a coral reef and the ingestion (53 to 500  $\mu\text{m}$ ) by coral reef clams have been reported and microplastic removal by giant clams has been proposed [4,27]. To simulate nano-plastic accumulation in coral reefs and in the host organisms, three different concentrations (0.01 mg/L, 0.1 mg/L, and 10 mg/L) of nano-plastic (42-nm pristine polystyrene beads, nanoPS<sub>42</sub>, from Bangs Laboratories Inc., catalog number FSDG001, polystyrene density 1.05 g/cm<sup>3</sup>, nanoPS)

were added to the treatment tanks (Table S1). Preliminary tests were run to confirm no leaching of the fluorescent dye (data not shown). Concentrations were chosen to span a range of possible environmental concentrations, starting at 0.01 mg/L with a surface area of  $1.36 \times 10^6 \mu\text{m}^2/\text{L}$  and  $2.46 \times 10^8$  particles per L. The next highest concentration is just one magnitude higher (0.1 mg/L, surface area  $1.36 \times 10^7 \mu\text{m}^2/\text{L}$  and  $2.46 \times 10^9$  particles per L). This middle range concentration corresponds to actually observed lower concentrations of microplastic particles [28]. Just as microplastic concentrations are highly variable, nanoplastic concentrations are assumed to change depending on the proximity to human activity. To account for these variables, but not at the highest measured microplastic concentration, we placed our highest concentration at 10 mg/L with a surface area of  $1.36 \times 10^9 \mu\text{m}^2/\text{L}$  and  $2.46 \times 10^{11}$  particles per L (Table S1). Treatment tanks as well as control tanks (no nanoPS) were established in triplicate. Three tanks without algae were prepared as negative controls (at 10 mg/L, 0.01 mg/L, and 0 mg/L nano-plastic). In each culture tank, the final cell density of the two strains was adjusted to  $\sim 7 \times 10^5$  cells/mL. Tanks were harvested after 9–11 days for logistical reasons, making replicates a day apart (Table S2).

## 2.2. Measurements of Cell Density and Aggregation

Cells for growth rates were counted using hemocytometers (C-Chip DHC-N01) under a Zeiss Axio Imager Z1 microscope (Jena, Germany). At least two subsamples and 200 cells were counted per sample.

Aggregates were imaged and counted in each tank and for five size classes, as follows: 0.2–0.5 mm, 0.5–1 mm, 1–2.5 mm, 2.5–3.5 mm, and >3.5 mm in the longest dimension. Tanks of the same concentration were sampled at the same time of day. Controls were sampled first and then in order of increasing nanoPS<sub>42</sub> concentration to avoid nano-plastic carry over from higher concentrations to lower. In order to examine how nanoPS<sub>42</sub> affects aggregate formation, aggregates were collected for different measurements after the approximate total number of aggregates in each tank had been determined. Aggregation of algae and plastic was confirmed with 3D imaging using a Zeiss Lightsheet Z.1 and Imaris software. NanoPS<sub>42</sub> was observed with a band-pass filter (excitation: 405 nm, emission: 505–545 nm) and chloroplasts were visualized using a long-pass red filter (excitation: 488 nm, emission: 660 nm).

One fourth of all aggregates were collected for RNA analysis (2 min spin down at 12,000 rpm and discarding the supernatant, freezing in liquid nitrogen, and storage at  $-80^\circ\text{C}$ ). For all other measured factors, harvest included separate sampling of the aggregate fraction (aggregates >0.5 mm, Agg) and the surrounding sea water fraction (aggregates <0.5 mm and un-aggregated cells) [29]. Aggregates for sinking velocity (three aggregates per size class for 11.5 cm in a 100-mL glass graduated glassware cylinder) was collected in artificial seawater at the same temperature as experiments were conducted.

## 2.3. RNA Extraction, Library Construction, and Sequencing

Frozen cells were broken mechanically using a polytron (KINEMATICA Inc., Luzern, Switzerland) in tubes chilled with liquid nitrogen. RNAs were extracted using Trizol reagent (Invitrogen) according to the manufacturer's protocol. The quantity and quality of total RNA were checked using a Qubit fluorometer (ThermoFisher, Waltham, MA) and a TapeStation (Agilent, Santa Clara, CA), respectively. Libraries for RNA-seq were constructed using the NEBNext Ultra II Directional RNA Library Prep Kit for Illumina (#E7760, NEB). Sequencing was performed on a NovaSeq6000 SP platform. Nine mRNA-seq libraries from nanoPS-exposed photosymbiotic algae were sequenced (3 concentrations  $\times$  3 exposure times) plus three controls (Table S2).

