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Toxic dinoflagellates produce true grazer deterrents

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Abstract. Many phytoplankton species produce toxic substances, but their functional role is unclear. Specifically, it remains uncertain whether these compounds have a toxic or deterrent effect on grazers; only, the latter is consistent with toxins as defensive tools. Here, we show that 10 of 12 species or strains of toxic dinoflagellates were consumed at lower rates than a similarly sized nontoxic dinoflagellate by a copepod. Through video observations of individual prey–grazer interactions, we further demonstrate that the dominating mechanism is through capture, examination, and subsequent rejection of vital cells, that is, a true deterrent effect that offers a straightforward explanation to its evolution. We argue that the diversity of grazer responses to toxic phytoplankton reported in the literature, including toxic effects, and the high diversity of toxin profiles between strains of the same phytoplankton species reflect different stages of an ever-ongoing evolutionary arms race, facilitated by rapid adaptation of grazers to toxic substances. We further argue that defensive toxicity requires a chemical signal exterior to the cell that informs the grazer about the toxicity of the cell. The signal can be the toxin itself or just an aposematic signal of toxicity. In the former case, allelochemical effects may emerge at high cell concentrations as a nonadaptive side effect of a predator defenses.

Key words: coevolution; feeding behavior; grazer deterrent; prey rejection; Temora longicornis; toxic dinoflagellates.

INTRODUCTION

Many species of phytoplankton produce substances that negatively impact other organisms. These substances may be allelochemical compounds that are released into the environment and may affect competing phytoplankton, potential grazers, potential prey, and other organisms (Tillmann and John 2002, Tillmann 2003, Granéli and Hansen 2006, Blossom et al. 2012, John et al. 2015). Alternatively, such substances may be intracellular compounds that directly impact only organisms that consume the phytoplankton (see reviewed by Turner 2014, Uye and Takamatsu 1990, Xu et al. 2017). Chemical warfare and defense are well described and understood in terrestrial plants that are sedentary and can afford sacrificing a few leaves to deter grazers (Wittstock and Gershenzon 2002). Terrestrial models of chemical warfare and defense do not, however, necessarily apply well to phytoplankton. A toxic phytoplankton cell consumed by a grazer may benefit its genetically identical siblings in a population consisting of clonal cells (Thornton 2002), but phytoplankton populations are typically genetically diverse (Alpermann et al. 2010) and cheaters that do not produce the substance as well as competing species may be equally benefitted. Thus, cheaters may invade and this appears not to be an evolutionary stable strategy (ESS) (West et al. 2006, Lee et al. 2010). Only in toxic cells with grazer-deterrent effects can toxicity be considered a true defense. However, most incubation experiments have been unable to distinguish toxic from deterrent effects (Turner 2014). Among experiments that have been able to separate the two effects, only a few have demonstrated selection against toxic cells (Teegarden 1999, Selander et al. 2006), while many reports demonstrate toxic effects (Sykes and Huntley 1987, Lasley-Rasher et al. 2016, Rasmussen et al. 2017, Xu et al. 2017, 2018). Similarly, leaked substances with allelopathic effects on competitors and predators are “public goods” that benefit not only the producer but also nontoxic species and thus do not constitute an ESS (Lewis 1986, Driscoll et al. 2016). Understanding the evolution of the production of toxic substances in phytoplankton is incomplete, and toxin functional roles remain unclear. The fact that the toxic compounds may often negatively impact grazers (as reviewed by Turner 2014), and that the production of the substances may be induced or upregulated in the presence of grazers due to chemical signals (Selander et al. 2006, 2012, 2015) strongly suggests; however, that in many cases, these substances play a defensive role.

One key to further the understanding of the function and evolution of harmful substances as defense tools is to examine the impact at the individual level as this may allow one to distinguish toxic from deterrent effects. There have been only a handful of studies that have examined the effects of a few species of toxic algae on copepod grazers at the individual level. These studies have revealed a number of different responses: (1) aberrant swimming or appendage behavior in copepods subsequent to exposure to toxic algae (Sykes and Huntley 1987, Cohen et al. 2007, Hong et al. 2012, Lasley-Rasher et al. 2016, Xu et al. 2018), likely a consequence of the consumption of some cells and subsequent intoxication and reduced feeding rate; (2) cells are captured but subsequently regurgitated (Sykes and Huntley 1987, Xu et al. 2017), resulting in mortality of ingested cells of toxic algae, initially similar to that of nontoxic cells, but with reduced benefits to the grazer and reduced grazing overall in the longer term; and (3) toxic cells are captured but subsequently rejected, leaving the toxic cell intact and vital (Xu et al. 2018).
et al. 2017, 2018). While the grazers’ behavioral responses in all cases lead to reduced grazing on toxic cells, only the third response is a true grazing deterrent response that yields the toxic cells with a clear individual advantage and that can be easily understood as a defense mechanism.

