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Effects of auxin derivatives on phenotypic plasticity and stress tolerance in the alga *Desmodesmus* (Chlorophyceae, Chlorophyta)

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The green microalga *Desmodesmus* is characterized by a high degree of phenotypic plasticity, allowing them to be truly cosmopolitan and withstand environmental fluctuations. This flexibility enables *Desmodesmus* to produce a phenotype-environment match across a range of environments broader than that possible if the phenotypic traits were fixed. Indoles and their derivatives are a well-known crucial class of heterocyclic compounds and are widespread in different species of plants, animals, and microorganisms. Indole-3-acetic acid (IAA) may behave as a signaling molecule in microorganisms, and the physiological cues of IAA may also trigger phenotypic plasticity responses in *Desmodesmus*. In this study, we demonstrated that the changes in colonial morphs of Desmodesmus were specific to IAA but not to chemically more stable synthetic auxins, naphthalene-1-acetic acid and 2,4-dichlorophenoxyacetic acid. Moreover, inhibitors of auxin biosynthesis and polar auxin transport inhibited cell division. Notably, different algal species (even different intraspecific stains) exhibited phenotypic plasticity different to that of IAA. Thus, the plasticity involving individual-level heterogeneity in morphological characteristics may be crucial for microalgae to adapt to changing or novel conditions, and IAA treatment potentially increases the tolerance of *Desmodesmus* to several stress conditions. In summary, our results provide circumstantial evidence for the hypothesized role of IAA as a diffusible signal in the communication between the microalga and its associated microorganisms. This information is crucial for elucidation of the role of plant hormones in microalgal ecology.

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Abstract

The green microalga Desmodesmus is characterized by a high degree of phenotypic plasticity, allowing them to be truly cosmopolitan and withstand environmental fluctuations. This flexibility enables *Desmodesmus* to produce a phenotype–environment match across a range of environments broader than that possible if the phenotypic traits were fixed. Indoles and their derivatives are a well-known crucial class of heterocyclic compounds and are widespread in different species of plants, animals, and microorganisms. Indole-3-acetic acid (IAA) may behave as a signaling molecule in microorganisms, and the physiological cues of IAA may also trigger phenotypic plasticity responses in *Desmodesmus*. In this study, we demonstrated that the changes in colonial morphs of *Desmodesmus* were specific to IAA but not to chemically more stable synthetic auxins, naphthalene-1-acetic acid and 2,4-dichlorophenoxyacetic acid. Moreover, inhibitors of auxin biosynthesis and polar auxin transport inhibited cell division. Notably, different algal species (even different intraspecific stains) exhibited phenotypic plasticity different to that of IAA. Thus, the plasticity involving individual-level heterogeneity in morphological characteristics may be crucial for microalgae to adapt to changing or novel conditions, and IAA treatment potentially increases the tolerance of *Desmodesmus* to several stress conditions. In summary, our results provide circumstantial evidence for the hypothesized role of IAA as a diffusible signal in the communication between the microalga and its associated microorganisms. This information is crucial for elucidation of the role of plant hormones in microalgal ecology.

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Keywords Coenobial algae · *Desmodesmus* · Indole derivatives · Microalgae · Phenotypic plasticity



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Introduction

37 Phenotypic plasticity can be broadly defined as the capacity of a single genotype to exhibit variable phenotypes in different environments and implies that a species can conquer diverse 38 39 environments. A well-known example of phenotypic plasticity is changes in multicelled 40 structures in the coenobial algae. Most studies on phenotypic plasticity in coenobial algae have 41 been conducted considering morphological responses to an abiotic factor. Neustupa and Hodač 42 demonstrated that morphological plasticity of *Pediastrum duplex* var. duplex is related to the pH 43 dynamics of freshwater lakes (Neustupa & Hodac, 2005). Peña-Castro et al. also reported the 44 phenotypic plasticity in Scenedesmus incrassatulus in response to heavy metal stress (Pena-Castro et al., 2004). However, microalgae are typically associated with other microorganisms, 45 such as zooplankton, fungi, and bacteria. Thus, studies on phenotypic plasticity of the coenobial 46 47 algae have increased in number and broadened their scope from the initial focus on abiotic factors to that on biotic ones. Hessen and Van Donk first indicated that the presence of the 48 49 grazing pressure from water flea (Daphnia magna) can induce colony formation in Scenedesmus algae (Hessen & Van Donk, 1993). Furthermore, Lurling et al. proved that the induced colony 50 51 formation in the presence of herbivores is considered a strategy more efficient than constitutive 52 defenses under variable grazing risk (Lürling & Van Donk, 1996; Lürling, 2003). Wu et al. 53 further revealed that the number of cells per coenobium of Scenedesmus increased with the 54 density of *Daphnia* growth, thus indicating a grazer density-dependent response (Wu et al., 55 2013).

Auxins, which constitute a class of plant hormones, have previously been suggested to regulate physiological responses and gene expression in microorganisms (Spaepen et al., 2007). Furthermore, indole-3-acetic acid (IAA) is one of the most physiologically active auxins that can be produced by a numerous microbial species (Spaepen et al., 2007; Fu et al., 2015). Furthermore, the phylogenetic analyses have revealed that IAA biosynthetic pathways evolved independently in plants, bacteria, algae, and fungi (Fu et al., 2015). The convergent evolution of IAA production leads to the hypothesis that natural selection might have favored IAA as a widespread physiological code in these microorganisms and their interactions. In natural water bodies, the crucial physical associations and biochemical interactions between microalgae and other microorganisms are generally well recognized (Natrah et al., 2014). Piotrowska-Niczyporuk and Bajguz found that IAA plays a crucial role in the growth and metabolism of Chlorella vulgaris during a 72-hour culture period (Piotrowska-Niczyporuk & Bajguz, 2014). Jusoh et al. indicated that IAA can induce changes in oil content, fatty acid profiles, and expression of four genes responsible for fatty acid biosynthesis in C_k vulgaris at early stationary growth phase. In addition, the significance of these interactions in algal phenotypic plasticity has attracted considerable scientific attention (Lürling & Van Donk, 1996; Lürling & Van Donk,



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2000; Lürling 2003 Ve previously used IAA as a signal molecule in microorganisms to simulate a selection pressures caused by interspecific competition. The results indicated that the mean number of cells per particle of Desmodesmus opoliensis and D. komarekii decreased gradually as the IAA concentration increased gradually. The proportion of *Desmodesmus* unicells in monocultures increased with IAA concentration. We also demonstrated that these unicells exhibited a lower tendency to sedimentation than did large cells and that shrinkage may facilitate nutrient uptake and light capture (Chung et al., 2018). However, whether other coenobial algal species of *Desmodesmus* use the same strategy to overcome stress remains unknown. Hence, the objective of the present study was to compare the effects of IAA at different concentrations on phenotypic responses in different Desmodesmus species. Moreover, to address the auxin specificity of these processes and obtain an insight into the complex auxinrelated regulatory mechanism(s) in algal physiology, we have selected a group of compounds called "auxin analogs," such as synthetically produced naphthalene-1-acetic acid (NAA) and 2,4dichlorophenoxyacetic acid (2,4-D), which are structurally related to IAA. We thus aim to determine the differential effects of auxins and auxin-like compounds on the morphological responses of these coenobial algae. Furthermore, IAA has been detected in some species of Scenedesmaceae microalgae (Mazur et al., 2001; Prieto et al., 2011) Therefore, we investigated the effects of inhibitor of auxin biosynthesis and auxin transport in *Desmodesmus*. To elucidate the physiological changes induced by phytohormone treatment, we also investigated whether IAA pretreatment promotes an enhanced stress-tolerant phenotype. The obtained results are crucial for elucidation of the role plant hormones in microalgal physiology.

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Materials and Methods

95 Isolation and Culture of Microalgae

The algal strains used here were isolated from natural water bodies in Central Taiwan. Water samples with visible microalgal population were centrifuged at 3000 ×*g* for 10 minutes at room temperature to concentrate the cells and spread onto CA agai ates. For isolating an axenic single colony from field water samples, the streak plate method was used. The algae were cultured in CA medium, consisting of 2 mg/L Ca(NO₃)₂.4H₂O, 10 mg/L KNO₃, 5 mg/L NH₄NO₃, 3 mg/L β–Na₂glycerophosphate.5H₂O, 2 mg/L MgSO₄.7H₂O, 0.01 μg/L vitamin B12, 0.01 μg/L biotin, 1 μg/L thiamine HCl, and 0.1 mL/L PIV metals (1000 mg/L Na₂EDTA.2H₂O, 196 mg/L FeCl₃.6H₂O, 36 mg/L MnCl₂.4H₂O, 10.4 mg/L ZnCl₂, 4 mg/L of CoCl₂.6H₂O, and 2.5 mg/L of Na₂MoO₄.2H₂O), 0.1 mL/L Fe (as EDTA; 1:1 molar; 702 mg/L Fe(NH₄)₂(SO₄).6H₂O and 660 mg/L Na₂EDTA.2H₂O), and 40 mg/L of HEPES; the pH was then adjusted to 7.2 colated algal cells were stored at −80°C in 15%–20% glycerol. For each experiment, the alga was cultured axenically in liquid CA medium at 125 rpm in a tube rotator and grown at 25°C under cool white



108 fluorescent light (approximately 46.30 μmol m⁻² s⁻¹) with a 14:10-h light–dark period.