## 2.4. RNA-Seq Data Mapping and Clustering Analysis

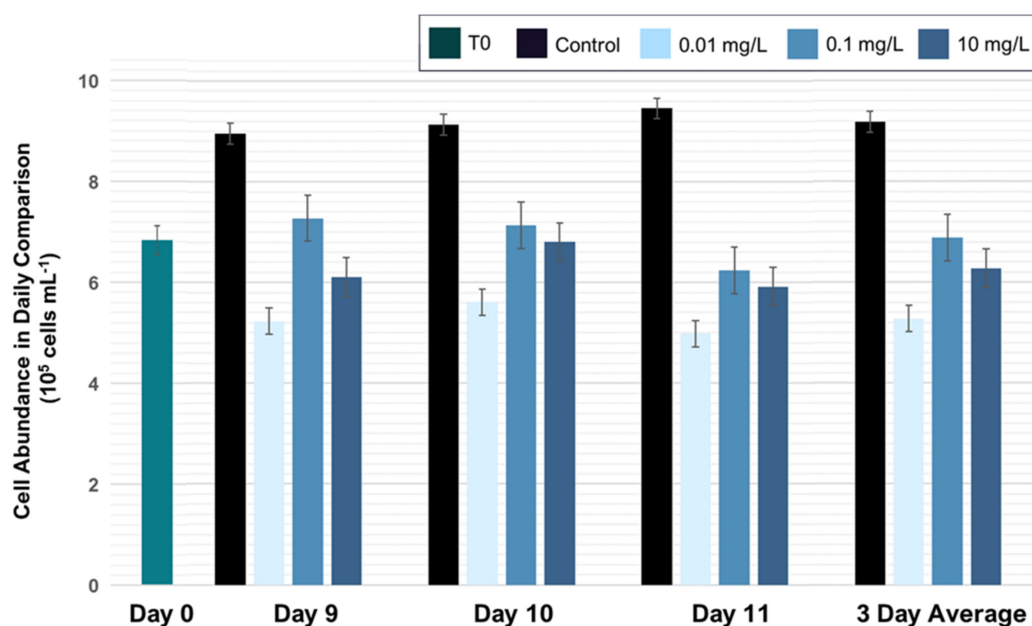
Raw sequencing data obtained from the NovaSeq6000 were quality trimmed with Trimmomatic (v. 0.32) in order to remove adapter sequences and low-quality reads. Paired reads that survived the trimming step (on average 92%) were mapped against reference transcripts of *Symbiodinium*

and *Cladocopium* sp. For each gene in the genomes of *Symbiodinium* and *Cladocopium* sp. a \*.t1 transcript form was used as a reference sequence. Mapping was performed using RSEM (RNA-Seq by Expectation-Maximization) [30] with the bowtie (v. 1.1.2) as an alignment tool. Expression values across all samples were normalized by the TMM (Trimmed Mean of *M*-values) method [31]. Genes with differential expression (two-fold difference and  $p < 0.001$ ) were identified with edgeR Bioconductor, based on the matrix of TMM normalized TPM Transcripts Per Kilobase) values. Experimental samples were clustered according to their gene expression characteristics using edgeR. Annotations were performed using BLAST2GO and Pfam databases [25] and are available at the genome browser site (<https://marinegenomics.oist.jp>).

### 3. Results and Discussion

#### 3.1. Suppression of Algal Growth by Nano-Plastic Exposure

Exposure to nanoPS<sub>42</sub> decreased the mean growth rate of photosymbiotic algae (Figures 1 and S1). The greatest reduction in growth rate was seen at the lowest nanoPS<sub>42</sub> treatment (0.01 mg/L) with cell densities reduced from starting values by  $-0.062 \pm 0.02$  (Holm-Sidak,  $p = 0.002$ ), which was followed by the highest nanoPS<sub>42</sub> treatment (10 mg/L) with  $-0.013 \pm 0.05$  (Holm-Sidak,  $p = 0.026$ ). In the 0.1 mg/L treatment, cell densities increased slightly by  $0.028 \pm 0.04$ . Thus, nanoPS<sub>42</sub> either inhibited algal growth in a non-linear manner or had a limited effect [32]. Reductions in growth rates have also been reported in the micro-plastic study of Reference [13] in *Cladocopium goreau* and in other microalgae exposed to micro-plastics (*Chlamydomonas reinhardtii* [33] and *Skeletonema costatum* [34]).



