

Grazer cues induce stealth behavior in marine dinoflagellates

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Chain formation is common among phytoplankton organisms but the underlying reasons and consequences are poorly understood. Here we show that chain formation is strongly impaired by waterborne cues from copepod grazers in the dinoflagellate *Alexandrium tamarens*. Chains of *Alexandrium* cells exposed to copepod cues responded by splitting into single cells or shorter chains. Motion analysis revealed significantly lower swimming velocities for single cells compared with chains, with two- to fivefold higher simulated predator encounter rates for two- and four-cell chains, respectively. In addition, the few remaining two-cell chains in grazed treatments were swimming at approximately half the speed of two-cell chains in treatments without grazers, which reduced encounter rates with grazers to values similar to that of single cells. Chain length plasticity and swimming behavior constitute unique mechanisms to reduce encounters with grazers. We argue that dinoflagellates can regulate the balance between motility and predator avoidance by adjusting chain length. The high predator encounter rate for motile chains may have contributed to the low prevalence of chain formation in motile phytoplankton compared with in nonmotile phytoplankton where chain formation is more common.

inducible defense | chemical ecology | hydrodynamics | plankton ecology

Oceanic primary producers contribute ~50% of the global carbon dioxide fixation and have profound effects on biogeochemical cycles (1), yet our understanding of their functional morphology is still rudimentary. As an example, chain formation is common mainly among nonmotile groups of marine phytoplankton, e.g., diatoms and cyanobacteria, but less common in motile groups like dinoflagellates. The ultimate reasons for chain formation are poorly understood and several alternative explanations have been proposed (2). Chains and colonies have been suggested to provide lower sinking rates, allowing nonmotile phytoplankters to remain in surface waters, although this result has little theoretical and experimental support (3). Size changes dramatically with chain and colony formation, allowing both motile and nonmotile phytoplankton organisms to enter size-limited grazer refuges. For example, a large *Phaeocystis* colony has a diameter of $>10^2$ times the diameter of a single cell. With the exception of parasites and pathogens, pelagic consumers are typically not able to feed on such a large size range of prey (4). Thus, it is likely that size selective grazing contributed to the evolution of size and colony formation in phytoplankton organisms (5). This relationship is further supported by the ability of *Phaeocystis globosa* and *Scenedesmus subspicatus* to sense and respond to grazer presence by forming colonies larger than the capture size of the inducing grazer (6, 7) or by breaking up colonies into sizes too small to be retained (8). Moreover, chain length correlates to growth rate in some diatoms and dinoflagellates (9, 10), suggesting that chain length may also depend on growth conditions. Finally, chains of dinoflagellates typically swim 40–60% faster than single cells, which has been suggested to enhance their ability to migrate vertically to collect nutrients at depth and harvest light at the surface (11) and to maintain depth in the face of turbulence (12–14).

Increased size and swimming velocity associated with chain formation, however, also leads to higher encounter rates with predators, and organisms must balance resource acquisition with the risk of predation (15). The encounter rate between a swimming dinoflagellate and an ambush-feeding predator scales with the swimming velocity and the square of the distance at which the cells can be perceived and attacked (16). Thus, chain formation is likely to result in an increased encounter rate with grazers, both due to the higher swimming velocity and due to an increased detection distance caused by increased hydrodynamic signal of larger and faster units (17).

Here we demonstrate in incubation experiments and through measurements of swimming velocities that a chain-forming motile dinoflagellate, *Alexandrium tamarens*, reduces its chain length and swimming velocity in response to waterborne grazer cues and that this response implies a substantially reduced risk of being eaten by a zooplankton grazer.

Results

Effect of Grazers on Chain Length. *A. tamarens* exposed to copepod grazers responded by splitting up chains into single cells and had a significantly lower proportion of cells in chains ($P < 0.05$) relative to that of control cultures (Fig. 1). Four-cell chains were common in the ungrazed treatment but observed only once in the grazed treatment. The concentration of cells (solitary or in chains) was reduced by $<15\%$ in the grazed treatment, showing that the reduction in chain length was not the result of grazing. This result was confirmed by a similar response in chain length in experiments where phytoplankton was exposed to copepods placed in plankton mesh (15- μm) cages (Fig. 2A) and demonstrates that the response was triggered by waterborne cues from the copepod grazers and not from physical contact with grazers. *Alexandrium* cultures exposed to copepod densities ranging from zero to eight copepods per liter revealed a density-dependent response with the highest proportion of single cells observed together with the highest number of grazers (Fig. 2B), corresponding to the upper range of naturally occurring copepod densities.