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- 110 Algae Identification
- 111 The algal cells were harvested by centrifugation at 3000 $\times g_{\overline{5}}$ at 25°C for 10 minutes. The
- 112 genomic DNA used for analysis was isolated using AccuPrep GMO DNA Extraction Kit
- 113 (Bioneer, Korea). The 18S rDNA was amplified through PCR by using the following primers:
- 114 18S forward-TTTCTGCCCTATCAACTTTCGATG and 18S reverse-
- 115 TACAAAGGCAGGACGTAAT, which yielded a fragment of approximately 1200 bp (Pan
- et al., 2011). The PCR conditions were as follows: initial denaturation at 96°C for 4 minutes; 36
- cycles of denaturation at 96°C for 30 s, annealing at 50°C for 30 s, and extension at 72°C for 1
- minutes; and final extension at 72°C for 6 minutes. The ITS1-5.8 S-ITS2 region rDNA was
- amplified using the primers ITS forward1 (ACCTAGAGGAAGGAGAAGTCGTAA) and ITS
- reverse1 (TTCCTCCGCTTATTGATATGC), which provided a fragment of approximately 1200
- bp (Pan et al., 2011). The PCR conditions are as follows: initial denaturation at 96°C for 4
- minutes; 36 cycles of denaturation at 96°C for 30 s, annealing at 48°C for 30 s, and extension at
- 72°C for 1 minutes; and final extension at 72°C for 6 minutes. DNA sequencing was performed
- by Tri-I Biotech, Incide he Basic Local Alignment Search Tool was used to find regions of local
- 125 similarity between sequences on the website of the National Center for Biotechnology
- 126 Information (http://www.ncbi.nlm.nih.gov).

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- 128 Experimental Design
- 129 Solutions containing different concentrations of phytohormones (IAA, NAA, and 2,4-D) and
- auxin-related compounds (4-biphenylboronic acid and 2,3,5-triiodobenzoic acid) were prepared
- to investigate their influence on the growth and morphological plasticity of *Desmodesmus*. The
- concentrations of each phytohormone and compounds used in each experiment depended on the
- sensitivity of each species. Algae were harvested after each experiment, and the proportions of
- 134 different-celled populations were calculated under an optical microscope (DMRB, Leica,
- 135 Germany). The proportion of different algal populations (including unicellular; two-, four-, and
- eight-celled; and other colonial morphs) were calculated, and the mean numbers of cells in
- different morphotypes were calculated. The numbers of cells per coenobium were counted by
- dividing the total cell number by the number of coenobia.

- 140 Transmission Electron Microscopy
- All specimens were prefixed in 2.5% glutaraldehyde/0.1 M sodium cacodylate buffer (pH 7.3)
- containing 1% tannic acid at 4°C overnight. After in 0.1 M sodium cacodylate buffer with 5%
- sucrose for 15 minutes were washed three times, specimens were postfixed with 1% osmium



144 tetroxide in 0.1 M sodium cacodylate buffer at 4°C overnight. Specimens were then washed in buffer, en bloc stained with 2% aqueous uranyl acetate, dehydrated through a graded series of 145 ethanol and two times with 100% acetone. Specimens were infiltrated with Spurr resin overnight 146 147 and embedded in fresh Spurr resin the next day. Serial ultrathin sections of approximately 70 nm 148 were cut with a diamond knife on a Leica Ultracut R ultramicrotome (Leica, Heerbrugg, Switzerland) and examined with a Hitachi H-7500 transmission electron microscope (Hitachi, 149 Tokyo, Japan) at 80 kV. Images were recorded using a 2048 × 2048 Macrofire monochrome 150 151 CCD camera (Optronics, Goleta, CA, USA).

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- 153 Stress Tests
- 154 The culture samples were harvested; the cells were then washed with CA medium and 155 resuspended in the CA medium with different treatments. For osmotic shock test, the cells were incubated in the CA medium with 0.5 M NaCl. For the acidic pH assay, the culture samples were 156 157 resuspended in CA medium at pH 3.0 (adjusted with HCl). For oxidative stress, the cells were 158 exposed to hydrogen peroxide at final concentration of 5 mM. The cell suspensions subjected to the aforementioned treatments were shaken at 25°C for 15 or 30 minutes. For inducing heat 159 shock, the cells were exposed to 40°C for 10, 15, or 20 minutes by immersing the cultures in a 160 161 shaking water bath. For cold treatment, the cultures were exposed to 4°C for 24 hours. Fractions of viable cells of each experiment were determined by plating appropriate dilutions of the 162 cultures on CA agar plates before and after treatments. The controls (without IAA treatment) 163 received the same treatments used throughout the procedure. 164

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166 Statistical Analysis.

Data are presented as means of three replicates ± their standard deviations (SDs). The proportions of colonies with different numbers of cells and mean number of cells per coenobium were compared using a one-way analysis of variance (ANOVA) with least significant difference post hoc test. In stress tests, the significance of differences between the groups was determined using the Student t test and ANOVA. A p of <0.05 was considered statistically significant.

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Results

- 174 Effect of Auxin Analogs on Algal Growth and Phenotypic Plasticity Induction
- 175 In a previous study, we performed a dose–response analysis to determine the fitness effects of
- 176 IAA on the coenobial algae D. komarekii (Chung et al., 2018). The results revealed that different
- concentrations of IAA had different effects on the growth and morphological changes of D.
- 178 komarekii. Thus, we concluded that Desmodesmus can respond to the external phytohormone
- 179 IAA signal and then integrate the information to initiate physiological changes. In this study, our



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aim was to determine whether the physiological cues of IAA-related compounds also trigger the growth and phenotypic plasticity responses in *Desmodesmus*. With respect to *D. komarekii* growth, we examined the effects of the natural auxin IAA as well as those of the synthetic auxins NAA and 2,4-D. At 300 μM, IAA, NAA, and 2,4-D clearly inhibited growth; however, IAA caused lower inhibition than did NAA and 2,4-D (Fig. 1a). This inhibitory effect was also observed at 100 and 200 μM NAA and 2,4-D, but not in the cells treated with 100 μM IAA (Fig. 1b, c). These observations indicated that these auxin-related compounds inhibit *D. komarekii* growth.

To measure phenotypic plasticity responses in algal populations, monocultures of D. komarekii were used. After 1 week of culturing, the monocultures of D. komarekii in the groups with exogenous 300 µM IAA and synthetic auxins were compared with those in the control environment (without treatment). We found that the monocultures of D. komarekii in the control groups (without IAA treatment) were dominated by one- and four-celled coenobia (Fig. 2a-c). The morphology of D. komarekii monocultures changed drastically compared with the control after exposure to IAA (Fig. 2b, c). The proportion of unicells increased rapidly from day 3, and the proportion of four-celled coenobia decreased (Fig. 2a, c). The mean number of cells per particle reached its minimum on approximately day 7 (Fig. 2a), the proportion of unicells increased from 37% to approximately 73%, and the proportion of four-celled coenobia decreased from 49% to approximately 16% on day 9. The proportion of two-celled coenobia changed only slightly from approximately 13% to approximately 7%. The mean number of cells per particle in the control groups remained at >2.5 during the 9-day period. In this experiment, the D. komarekii population of each culture was composed of unicells and two-, four-, and eight-celled colonies; a few three-, five-, six-, and seven-celled colonies were also present, but coenobia with more than eight cells were not observed.