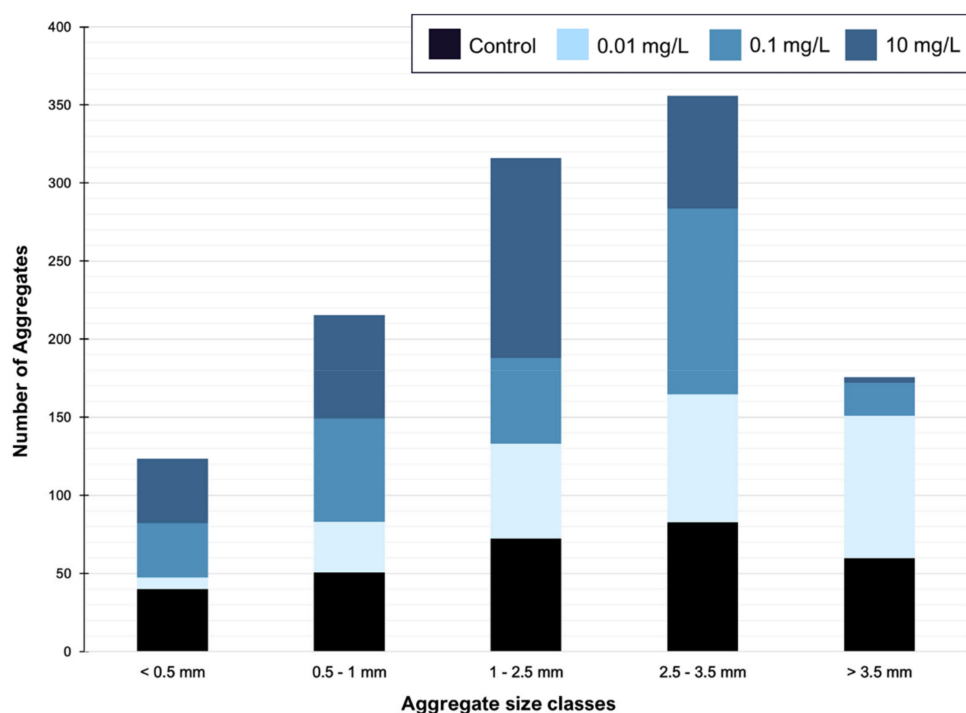
**Figure 1.** Treatment and control tanks were sampled after 9, 10, and 11 days. Experiments started with  $\sim 680,000$  cells/mL in all tanks. There are differences between the growth rate in the different treatments, but the ratio stays the same over all three sampling days. The cell density in the control was  $9.83 \pm 0.39 \times 10^5$  cells per mL, while treatment tanks were significantly lower: 0.01 mg/mL:  $5.69 \pm 0.12 \times 10^5$  cells per mL, 0.1 mg/mL:  $7.51 \pm 0.34 \times 10^5$  cells per mL, and 10 mg/mL:  $6.96 \pm 0.40 \times 10^5$  cells per mL. Bars display a confidence interval.

In addition, Su et al. [13] reported a reduction in cell size in *Cladocopium goreau*. Further investigations are needed to see if this is the case under nano-plastic exposure. The biggest growth rate reduction observed was at 0.01 mg/L nanoPS<sub>42</sub>, which is far below the 5 mg/L used by Su et al. [13]. The nutrient deficiency is also a reason discussed in Reference [23], which could

explain the larger effects on growth rates at lower concentrations. The reason for nutrient limitation induced by plastic is proposed to be interactions of the nutrients with the surface of the plastics [35]. NanoPS<sub>42</sub> self-aggregation could account for the higher nanoPS<sub>42</sub> treatments having less of an effect on the growth rates.

### 3.2. Nano-Plastic Exposure Influences the Number and Sinking Velocity of Cell Aggregates

To understand the impact of nanoPS<sub>42</sub> on aggregation in these two Symbiodiniaceae cultures, the total number of algal aggregates per tank and in five aggregate class sizes was recorded (Figures 2 and S2). All tanks showed aggregation, which was expected, as self-aggregation of Symbiodiniaceae has been observed previously [13].



**Figure 2.** NanoPS exposure leads to a change in aggregation. Aggregates sorted by class size show a significant change in the distribution pattern under nanoPS<sub>42</sub> exposure (Holm-Sidak,  $p = 0.05$ ). No differences are observed when exposure length is compared.