Motion analysis revealed higher swimming speed for chains in control treatments. Four-cell chains swam close to twice as fast as single cells and 33% faster than two-cell chains. Further, in grazed treatments, two-cell chains swam significantly slower compared with two-cell chains in control treatments ($P < 0.05$) and even slower than single cells in control containers ($P < 0.05$, Fig. 3 and Table 1), whereas single cells were swimming at a similar speed compared with control single cells ($P = 0.10$).

The directional persistence of swimming cells, quantified by the decorrelation timescale (τ), was comparable between treat-

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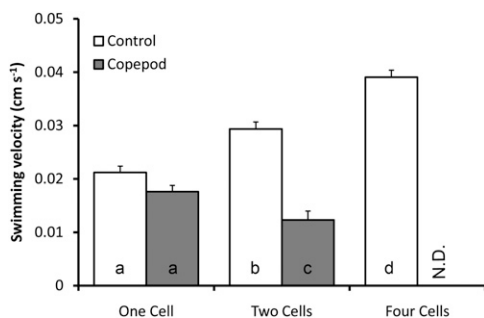


Fig. 3. Swimming velocity of single cells and two- and four-cell chains in grazed and control treatments. Only one single four-cell chain was observed in the grazed treatment, and it is not included in the analysis (shown as N.D. in the graph). Bars show mean values + SE of mean based on the number of observations (Table 1). Letters indicate statistically homogenous subsets according to the Student-Newman-Keuls post hoc procedure ($P < 0.05$).

control treatment because their larger size is compensated by their lower swimming velocity (Fig. 4). Two-cell chains were, however, scarce in the grazed treatment and it is possible that they represent recently divided cells that had not yet had time to regain speed after division or to part into single cells, rather than a behavioral response to copepod cues.

If we instead consider a moving predator, dinoflagellate velocity will become less important as predator velocity increase and the difference in encounter rate will approach the difference solely driven by the longer reaction distances (R) to chains, i.e., 2.2 and 4.6 times higher for two- and four-cell chains, respectively (Eq. 1 and Fig. 4B). This reasoning holds for rheotactic predators detecting prey by hydrodynamical signals. For grazers using chemical prey detection, such as many copepods (24), the reaction distance will also increase with chain length, simply because more cells leak more chemical signals. For moving and non-moving cells the enhancement is proportional to the leakage rate and, hence to the number of cells in the chain (25, 26). For grazers with a feeding current the effect may be of similar magnitude, although the exact effect is difficult to predict. Volumetric encounter rates may thus increase by a factor of up to 4^2 in four-cell chains compared with single cells (Eq. 1). The density dependence experiment (Fig. 2B) suggests that the response is well tuned to the naturally occurring densities of copepod grazers and that the proportion of single cells depends directly on the density of copepods grazers. It is possible that the reactions would have been more pronounced if the dinoflagellates were introduced to water already tainted by copepods because copepod cues would not have to build up during the first part of the experiments. If copepods are removed, the dinoflagellates need time to undergo one to two divisions to recover chain length as chains are formed by daughter cells staying attached after division.

The observed swimming velocities for single cells and chains in the control treatment are in good agreement with previous observations of swimming behavior in chain-forming dinoflagellates.

Fraga and coworkers (12) reported a 50–60% increase in swimming velocity for four-cell chains compared with single cells for *Alexandrium affine* and *Gymnodinium catenatum*, slightly less than the 84% observed here. Lewis and coworkers (13) found 45% higher swimming velocity for two-cell chains of *A. tamarense* compared with single cells, similar to the 38% observed in the present study.