Through transmission electron microscopy (TEM) analysis, we confirmed that the morphological changes in coenobia were not caused by cell aggregation but by the vegetative growth of a mother cell (Fig. 3a). No extracellular matrix was seen on or around the eells, and the connecting strands between cells were highly visible (Fig. 3b). Dense section of warty layer, can be seen over each coenobial junction (yellow circle; Fig. 3c). Notably, we observed that specific large unicells were formed in the monocultures of *D. komarekii* after day 5 under IAA treatment. Thus, the samples collected at day 7 after IAA treatment and the cells in the control groups were used for observation of morphology through TEM. The accumulation of many starch granules and lipid bodies was observed in the large unicells compared with the cells in control groups (Fig. 3d). By contrast, we found that the auxin-related compounds NAA and 2,4-D both inhibit the growth of *D. komarekii* in dose-dependent manner, but they did not influence their number of cells of individual Fig. 2d, e). We next tested the effects of an auxin



biosynthesis inhibitors and a polar auxin transport inhibitor. Here, 4-biphenylboronic acid (BBo), a potent YUCCA enzyme inhibitor and *Arabidopsis* growth inhibitor, and 2,3,5-triiodobenzoic acid (TIBA), a polar auxin transport inhibitor, were used [(Dhonukshe et al. 2008; Kakei et al. 2015)]. BBo strongly inhibited growth even at 100 μM and its inhibitory effect increased with its concentration (Fig. 1c). At 200 and 300 μM, both of TIBA and BBo inhibited *D. komarekii* growth (Fig. 1a, b). These results suggested that inhibition of auxin transport and inhibition of YUCCA function both inhibit cell growth.

The auxin-like physiological competence of selected compounds was analyzed in *Desmodesmus* based on the inhibition of growth in liquid cultures and morphological changes. Thus, we performed a dose–response analysis to determine the fitness effects of IAA and other analogs on eight other *Desmodesmus* strains. The results revealed that different concentrations of indole derivatives had divergent effects on the growth of different *Desmodesmus* species (Figs. 4–11). In general, high concentrations (>300 μM) of IAA and other analogs inhibited the growth of the algal population. Thus, *Desmodesmus* can respond to the external phytohormone signal of IAA and other analogs and then integrate the information to initiate physiological changes. In the subsequent experiment, our aim was to determine whether the physiological cues of IAA and other analogs in these cultures also trigger phenotypic plasticity responses.

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Plastic Phenotypic Changes in Response to IAA Are Strain-Dependent Behaviors

235 To measure the phenotypic plasticity responses to indole derivatives on four strain of D. armatus, 236 two strains of D. communis, and one strain each of D. intermedius and D. opoliensis were used in 237 this study. After 1 week of treatment, the monocultures of D. armatus in the control groups 238 (without indole derivatives treatment) were compared with indole derivatives. We found that the 239 changes in colonial morphs in D. armatus are specific to IAA but not to chemically to synthetic auxins, NAA and 2,4-D, which are chemically more stable than IAA (Figs. 4-6). Moreover, we 240 241 found the different algal strains of D. armatus demonstrating phenotypic plasticity different to 242 IAA. In D. armatus JYCA037, the monocultures in the control groups were dominated by twoand four-celled coenobia, with <2% unicells (Fig. 4a, b). The morphology of D. armatus 243 244 JYCA037 populations considerably changed under high concentration of IAA treatment 245 compared with that in the control environment (without IAA addition). When the IAA 246 concentration increased, the proportion of four-celled coenobia declined from >90% to approximately 21%, and the number of unicells increased from <2% to approximately 12% and 247 two-celled coenobia increased from approximately 7% to 66%. The mean number of cells per 248 249 particle of D. armatus JYCA037 decreased gradually as the IAA concentration gradually 250 increased, and the cell number reached its minimum level at an IAA concentration of 400 µM (Fig. 4a). Similar results were observed in the monocultures of D. communis JYCA040; it was 251



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252 dominated by two- and four-celled coenobia in the control groups (Fig. 8). When IAA 253 concentration increased, the proportion four-celled coenobia decreased, and the number of unicells increased. The mean number of cells per coenobium particle in the control groups of 254 255 these two strains remained for >3 days after 7-day culturing. By contrast, in D. armatus 256 JYCA041, the monocultures in the control groups were dominated by unicells (47%), with 47% two- and <7% four-celled individuals (Fig. 4c, d). The morphology of D. armatus JYCA041 populations changed considerably under high concentration of IAA treatment compared with that 258 259 in the control environment (without IAA addition). When IAA concentration increased, the 260 proportion of two-celled coenobia increased from approximately 47% to approximately 69% and the number of unicells declined from approximately 47% to approximately 28%. The proportion 262 of four-celled coenobia only slightly changed from approximately 6% to approximately 3%. The 263 mean number of cells per particle of D. armatus JYCA041 increased gradually as the IAA concentration gradually increased and reached its maximum level at an IAA concentration of 400 264 μM (Fig. 4b) Similarly, the mean number of cells per particle of D. armatus JYCA039 increased 266 gradually as the IAA concentration gradually increased and reached its maximum level when the IAA concentration was approximately 200 µM (Fig. 7a, b). Notably, the aformentioned morphological changes were not observed in D. armatus JYCA045 even under treatment with 268 high concentrations with IAA (Fig. 4e, f). By contrast, we found that the auxin-related 270 compounds NAA and 2,4-D both inhibit *Desmodesmus* growth in a dose-dependent manner, but the treatment did not influence their number of cells of individuals in these four D. armatus 272 strains (Figs. 4 and 5). The strain-dependent response to IAA but not to NAA and 2.4-D also occurred in one strain of D. communis (JYCA040; Fig. 8) and D. opoliensis (JYCA043; Fig. 9). 273 274 However, the phenotypic plasticity caused by auxin analogs was not obviously shown in one strain of D. communis (JYCA044; Fig. 10) and D. intermedius (JYCA042; Fig. 11).

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277 IAA Improves Algal Defenses Against Stress

> In this study, we found that starch granules and lipid bodies accumulated in algal cells grown at a high IAA concentration. In this environment, algal cells also demonstrated slow growth. Thus, algae contain storage in the form of natural oils, such as neutral lipids or triglycerides, and algal growth diminishes when exposed to stresses. Thus, we propose that the morphological responses and the associated physiological changes provide some fitness advantages to *Desmodesmus*, such as the ability to survive in the water bodies often exposed to fluctuations in environmental factors. The data reported in Table 1 showed that IAA-treated cells could withstand sudden changes in the environment, demonstrating significantly longer survival rates in the media subjected to temperature shock (40°C, 15 minutes and 4°C, 24 hours) and acid treatment (pH 3.0, 15 minutes). The data also shows that IAA treatment also marginally but significantly increased



the survival rate of microalgae treated with 5 mM H_2O_2 for 30 minutes (p = 0.052). However, although the survival rates did not increase significantly, the average survival rates of IAA-treated cells were higher than those of the controls in many treatments.

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Discussion

A central-orienting question in biodiversity theory and ecology is the "paradox of the plankton," which indicates that the number of coexisting planktonic species far exceeds the expected and explicable number based on competition theory (Hutchinson, 1961). Ecologists have provided multiple solutions to the paradox by applying game theory, chaos, tradeoffs, and many other concepts in the past five decades (Tilman, 1994; Huisman & Weissing, 1999; Károlyi et al., 2000; Kerr et al., 2002; Goyal & Maslov, 2018). A leading theory to explain the paradox is that individual variability maintains high biodiversity in planktonic microorganisms (Menden-Deuer & Rowlett, 2014). In aquatic ecosystems, significant evidence supports individual variability; in individual behaviors or physiology, among planktonic microorganisms. This phenotypic plasticity has played a central role in studies on the evolution of diversity. Ecologically, phenotypic plasticity has been considered particularly crucial when environmental changes occur and different phenotypes have different fitness values across environments that decide the survival an individual in the face of environmental changes (West-Eberhard, 1989). The plasticity even can potentiate evolvability of microorganisms by opening up new regions of the adaptive landscape (Yi & Dean, 2016).

In our study, we revealed that the morphological characteristics of *Desmodesmus* changed considerably when exposed to IAA compared with the algal cells in the control environment. We found that the algal strains we assayed here have different response patterns to the external IAA. In this study and our previous study, we found that when IAA concentration increased, the mean number of cells per particle of some *Desmodesmus* species decreased (Chung et al., 2018). The surface-to-volume ratios of the unicells was larger than the colony cells in microalgae. Previous literatures have reported that the changes in colony size influence algal surface-to-volume ratios, and the surface-to-volume ratio can affect light capture and nutrient uptake (Reynolds, 2006; Steele et al., 2009). Notably, in this study, we found that in some algal strains, this trend was reversed: the mean number of cells per particle of some Desmodesmus strains was increased when IAA concentration increased. These colonial populations have higher sinking velocities than the unicells and two-celled coenobia. The colonial populations have higher sinking velocities than unicellular cells; consequently, their competitive ability of microalgae might be altered (Lürling, 2003). Thus, plasticity involving individual-level heterogeneity in behaviors and physiological characteristics is crucial for planktonic microorganisms to adapt to changing or novel conditions. This may suggest that individual variability is perhaps the key mechanism



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supporting planktonic biodiversity.