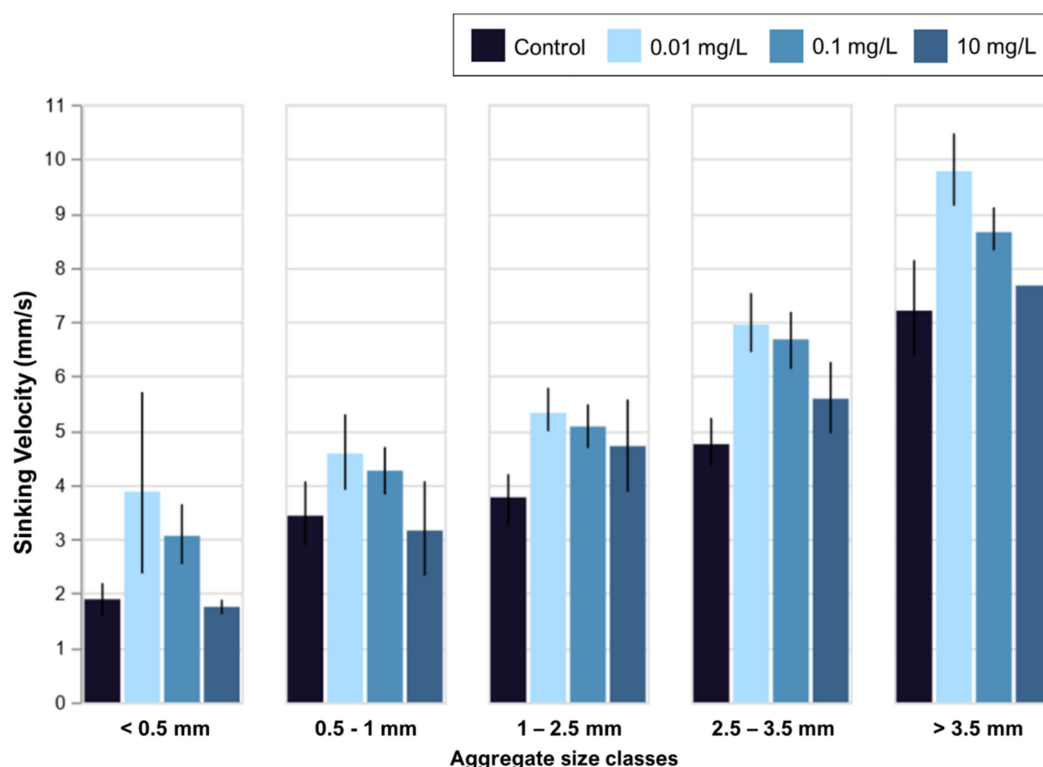
The majority of aggregates exhibited an ovoid form. A significant difference can be observed when aggregate numbers are compared over all class sizes and all treatments, showing that the nanoPS has an influence on the aggregation process. The lowest nanoPS<sub>42</sub> treatments (0.01 mg/L) shows significant reduction in the total aggregates count by 10% (Holm-Sidak,  $p = 0.003$ ), but aggregation was enhanced overall in that treatment to have a higher percentage of huge aggregates than in the control treatment (Holm-Sidak,  $p = 0.001$ ). While there is also a reduction of 3% in the intermediate nanoPS<sub>42</sub> treatment (0.1 mg/L), this is not significant (Holm-Sidak,  $p = 0.319$ ). In the highest plastic treatment at 10 mg/L, this is reversed, leading to more aggregates overall, and more of those being of smaller sizes. The different aggregate class sizes show significantly different distributions in all three treatments and the control (ANOVA,  $p < 0.001$ ) (Figure S2). In the control, the self-aggregation led to a specific distribution pattern of aggregate sizes, which was not repeated in the treatments. Self-aggregation was also observed in the microplastic experiments of Su et al. [13]. The fact that the presence of nanoPS changes the aggregation between the cells and leads to more aggregates in the bigger size classes is possible due to higher production of extracellular polymeric substances (EPS) with sticky properties, trapping more cells in one aggregate and keeping aggregates closer together.

Nutrient depletion, which has been linked to the presence of micro-plastics in algae cultures [35], is associated with increased stickiness of the EPS [36,37]. Differences in the EPS production due to the presence of nanoPS is a likely factor contributing to the differences in aggregation seen in the study. EPS production was not measured, so further studies are needed to confirm this hypothesis linking the aggregation process and EPS production in Symbiodiniaceae under nanoPS influence. Lagarde et al. [33] notices different aggregate formation under different plastic treatment and sizes, which matches with our results. In addition, in *Symbiodinium tridacnidorum*, genes encoding a protein with a TIG (Transcription factor immunoglobulin) domain were upregulated. Since this protein is found in surface cell receptors, it may influence changes in hetero aggregation.

Significant differences are evident when aggregate numbers are compared over size classes and treatments, showing that nanoPS influences aggregation. Aggregate size classes show significantly different distributions in all three treatments vs. controls (ANOVA,  $p < 0.001$ ) (see Figure 2). These differences in aggregation could be due to changes of the cell surface receptors, as nanoPS increases genes related to those two-fold (see Section 3.3. NanoPS effects on gene expression).