Whereas the above considerations suggest that there are fitness benefits associated with reduced swimming velocity (lower predation risk), one may ask what the costs and benefits of elevated swimming speeds are in predator-free environments. The energetic cost of swimming at low Reynolds numbers is generally low (27) and the higher swimming velocity of chains may result mainly from the changed ratio between propulsion and drag forces (12). The difference in velocity between single cells and chains is actually less than predicted from propulsion to drag force calculations (12), suggesting that the higher velocity of chains comes for free or even at a reduced cost. Sperms from the wood mouse similarly gain higher path velocity by forming cooperative “trains” (28). Cells in chains may, however, experience increased competition from adjacent cells when nutrients are limiting. Assuming self-propelled spherical cells, using observed swimming velocities and cell chain sizes, and applying the numerical advection-diffusion model of Langlois et al. (26), we find that the specific transport-limited nutrient uptake may decrease by 13% in two-cell chains and by 45% in four-cell chains relative to solitary cells. The reduced nutrient uptake of cells in chains is relaxed because chains are elongated, whereas solitary cells are near spherical. In a patchy environment, however, most of the nutrient uptake may occur while cells are inside nutrient patches and nutrient uptake is not transport limited. *A. tamarense* cells perform nocturnal vertical migrations at speeds of up to $2 \text{ m}\cdot\text{h}^{-1}$ and an amplitude of $\sim 10 \text{ m}$ to retrieve nutrients from depth (29), a strategy that appears critical to the survival of many dinoflagellates in nutrient-limited environments (11). On the basis of the swimming speeds from the current experiment, single cells would not be able to complete such migrations in 24 h, whereas chains would, provided that they are able to maintain directional swimming during migrations. The benefits of vertical migration have been calculated to greatly exceed the costs (30), suggesting that there is a trade-off between nutrient acquisition and grazer avoidance in chain-forming dinoflagellates and that the dinoflagellates can use grazer-induced chain-length plasticity to adjust the balance between the two. Experimental evidence that chains actually benefit from their higher velocity is, however, still lacking.

Grazing pressure on phytoplankton is extremely high (31) and large motile cells have to compensate for their motility-dependent higher encounter rates with grazers. They can do so either by higher growth rates, which is typically not the case for large dinoflagellates (32), or by avoiding grazers by other means. Escape responses is one such mechanism that occurs among motile dinoflagellates (33). The production of poisonous secondary metabolites appears particularly common among large chain-

Table 1. Motion analysis results from single cells and chains in control and grazed treatments

Treatment	Length, cells	ESD, μm	Velocity, $\mu\text{m}\cdot\text{s}^{-1}$, mean \pm SE	Observations, tracks	Decorrelation time, τ , s, mean \pm SE
Control	1	28	212 ± 12	23	1.3 ± 0.01
	2	34	294 ± 13	23	1.1 ± 0.02
	4	44	391 ± 13	24	2.0 ± 0.04
Grazed	1	28	176 ± 12	23	1.2 ± 0.02
	2	34	123 ± 17	7	1.7 ± 0.03
	4	44	—	—	—

Four-cell chains were observed only once in the grazed treatment and are not included.

The swimming behavior of dinoflagellates was analyzed using a MatLab algorithm that returned the positions, velocity, and net and gross displacement over time for individual cells or chains every 0.12 or 0.24 s. Trajectories averaged 14 ± 6.5 s (mean \pm SD) in length. Convection was slow compared with the swimming velocity of *Alexandrium* cells, and tracks of single cells and chains from each treatment were obtained from the same video sequences, assuming that any contribution of convection would cancel out in the comparison between chains and single cells. Four-cell chains were observed only once in the grazed treatment recording and therefore not included in motion analysis. 3D swimming speeds were calculated by multiplying by $\sqrt{3}/\sqrt{2}$, thus assuming that the cells on average were swimming isotropically in all directions. Highly convoluted swimming patterns may reduce encounter rates significantly (43). To control for any major changes in the rate of change of direction we estimated the decorrelation timescale (τ) as a measure of directional persistence from curve fits to Taylor's formula for diffusion by continuous movement (44) to plots of root mean square displacement over time (t):

$$\text{Root mean square displacement (RMS)} = \left\{ 2v^2\tau \left[t - \tau(1 - e^{-(t/\tau)}) \right] \right\}^{0.5}. \quad [3]$$

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Statistical Analysis. The proportion of cells present in chains in grazed and nongrazed treatments for motion analysis was compared with a χ^2 -test using the control distribution as the expected frequency. The percentage of cells present in chains in the cage experiment was compared with Student's *t* test for two samples assuming equal variances. A single-factor ANOVA was used to evaluate the effect of grazer density. The effect of chain formation and grazer presence on swimming velocity was compared using a two-factor ANOVA, with the factors "grazer presence" (two levels: present and absent) and chain formation (three levels: single cells, two-cell chains, and four-cell chains) as fixed factors. The average velocity from each trajectory was used in the analysis. The Student–Newman–Keuls post hoc procedure was used to identify statistically homogenous subsets when ANOVA indicated significant effects.

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