In this study, two widely used auxins in plant tissue culture, NAA and 2,4-D, were also used to investigate their effect on algal growth and physiological responses. These synthetic auxins show varying degrees of auxin-like activity in different bioassays (Abebie et al., 2007; Savaldi-Goldstein et al., 2008). For instance, in Simon et al., the seedlings of Arabidopsis thaliana and suspension-cultured cells of *Nicotiana tabacum* BY-2 were used to investigate the physiological activity of several auxin analogs, along with their capacity to induce auxin-dependent gene expression, to inhibit endocytosis and to be transported across the plasma membrane (Simon et al., 2013). The authors concluded that the major determinants for the auxin-like physiological potential of a particular compound are highly complex and involve its chemical and metabolic stability, its ability to distribute in tissues in a polar manner, and its activity toward auxinsignaling machinery. Thus, the distinct behavior of some synthetic auxin analogs suggests that they might be useful tools in investigations of the molecular mechanism of auxin action. Ohtaka et al. also examined the responses of the natural auxin (indole-3-butylic acid; IBA) as well as the synthetic auxins (NAA and 2,4-D) on the charophyte alga Klebsormidium nitens (Ohtaka et al., 2017). Consistent with our results, the authors indicated that these auxin-related compounds all inhibit K. nitens growth in a dose-dependent manner. Notably, the IAA was detected in cultures of K. nitens, but K. nitens lacks the central regulators of the canonical auxin-signaling pathway found in land plants. However, the authors found that the exogenous IAA inhibited cell division and elongation, and this treatment rapidly induced expression of the transcription factor lateral organ boundaries-domain. During evolution, K. nitens may have acquired a primitive auxinresponse pathway to regulate transcription and cell growth. Here, we found that the natural auxin IAA and the synthetic auxins NAA and 2,4-D can all influence Desmodesmus growth rate. However, the changes in the colonial morphs in *Desmodesmus* are specific to IAA, but not to chemically more stable synthetic auxins. These studies have suggested that structure-activity relationships determined precisely at the level of a particular protein (e.g., receptor or carrier) may not correspond completely to the final auxin-like physiological activity of a particular compound in the streptophytes and their sister group, the chlorophytes. Thus, the comparison of the structure-activity relationships for the aforementioned phenotypic changes highlights differences in the structural requirements of these auxin-related physiological processes, thus making the differential (or the same) phenotypic outcome of the same (or different) compound a very crucial aspect of auxin biology.

Microalgae are unicellular photosynthetic microorganisms, typically found in freshwater and marine systems. The high flexibility and adaptability of this extremely diverse group of eukaryotic organisms enable it to grow in diverse environments, including fresh saltwater, blackish, marine, and soil environments. These coexist with heterotrophic microorganisms, and



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the exchange of chemical compounds is central to the interactions of microalgae with other microorganisms. How microalgal-microbial interactions and participating chemical compounds shape their communities and considerably affect their fitness remains unknown (Hom et al., 2015). Notably, not only plants but also bacteria, fungi (including yeast), and even some microalgae produce or respond to IAA (Fu et al., 2015). Researchers have hypothesized that the microbes sense environmental IAA concentrations to determine the cell density of its competitors (Spaepen et al., 2007; Fu et al., 2015; Chung et al., 2018). Thus, IAA has been speculated to be a signal that coordinates microbial behavior to enhance protection against damage by adverse conditions (Bianco et al., 2006; Chung et al., 2017). Here, we confirmed that the physiological changes in response to IAA confers a fitness advantage by promoting the ability of Desmodesmus to survive in their niches that often undergo fluctuations in environmental factors, such as temperature, osmotic pressure, reactive oxygen species, and pH changes. Under unfavorable stress conditions, such as nutritional starvation, salinity stress and high light intensity, lipid production is usually enhanced in algal cells, due to shifts in lipid biosynthetic pathways toward neutral lipid accumulation (Sun et al., 2018). Microalgae generally accumulate neutral lipids, mainly in the form of triacylglycerols (TAG) under environmental stress conditions. The accumulation of TAG likely occurs as a means of creating an energy deposit that can be readily used in response to a more favorable environment allowing for rapid growth (Tan & Lee, 2016). In green algae, stress conditions also trigger the accumulation of starch granules in the cells, with starch accumulation preceding the accumulation of lipid bodies following stress onset (Siaut et al., 2011). It is generally assumed that the starch and TAG serve as electron sinks under conditions where photosynthesis or metabolism of an exogenous carbon source remains active but the growth is limited (Hu et al., 2008). This phenomenon suggests that carbon sources in algal cells during stress conditions were allocated to not only storage lipid production but also starch biosynthesis, and this finding demonstrates the possibility of partitioning manipulation in the cells. To link physiological changes to phenotype, we performed various cell viability assays in response to heat, cold, osmotic stress, oxidative stress, reactive oxygen species, and pH changes. We found an increased ability to tolerate these stresses, thereby confirming the inferred enhanced stress-tolerant phenotypes when exposed to IAA. The results are consistent with earlier research that on bacteria and found enhanced stress tolerance when the bacteria were pretreated with IAA across various stress conditions (Bianco et al., 2006; Imperlini et al. 2009; Donati et al., 2013).

In natural water bodies, the importance of physical associations and biochemical interactions between microalgae and microorganisms is generally well appreciated, but the significance of these interactions to microbial ecology has not been investigated. In our previous study, we found that a low concentration of IAA promoted the growth of algal cells, but high





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concentrations of IAA inhibited cell growth (Chung et al., 2018). Herein, we further proved that the effects of exogenous IAA and on algal growth and phenotypic changes is species- and even strain-dependent. IAA can exert stimulatory and inhibitory effects on not only algae, fungi, and yeast but also bacteria (Prusty et al., 2004; De-Bashan et al., 2008; Hu et al., 2010; Kerkar et al. 2012; Kulkarni et al., 2013; Sun et al., 2014; Liu et al., 2016; Fu et al., 2017). Bagwell et al. reported the frequency of co-occurrence between IAA-producing bacteria and green algae in natural and engineered ecosystems and revealed that the chlorophyll content and dry weight of algal cells were IAA concentration-dependent responses (Bagwell et al., 2014). A recent study also indicated that IAA produced by associated bacteria was transferred to diatom and influence their growth in exchange for organosulfur compounds (Amin et al., 2015). Thus, exposure to IAA could be likely to affect the outcome of competition among these coexisting organisms. We finally suggested that both algae and microorganisms altered their metabolism to defend themselves form their competitors (or suit each other's needs), and this interaction is potentially very prevalent in the aquatic ecosystems. These findings indicated that IAA is a major factor determining the competition (or mutualistic interactions) between microbial species occupying the same niche.

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Table 1 Increased resistance of *D. komarekii* cells to various stress conditions after exposure to IAA

	Survival (%)	
	Control	IAA-treated
Heat-shock (40°C, 10 mins)	88.0±5.7	93.4±5.2
Heat-shock (40°C, 15 mins)	74.9±2.5	83.9±4.1*
Heat-shock (40°C, 20 mins)	61.2±19.3	79.5±2.5
Cold-shock (4°C, 24 hrs)	25.2±16.4	50.0±2.6*
Osmotic shock (0.5 M NaCl, 15 mins)	46.4 ± 6.4	48.6±4.9
Osmotic shock (0.5 M NaCl, 30 mins)	34.4 ± 6.9	40.6±6
Oxidative stress (2 mM H ₂ O ₂ , 15 mins)	74.7 ± 24.4	79.6±5.7
Oxidative stress (2 mM H ₂ O ₂ , 30 mins)	54.5±9.7	68.6 ± 7.3
Acid shock (pH 3.0, 15 mins)	76.8±1.2	87.9±6.7*
Acid shock (pH 3.0, 30 mins)	54.8 ± 20.2	59.4±16.5
Alkaline shock (pH 8.0, 15 mins)	83.7±9.5	79.6±7
Alkaline shock (pH 8.0, 30 mins)	62.6±13.9	77.3±4.9

Reported values are mean of three measurements \pm their standard deviations. The significance of differences between groups was determined using Student t tests and analyses of variance. *p < 0.05.



571 Figure legends

- 572 Fig. 1 Growth of coenobial algae Desmodesmus komarekii in the presence of several auxins,
- 573 inhibitor of auxin biosynthesis and auxin transport. D. komarekii was cultured in the presence of
- a 300, b 200, and c 100 μM indole-3-acetic acid (IAA), 2,4-dichlorophenoxyacetic acid (2,4-D),
- 575 naphthalene-1-acetic acid (NAA), 2,3,5-triiodobenzoic acid (TIBA), or 4-biphenylboronic acid
- 576 (BBo). Growth curves of *D. komarekii* for each compound were measured at 1, 3, 5, 7 and 9 days.
- 577 Error bars represent standard deviation of values for three replicates.