Due to nanoPS exposure, aggregation and sinking velocities are impacted, which, in turn, leads to change in sedimentation. Since the majority of the host animals obtain their symbiotic dinoflagellates from the sand and water column [20], these changes in dinoflagellate sedimentation might lead to problems in acquisition of symbionts for the host animals. The lowest plastic treatment used, which is environmentally possible, already induces changes to the sedimentation. This lowest treatment led to bigger aggregates, which, at the same time, sank faster, possibly removing the symbionts from the water column faster than required from the host animals and reducing chances of encountering symbionts.

Changes in aggregation and resulting sedimentation were observed under nanoPS exposure (Figure 3). The biggest changes in sinking velocity correspond to increases in aggregation and are observed in the lowest plastic treatment at 0.01 mg/L. On the other hand, the 10 mg/L treatment did not have any significant effect on the sinking rates but did affect sedimentation indirectly through changes in the aggregate size distribution (Figure 3). These changes, including both sinking velocities and aggregate size distribution, are most likely due to hetero-aggregation between algae and nanoPS. Under different treatments, the size distribution of aggregates was significantly different (Figure 2). In combination, it is likely that the same effect that led to that difference in aggregation is also responsible for the difference in sinking velocities. Changes in EPS production and stickiness will lead to different cell packaging within the aggregates, possibly creating tighter packed aggregates in the lowest and intermediate treatment. This effect might be counteracted under the highest nanoPS exposure by the sheer volume of EPS, which is lighter than seawater. The nano-plastic itself trapped in these could also add to the sinking velocity returning back to control levels in the high plastic treatments. Since these symbionts are paired with the mobile larvae of the host animals, a higher sinking velocity would remove the potential symbiont from the pelagic area and reduce the chance of a match.

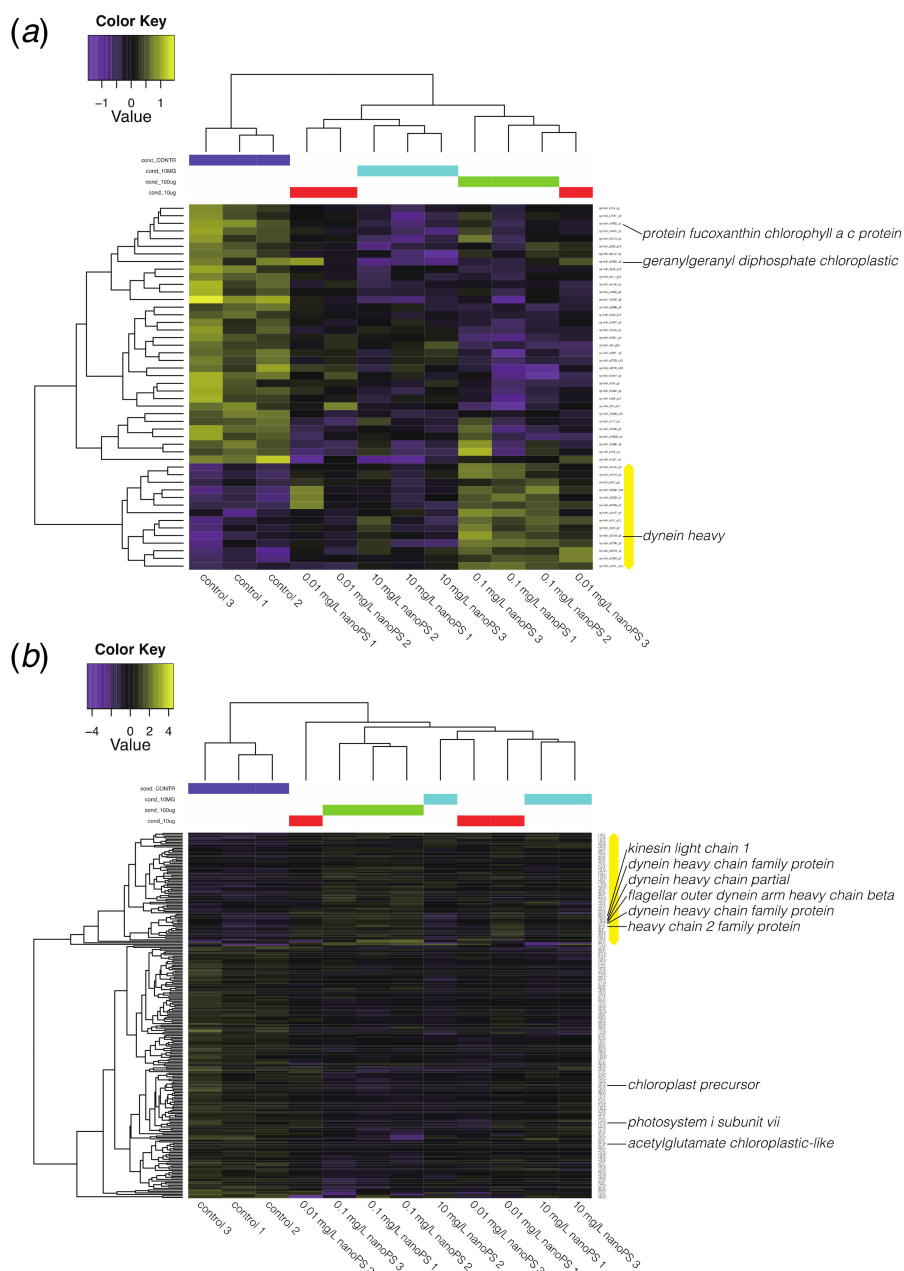


**Figure 3.** Sinking velocity change with nanoPS exposure. Sinking velocities decrease with aggregate size from more than 7 mm/s (>3.5 mm) to less than 2 mm/s (<0.5 mm). In all class sizes, the control was similar in sinking velocity to the highest nanoPS treatment (10 mg/L). The low nanoPS treatment (0.01 mg/L) differed significantly from both controls ( $t$ -test, two-tailed  $p = 5.56 \times 10^{-4}$ ) and the highest nanoPS treatment ( $t$ -test, two-tailed  $p = 9.03 \times 10^{-4}$ ). This was also true for the intermediate nanoPS treatment (darker blue, 0.1 mg/L). Error bars are 95% confidence intervals. Only one huge aggregate was measured in the highest nanoPS treatment. No differences in sinking velocity were observed in relation to exposure length.

### 3.3. NanoPS Effects on Gene Expression

Analysis of differential gene expression showed that, in *Symbiodinium*, 14 genes were upregulated after nanoPS<sub>42</sub> exposure, and 34 were downregulated relative to controls (Figure 4a). In *Cladocopium*, 75 genes were upregulated, and 169 genes were downregulated (Figure 4b). *Cladocopium* seems more sensitive to nanoPS<sub>42</sub> exposure, as overall more genes responded than in *Symbiodinium*. Since Pfam analysis had more annotations than BLAST2GO in differentially expressed genes (DEGs) of *Cladocopium*, we list the major domains encoded by the DEGs of *Cladocopium* (Tables S3–S6).

The largest group of upregulated genes was a subfamily of dynein-related proteins having an AAA\_5 domain (Table 1). Dynein is a microtubule-associated motor protein. Ten genes for dynein-related proteins with AAA and/or DHC (Dynein heavy chain) were upregulated in *Cladocopium* sp. by nanoPS<sub>42</sub> (Table 1 and Table S4). It has been shown that microplastic exposure induces production of reactive oxygen species (ROS) in microalgae [13,33] and dynein upregulation. Therefore, it might be needed to balance cytoskeletal dynamics as microtubule polymerization is impaired by oxidative stress [38]. Dynein light chain genes were also shown to be upregulated in gill cells of zebra mussels exposed to polystyrene micro-plastic [39].



**Figure 4.** Heatmap and clustering of differentially expressed genes (2-fold changes,  $p < 0.001$ ) between dinoflagellates exposed to nano-plastics and controls. **(a)** DEGs in *Symbiodinium tridacnidorum*. **(b)** DEGs in *Cladocopium* sp. Values indicate the relative gene expression level with purple and yellow showing downregulation and upregulation, respectively. The yellow bar shows a cluster of upregulated genes. Annotations by Blast2GO show the presence of microtubule-related or photosynthesis-related genes among DEGs.

**Table 1.** Domains encoded by more than three up-regulated genes in *Cladocopium* sp.

Domain Name	Summary from Pfam Database	Gene Number
AAA_5	AAA domain (dynein-related subfamily)	6
DHC_N2	Dynein heavy chain, N-terminal region 2	5
AAA	ATPase family associated with various cellular activities	4
AAA_6	Hydrolytic ATP binding site of dynein motor region	4
TIG	IPT/TIG domain	4