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- Fig. 2 Mean number of cells per coenobium and proportions of unicells and of two- and four-
- 580 celled coenobia of Desmodesmus komarekii cultured at 300 μM indole-3-acetic acid (IAA), 2,4-
- 581 dichlorophenoxyacetic acid (2,4-D), and naphthalene-1-acetic acid (NAA) concentrations and
- cells without treatment. Data are presented as means (n = 3) for each group, and morphotype
- percentages and cell types were based on 200 cell counts in each repeat. Means with the same
- letter are not significantly different from each other according to a one-way analysis of variance
- and least significant difference post hoc test.

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- Fig. 3 Transmission electron micrographs of *Desmodesmus komarekii* cells under indole-3-acetic
- 588 acid (IAA) treatment. a Through transmission electron microscopy, we confirmed that the
- 589 morphological changes in coenobia were not caused by cell aggregation but by the vegetative
- 590 growth of a mother cell. **b** No extracellular matrix was seen on or around the cells, and the
- 591 connecting strands between cells were highly visible. c Dense section of the warty layer can be
- seen over each coenobial junction (yellow circle). d The accumulation of many starch granules
- 593 (S) and lipid bodies (L) was observed in the large unicells at day 7 after IAA treatment compared
- with the cells in control groups.

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- Fig. 4 Mean number of cells per coenobium and proportions of unicells and two- and four-celled
- 597 coenobia in three strains of Desmodesmus armatus cultured at different indole-3-acetic acid
- 598 (IAA) concentrations. Data are presented as means (n = 3) for each group, and morphotype
- 599 percentages and cell types were based on 200 cell counts in each repeat. Means with the same
- percentages and cen types were based on 200 cen counts in each repeat. Wearis with the same

letter are not significantly different from each other according to the results of a one-way

- analysis of variance and least significant difference post hoc test. a, b D. armatus JYCA037. c, d
- 602 D. armatus JYCA041. e, f D. armatus JYCA045.

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- Fig. 5 Mean number of cells per coenobium and proportions of unicells and two- and four-celled
- 605 coenobia in three strains of Desmodesmus armatus cultured at different 2,4-
- dichlorophenoxyacetic acid (2,4-D) concentrations. Data are presented as means (n = 3) for each



- group, and morphotype percentages and cell types were based on 200 cell counts in each repeat.
- Means with the same letter are not significantly different from each other according to a one-way
- analysis of variance and least significant difference post hoc test. a, b D. armatus JYCA037. c, d
- 610 D. armatus JYCA041. e, f D. armatus JYCA045.

- Fig. 6 Mean number of cells per coenobium and proportions of unicells and two- and four-celled
- 613 coenobia in three strains of Desmodesmus armatus cultured at different naphthalene-1-acetic
- acid (NAA) concentrations. Data are presented as means (n = 3) for each group, and morphotype
- percentages and cell types were based on 200 cell counts in each repeat. Means with the same
- letter are not significantly different from each other according to a one-way analysis of variance
- and least significant difference post hoc test. a, b D. armatus JYCA037. c, d D. armatus
- 618 JYCA041. **e**, **f** *D*. *armatus* JYCA045.

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- 620 Fig. 7 Mean number of cells per coenobium and proportions of unicells and two- and four-celled
- 621 coenobia of Desmodesmus armatus JYCA039 cultured at different indole-3-acetic acid (IAA),
- 622 2,4-dichlorophenoxyacetic acid (2,4-D), and naphthalene-1-acetic acid (NAA) concentrations.
- Data are presented as means (n = 3) for each group, and morphotype percentages and cell types
- were based on 200 cell counts in each repeat. Means with the same letter are not significantly
- 625 different from each other according to a one-way analysis of variance and least significant
- 626 difference post hoc test.

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- 628 Fig. 8 Mean number of cells per coenobium and proportions of unicells and two- and four-celled
- 629 coenobia of Desmodesmus communis JYCA040 cultured at different indole-3-acetic acid (IAA),
- 630 2,4-dichlorophenoxyacetic acid (2,4-D), and naphthalene-1-acetic acid (NAA) concentrations.
- Data are presented as means (n = 3) for each group, and morphotype percentages and cell types
- were based on 200 cell counts in each repeat. Means with the same letter are not significantly
- 633 different from each other according to a one-way analysis of variance and least significant
- 634 difference post hoc test.

- **Fig. 9** Mean number of cells per coenobium and proportions of unicells and two- and four-celled
- 637 coenobia of *Desmodesmus opoliensis* JYCA043 cultured at different indole-3-acetic acid (IAA),
- 638 2,4-dichlorophenoxyacetic acid (2,4-D), and naphthalene-1-acetic acid (NAA) concentrations.
- Data are presented as means (n = 3) for each group, and morphotype percentages and cell types
- 640 were based on 200 cell counts in each repeat. Means with the same letter are not significantly
- 641 different from each other according to a one-way analysis of variance and least significant
- 642 difference post hoc test.



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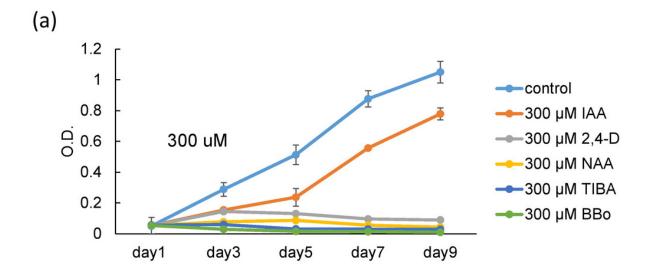
 Fig. 10 Mean number of cells per coenobium and proportions of unicells and two- and four-celled coenobia of *Desmodesmus communis* JYCA044 cultured at different indole-3-acetic acid (IAA), 2,4-dichlorophenoxyacetic acid (2,4-D), and naphthalene-1-acetic acid (NAA) concentrations. Data are presented as means (n = 3) for each group, and morphotype percentages and cell types were based on 200 cell counts in each repeat. Means with the same letter are not significantly different from each other according to a one-way analysis of variance and least significant difference post hoc test.

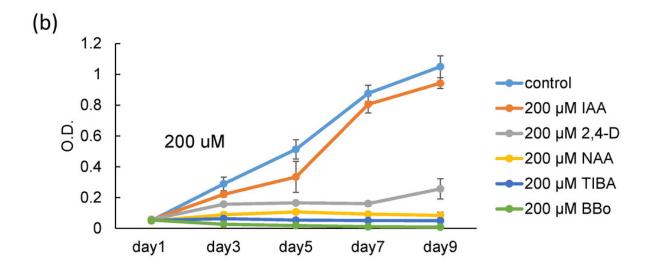
Fig. 11 Mean number of cells per coenobium and proportions of unicells and two-, and four-celled coenobia of *Desmodesmus intermedius* strain JYCA042 cultured at different indole-3-acetic acid (IAA), 2,4-dichlorophenoxyacetic acid (2,4-D), and naphthalene-1-acetic acid (NAA) concentrations. Data are presented as means (n = 3) for each group, and morphotype percentages and cell types were based on 200 cell counts in each repeat. Means with the same letter are not significantly different from each other according to a one-way analysis of variance and least significant difference post hoc test.

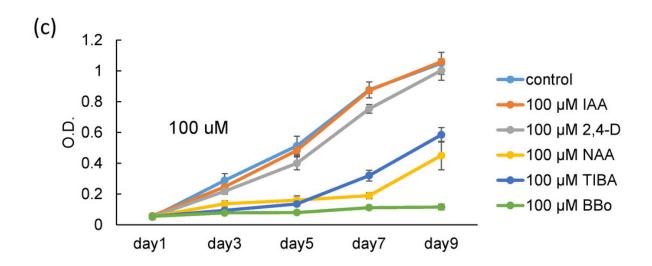


Growth of coenobial algae *Desmodesmus komarekii* in the presence of several auxins, inhibitor of auxin biosynthesis and auxin transport.

Growth of coenobial algae *Desmodesmus komarekii* in the presence of several auxins, inhibitor of auxin biosynthesis and auxin transport. *D. komarekii* was cultured in the presence of **a** 300, **b** 200, and **c** 100 µM indole-3-acetic acid (IAA), 2,4-dichlorophenoxyacetic acid (2,4-D), naphthalene-1-acetic acid (NAA), 2,3,5-triiodobenzoic acid (TIBA), or 4-biphenylboronic acid (BBo). Growth curves of *D. komarekii* for each compound were measured at 1, 3, 5, 7 and 9 days. Error bars represent standard deviation of values for three replicates.