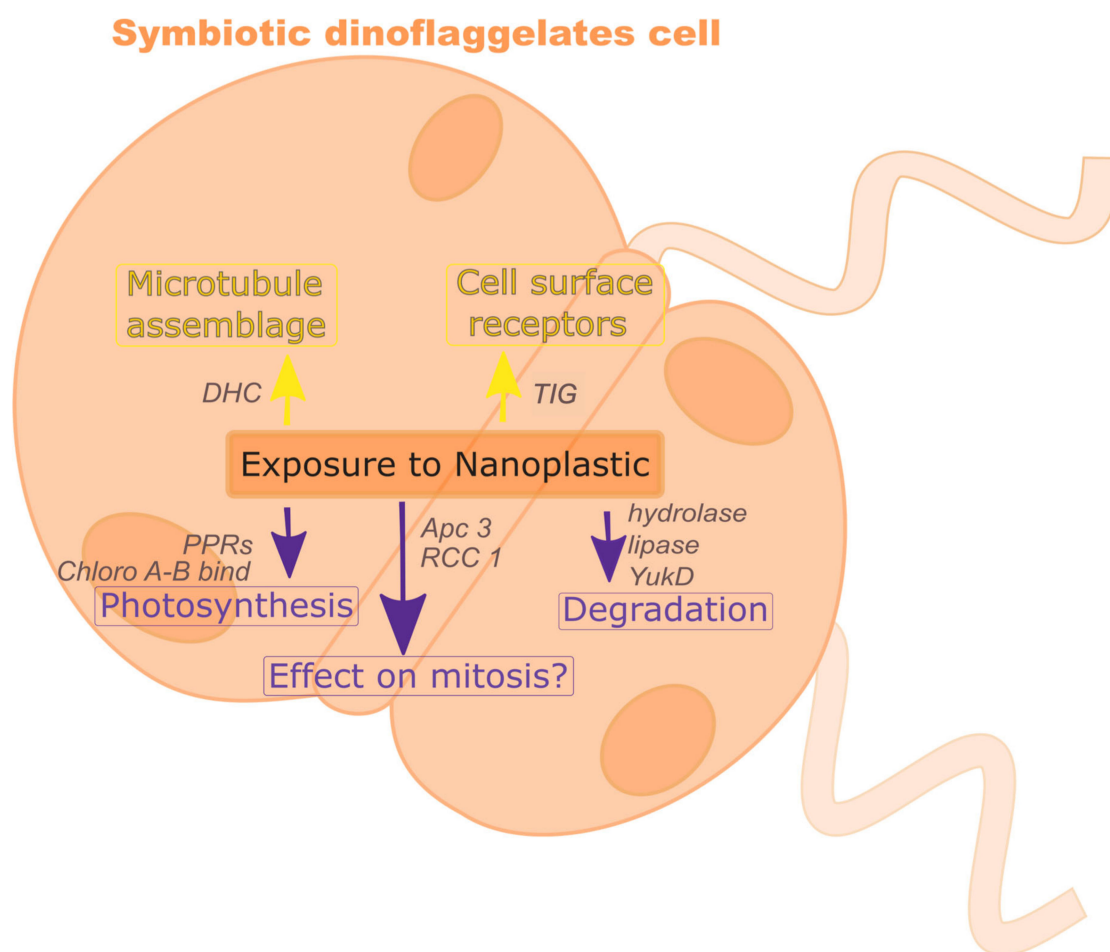
Four upregulated genes in *Cladocopium* (Table 1) encoded proteins with TIG domains that have an immunoglobulin-like fold and are found in cell surface receptors that control cell dissociation [40,41]. This might contribute to adhesion between neighboring cells and to the extracellular matrix composition and explain some of the changes observed in cell aggregations.

There were more downregulated genes than upregulated genes in both *Symbiodinium* and *Cladocopium* (Figure 4). PPR (pentatricopeptide repeat) protein (Table 2) is involved in RNA editing [42] and extensive RNA editing has been reported in organelles of Symbiodiniaceae [43,44]. Five genes for photosynthesis were downregulated (Figure 4). These changes may explain observed reductions in photosystem efficiency in *C. goreau* [13].

**Table 2.** Domains encoded by more than three down-regulated genes in *Cladocopium* sp.

Domain Name	Summary from Pfam Database	Gene Number
Ank	Ankyrin repeat	10
Ank_2	Ankyrin repeats (3 copies)	10
Ank_3	Ankyrin repeat	10
Ank_4	Ankyrin repeats (many copies)	10
Ank_5	Ankyrin repeats (many copies)	10
PPR_2	PPR repeat family	6
RCC1_2	Regulator of chromosome condensation (RCC1) repeat	6
ANAPC3 (Apc3)	Anaphase-promoting complex, cyclosome, subunit 3	5
Pkinase	Protein kinase domain	5
PPR	PPR repeat	5
PPR_3	Pentatricopeptide repeat domain	5
Abhydrolase_5	Alpha/beta hydrolase family	4
Abhydrolase_6	Alpha/beta hydrolase family	4
Lipase_3	Lipase (class 3)	4
PPR_1	PPR repeat	4
TPR_14	Tetratricopeptide repeat	4
YukD	WXG100 protein secretion system (Wss), protein YukD	4

Other downregulated gene groups were related to intracellular degradation processes, including hydrolase and lipase, and to subunit 3 of the anaphase-promoting complex/cyclosome [45]. The downregulated gene (s3282\_g2) with abhydrolase and chlorophyllase domains is likely related to chlorophyll degradation [46]. The gene, s576\_g21, for cell division control (CDC) protein 2 is downregulated in *Cladocopium*. Downregulation of six genes with RCC1 (regulator of chromosome condensation) and three genes with CDC domains suggest some effect on cell division. Thus, several negative consequences of nanoPS<sub>42</sub> exposure are suggested by DEGs (summarized in Figure 5).