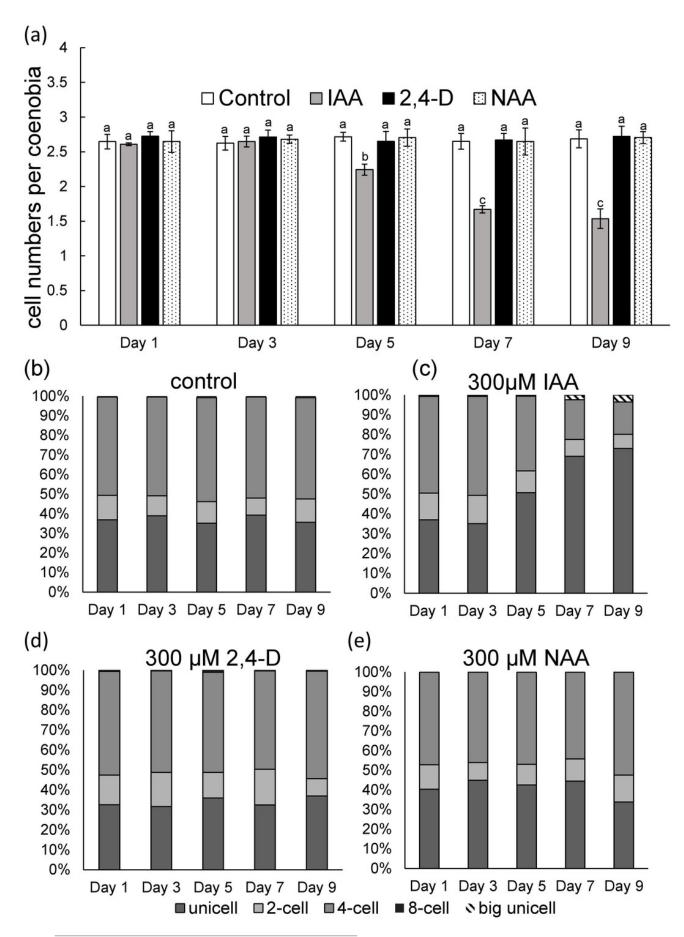






Mean number of cells per coenobium and proportions of unicells and of two- and four-celled coenobia of *Desmodesmus komarekii* cultured at 300 μ M IAA, 2,4-D, and NAA concentrations and cells without treatment.

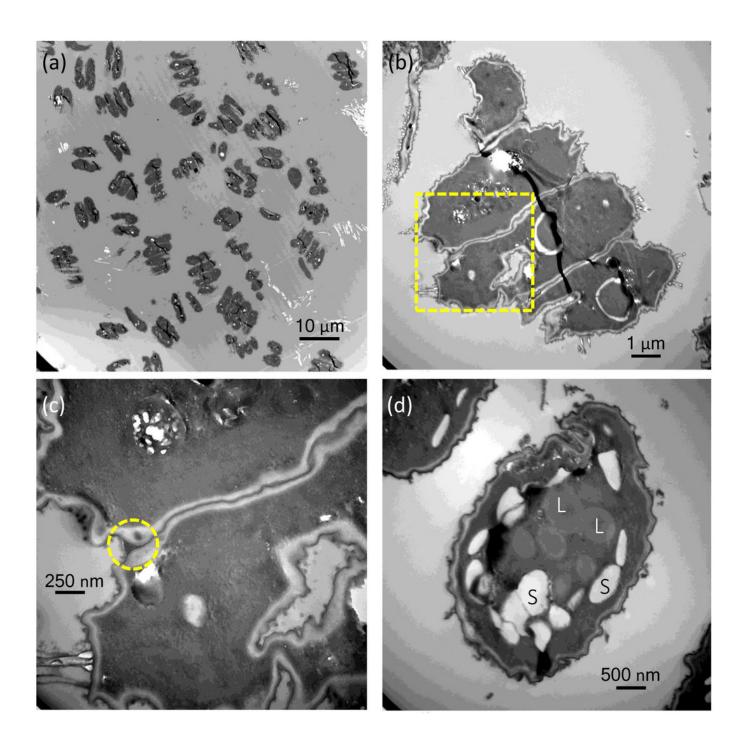
Mean number of cells per coenobium and proportions of unicells and of two- and four-celled coenobia of *Desmodesmus komarekii* cultured at 300 μ M indole-3-acetic acid (IAA), 2,4-dichlorophenoxyacetic acid (2,4-D), and naphthalene-1-acetic acid (NAA) concentrations and cells without treatment. Data are presented as means (n = 3) for each group, and morphotype percentages and cell types were based on 200 cell counts in each repeat. Means with the same letter are not significantly different from each other according to a one-way analysis of variance and least significant difference post hoc test.





Transmission electron micrographs of *Desmodesmus komarekii* cells under indole-3-acetic acid (IAA) treatment.

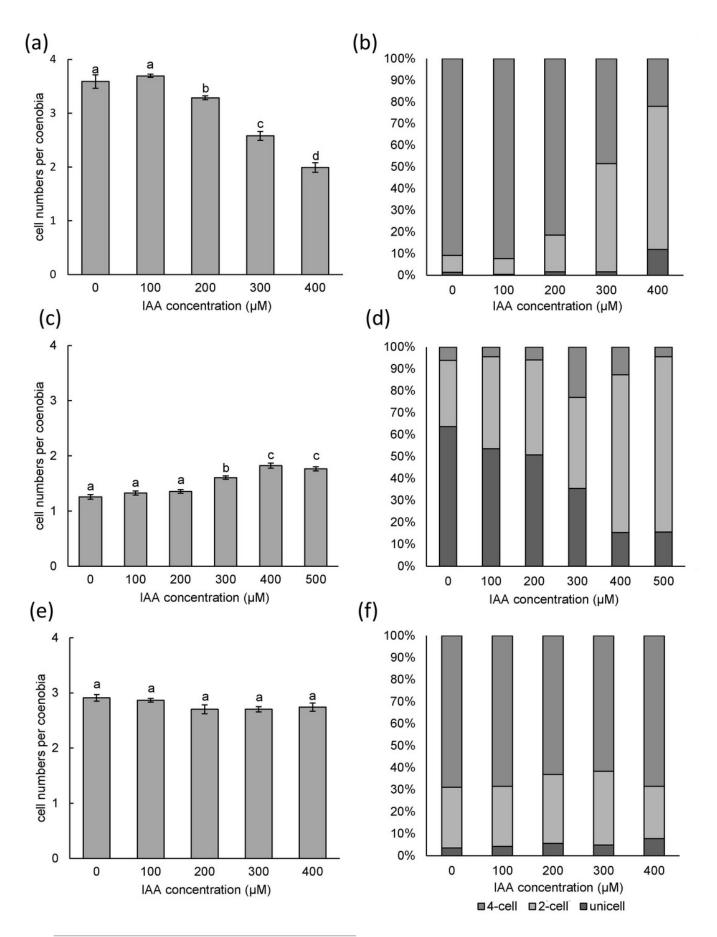
Transmission electron micrographs of *Desmodesmus komarekii* cells under indole-3-acetic acid (IAA) treatment. **a** Through transmission electron microscopy, we confirmed that the morphological changes in coenobia were not caused by cell aggregation but by the vegetative growth of a mother cell. **b** No extracellular matrix was seen on or around the cells, and the connecting strands between cells were highly visible. **c** Dense section of the warty layer can be seen over each coenobial junction (yellow circle). **d** The accumulation of many starch granules (S) and lipid bodies (L) was observed in the large unicells at day 7 after IAA treatment compared with the cells in control groups.





Mean number of cells per coenobium and proportions of unicells and two- and fourcelled coenobia in three strains of *Desmodesmus armatus* cultured at different IAA concentrations.

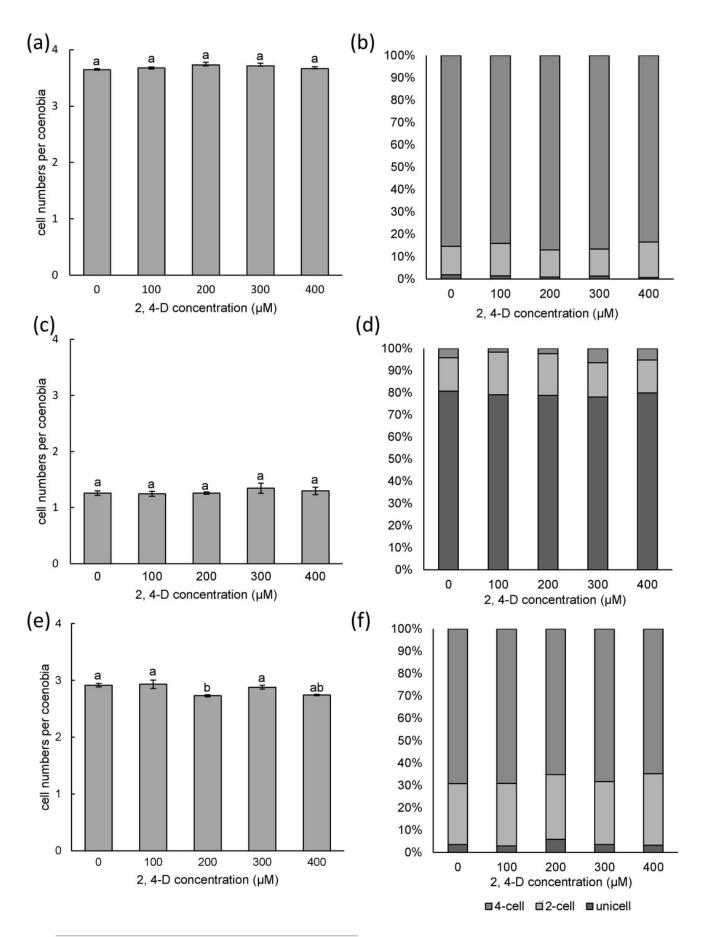
Mean number of cells per coenobium and proportions of unicells and two- and four-celled coenobia in three strains of *Desmodesmus armatus* cultured at different indole-3-acetic acid (IAA) concentrations. Data are presented as means (n=3) for each group, and morphotype percentages and cell types were based on 200 cell counts in each repeat. Means with the same letter are not significantly different from each other according to the results of a one-way analysis of variance and least significant difference post hoc test. **a**, **b** *D*. armatus JYCA037. **c**, **d** *D*. armatus JYCA041. **e**, **f** *D*. armatus JYCA045.





Mean number of cells per coenobium and proportions of unicells and two- and fourcelled coenobia in three strains of *Desmodesmus armatus* cultured at different 2,4-D concentrations.

Mean number of cells per coenobium and proportions of unicells and two- and four-celled coenobia in three strains of *Desmodesmus armatus* cultured at different 2,4-dichlorophenoxyacetic acid (2,4-D) concentrations. Data are presented as means (n = 3) for each group, and morphotype percentages and cell types were based on 200 cell counts in each repeat. Means with the same letter are not significantly different from each other according to a one-way analysis of variance and least significant difference post hoc test. **a**, **b** *D. armatus* JYCA037. **c**, **d** *D. armatus* JYCA041. **e**, **f** *D. armatus* JYCA045.

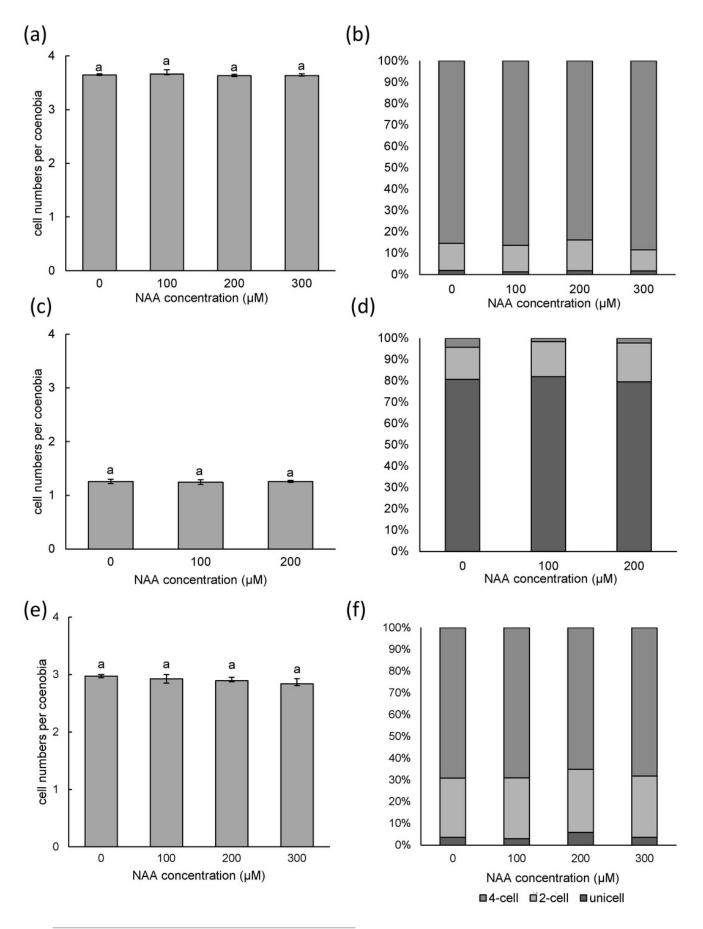




Mean number of cells per coenobium and proportions of unicells and two- and fourcelled coenobia in three strains of *Desmodesmus armatus* cultured at different NAA concentrations.

Mean number of cells per coenobium and proportions of unicells and two- and four-celled coenobia in three strains of *Desmodesmus armatus* cultured at different naphthalene-1-acetic acid (NAA) concentrations. Data are presented as means (n = 3) for each group, and morphotype percentages and cell types were based on 200 cell counts in each repeat. Means with the same letter are not significantly different from each other according to a one-way analysis of variance and least significant difference post hoc test. **a**, **b** *D*. *armatus* JYCA037. **c**, **d** *D*. *armatus* JYCA041. **e**, **f** *D*. *armatus* JYCA045.



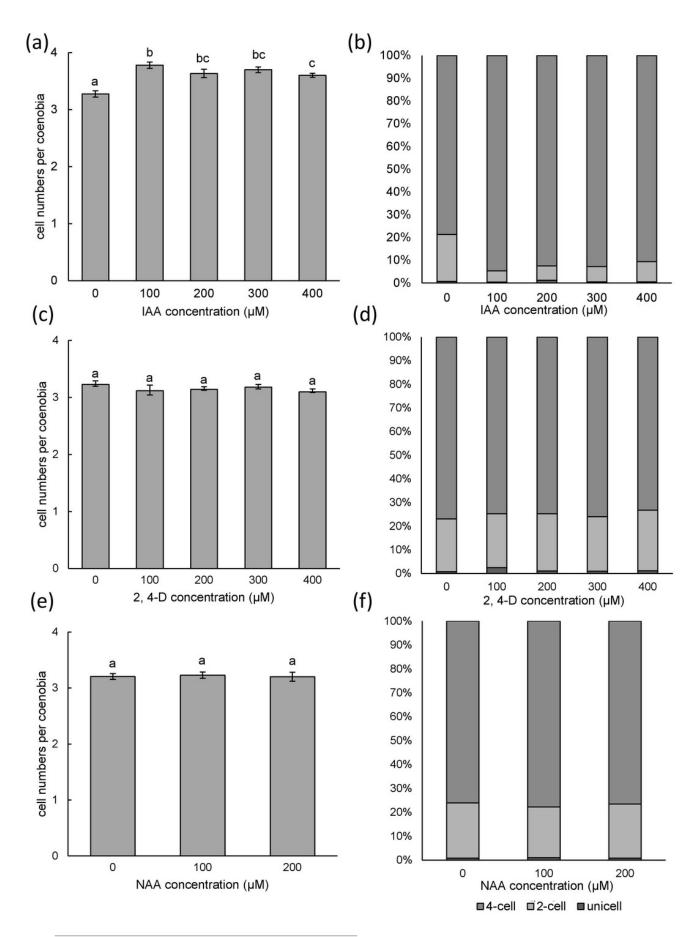




Mean number of cells per coenobium and proportions of unicells and two- and four-celled coenobia of *Desmodesmus armatus* JYCA039 cultured at different IAA, 2,4-D, and NAA concentrations.

Mean number of cells per coenobium and proportions of unicells and two- and four-celled coenobia of *Desmodesmus armatus* JYCA039 cultured at different indole-3-acetic acid (IAA), 2,4-dichlorophenoxyacetic acid (2,4-D), and naphthalene-1-acetic acid (NAA) concentrations. Data are presented as means (n=3) for each group, and morphotype percentages and cell types were based on 200 cell counts in each repeat. Means with the same letter are not significantly different from each other according to a one-way analysis of variance and least significant difference post hoc test.