**Figure 5.** Exposure to nanoPS<sub>42</sub> changes gene expression levels in symbiotic dinoflagellates. Yellow and purple arrows show up-regulation and down-regulation of gene expression, respectively.

#### 4. Conclusions

Previous studies have shown that nano-plastics have adverse effects on different algae groups [32,34,35,47,48], and a recent study shows that micro-plastics have similarly negative effects on an endosymbiotic dinoflagellate *Cladocopium goreauii* [13]. No previous studies have been conducted on nanoPS<sub>42</sub> effects on Symbiodiniaceae. We found significant changes in aggregation, photosystem efficiency, and aggregate sinking velocity of symbiotic dinoflagellates, which is coupled with variations in gene expression patterns after exposure to nanoPS<sub>42</sub>. The reduction in photosystem efficiency and photosystem gene expression patterns could have led to the observed reduced growth rates and are especially problematic given the obligate photosymbiotic nature of the host animals of the dinoflagellates.

**Supplementary Materials:** The following are available online at <http://www.mdpi.com/2076-2607/8/11/1759/s1>. Figure S1: Cell abundance in treatment tanks, control tanks, and outside controls. Figure S2: NanoPS exposure changes aggregation behaviour, reduces cell numbers, and alters size class distributions. Table S1: Relationship between nanoPS<sub>42</sub> concentration and particles per tank. Table S2: Sampling days of each tank. Table S3: Genes that responded to nano-plastic exposure in *Symbiodinium tridacnidorum*. Table S4: Genes that responded to nano-plastic exposure in *Cladocopium* sp. Table S5: Differentially expressed genes with a two-fold difference between the controls and nano-plastic exposure (*Symbiodinium tridacnidorum* cladeA, TMM FPKM values,  $p < 0.001$ ). Table S6: Differentially expressed genes with a two-fold difference between the controls and nano-plastic exposure (*Cladocopium* sp. cladeC, TMM FPKM values,  $p < 0.001$ ).

**Author Contributions:** C.R. designed the study and performed the experiment. E.S. carried out RNA analyses. K.K. and C.R. performed RNA-seq mapping and cluster analyses. All authors wrote the manuscript. All authors have read and agreed to the published version of the manuscript.

**Funding:** This research received no external funding. This research was funded by Okinawa Institute of Science and Technology Graduate University support for the Light-Matter Interactions for Quantum Technologies (SNC) and the Marine Genomics Unit (NS).

**Acknowledgments:** We are grateful to the DNA sequencing section of OIST for RNA preparation and sequencing and to the OIST imaging section for 3D image support. We thank members of the MGU, especially Haruhi Narisoko for cell culturing and Kanako Hisata for IT support. We are grateful for the help and support provided by K. Deasy from the Engineering Support Section of Research Support Division at OIST. We gratefully acknowledge Sile Nic Chormaic and Noriyuki Satoh for their continuing personal and material support and kind encouragement throughout the project.

**Conflicts of Interest:** The authors declare no conflict of interest.

**Availability of Data:** Data are available in the electronic supplementary material. Raw sequence data are available from PRJNA627564 in NCBI database. Symbiodinium (currently the family Symbiodiniaceae) clade A3 and C genomes: clade A3 ([https://marinegenomics.oist.jp/symb/viewer/info?project\\_id=37](https://marinegenomics.oist.jp/symb/viewer/info?project_id=37)) and clade C ([https://marinegenomics.oist.jp/symb/viewer/info?project\\_id=40](https://marinegenomics.oist.jp/symb/viewer/info?project_id=40)). Transcript models of Symbiodinium clades A3 and C: [https://marinegenomics.oist.jp/symb/download/syma\\_transcriptome\\_37.fasta.gz](https://marinegenomics.oist.jp/symb/download/syma_transcriptome_37.fasta.gz) and [https://marinegenomics.oist.jp/symb/download/symC\\_aug\\_40.fa.gz](https://marinegenomics.oist.jp/symb/download/symC_aug_40.fa.gz).

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