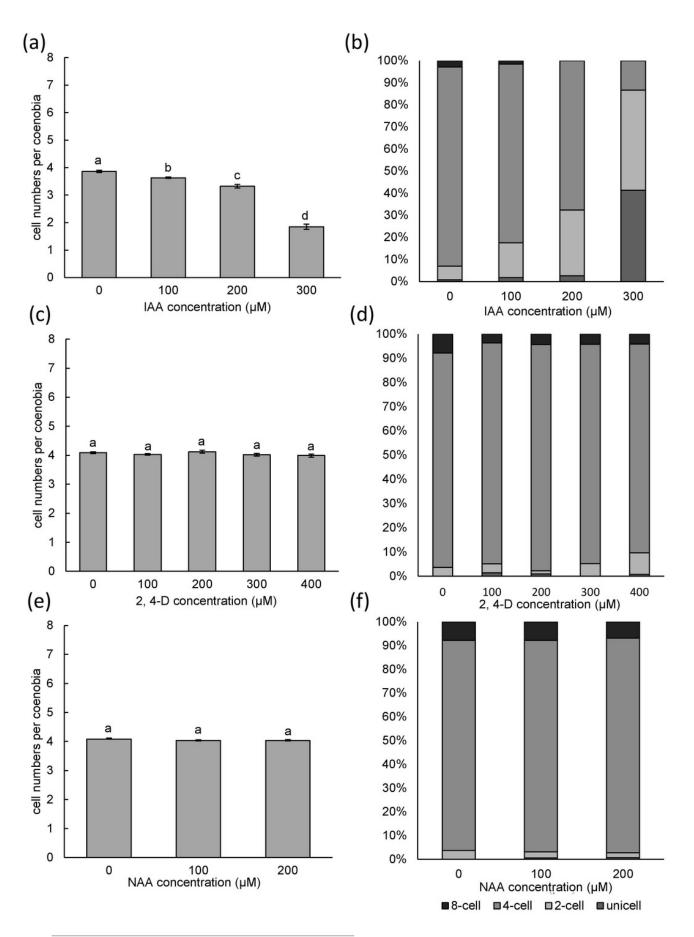




Mean number of cells per coenobium and proportions of unicells and two- and fourcelled coenobia of *Desmodesmus communis* JYCA040 cultured at different IAA, 2,4-D, and NAA concentrations.

Mean number of cells per coenobium and proportions of unicells and two- and four-celled coenobia of *Desmodesmus communis* JYCA040 cultured at different indole-3-acetic acid (IAA), 2,4-dichlorophenoxyacetic acid (2,4-D), and naphthalene-1-acetic acid (NAA) concentrations. Data are presented as means (n=3) for each group, and morphotype percentages and cell types were based on 200 cell counts in each repeat. Means with the same letter are not significantly different from each other according to a one-way analysis of variance and least significant difference post hoc test.

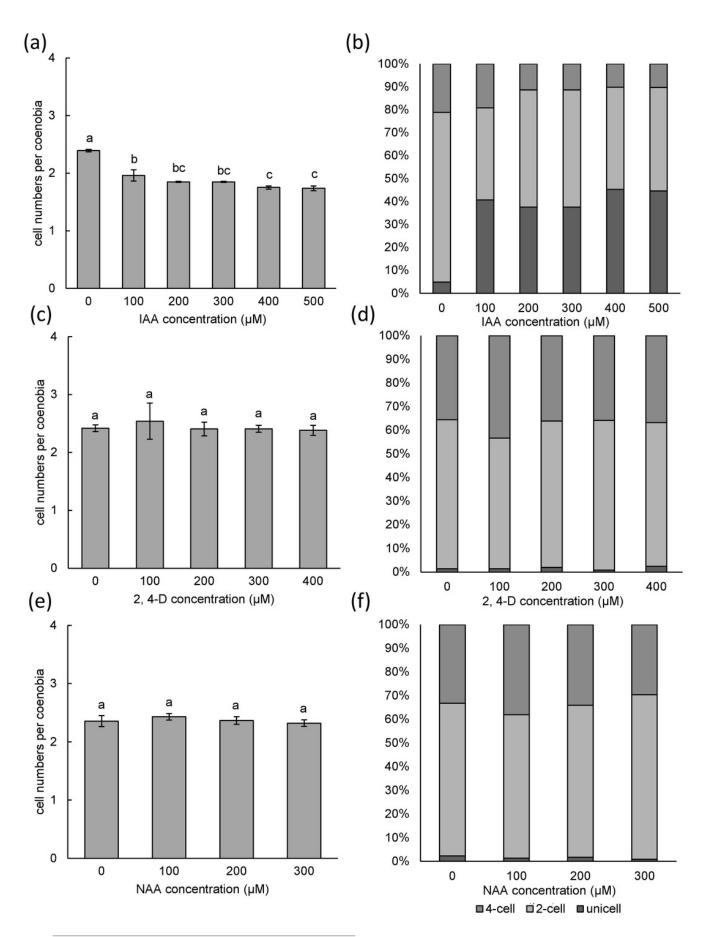






Mean number of cells per coenobium and proportions of unicells and two- and four-celled coenobia of *Desmodesmus opoliensis* JYCA043 cultured at different IAA, 2,4-D, and NAA concentrations.

Mean number of cells per coenobium and proportions of unicells and two- and four-celled coenobia of *Desmodesmus opoliensis* JYCA043 cultured at different indole-3-acetic acid (IAA), 2,4-dichlorophenoxyacetic acid (2,4-D), and naphthalene-1-acetic acid (NAA) concentrations. Data are presented as means (n=3) for each group, and morphotype percentages and cell types were based on 200 cell counts in each repeat. Means with the same letter are not significantly different from each other according to a one-way analysis of variance and least significant difference post hoc test.

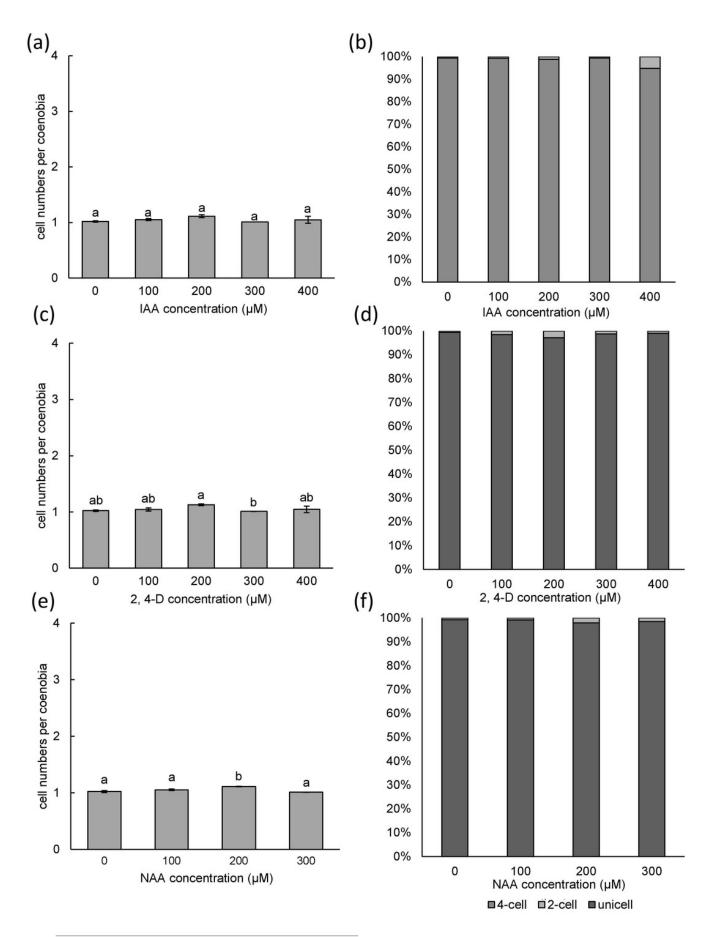




Mean number of cells per coenobium and proportions of unicells and two- and four-celled coenobia of *Desmodesmus communis* JYCA044 cultured at different IAA, 2,4-D, and NAA concentrations.

Mean number of cells per coenobium and proportions of unicells and two- and four-celled coenobia of *Desmodesmus communis* JYCA044 cultured at different indole-3-acetic acid (IAA), 2,4-dichlorophenoxyacetic acid (2,4-D), and naphthalene-1-acetic acid (NAA) concentrations. Data are presented as means (n=3) for each group, and morphotype percentages and cell types were based on 200 cell counts in each repeat. Means with the same letter are not significantly different from each other according to a one-way analysis of variance and least significant difference post hoc test.







Mean number of cells per coenobium and proportions of unicells and two-, and four-celled coenobia of *Desmodesmus intermedius* strain JYCA042 cultured at different IAA, 2,4-D, and NAA concentrations.

Mean number of cells per coenobium and proportions of unicells and two-, and four-celled coenobia of *Desmodesmus intermedius* strain JYCA042 cultured at different indole-3-acetic acid (IAA), 2,4-dichlorophenoxyacetic acid (2,4-D), and naphthalene-1-acetic acid (NAA) concentrations. Data are presented as means (n=3) for each group, and morphotype percentages and cell types were based on 200 cell counts in each repeat. Means with the same letter are not significantly different from each other according to a one-way analysis of variance and least significant difference post hoc test.