The above studies involved a relatively narrow selection of species and strains of dinoflagellates, and the different responses are not immediately comparable due to the use of different approaches. However, a genuine grazer-deterrent effect was demonstrated in only one case (Xu et al. 2017). Copepods are among the main grazers of dinoflagellates in the ocean, and most toxic species in the ocean are dinoflagellates (Turner 2014). To further explore the diversity of copepod behavioral responses to algae that produce toxic substance, we here examine, by a consistent approach, the effects of a diverse selection of 12 different species and strains of dinoflagellates on the foraging behavior of the copepod Temora longicornis. Through direct video observations, we demonstrate that the highly variable consumption rates observed between species and strains are accomplished by various mechanisms, but that the dominating response of the copepods to toxic algae is elevated rejection of captured cells, that is, a true grazer-deterrent effect that is immediately compatible with the role of toxic substances as a defense mechanism.

**Materials and Methods**

**Algal species and their toxin profiles**

The 13 dinoflagellate test species/strains varied somewhat in size, between 15 and 35 μm equivalent spherical diameter (Table 1), and they produce very different toxins. *Alexandrium tamarense* produces paralytic shellfish poisoning (PSP) toxins, a group of neurotoxins (Turner and Tester 1997, Cembella 1998, Turner 2014), and unknown lytic compounds (Ichimi et al. 2002, Tillmann et al. 2009). *Karenia brevis* can produce brevetoxins (PbTxs), a potent ichthyotoxic neurotoxins (Naar et al. 2007), brevetoxin antagonists (Baden et al. 2005, Satake et al. 2009), and hemolytic toxins (Neely and Campbell 2006). *Karenia mikimotoi* produces gymnocin-A, gymnocin-B, and some hemolytic toxins (Satake et al. 2002, 2005, Neely and Campbell 2006). *Karenia selliformis* is most closely related to *K. brevis* and *K. mikimotoi* and produces gymnodimines, brevetoxins, and hemolytic compounds (Seki et al. 1995, Miles et al. 2000, Tatters et al. 2010). The genus *Dinophysis* is known as the main source of okadaic acid (OA) and its derivatives, dinophysistoxins (DTX), while *Dinophysis acuminata* mainly produces pectenotoxin (PTX) rather than OA (Kamiyama and Suzuki 2009). In most cases, the effects of the various toxins on grazers are unknown. *Prototheca reticulata* served as a nontoxic control.

All the cultures were grown on B1 medium at 16°C, a salinity of 32, and an irradiance of 100–150 μmol photons·m⁻²·s⁻¹ on a 12 h light–dark cycle. The mixotrophic *D. acuminata* was additionally fed with the ciliate *Mesodinium rubrum*. We stopped feeding five days before experiments to have a monoculture of *D. acuminata* for the experiments. All algae for experiments were collected in the exponential growth phase.

Samples for toxin measurements of *A. tamarense*, *A. pseudogonyaulax*, *K. brevis*, *D. acuminata*, and *P. reticulata* were taken just prior to experiments, and the results shown in Table 2. Approximately 20 mL of *A. tamarense*, *A. pseudogonyaulax*, *K. brevis*, and *P. reticulata* cell culture (≥20,000 cells) was centrifuged at 2,150 × g for 15 min. Cell pellets were resuspended after removing most of the

**Table 1.** Phytoplankton used as prey for *Temora longicornis* including the strain number and isolation location.

<table>
<thead>
<tr>
<th>Algae</th>
<th>Strain</th>
<th>No.</th>
<th>Origin</th>
<th>ESD ± SD (μm)</th>
<th>Initial concentration (cells/mL)</th>
</tr>
</thead>
<tbody>
<tr>
<td><em>Alexandrium tamarense</em></td>
<td>Alex5</td>
<td>S1</td>
<td>North Sea of Scotland</td>
<td>33.8 ± 0.5</td>
<td>77</td>
</tr>
<tr>
<td></td>
<td>K-1471</td>
<td>S2</td>
<td>Offshore of Lysekil, Sweden</td>
<td>25.0 ± 2.3</td>
<td>192</td>
</tr>
<tr>
<td></td>
<td>CCMP 1598</td>
<td>S3</td>
<td>Daya Bay, China</td>
<td>30.0 ± 4.4</td>
<td>101</td>
</tr>
<tr>
<td><em>Alexandrium pseudogonyaulax</em></td>
<td>K-1344</td>
<td>S4</td>
<td>Limfjorden, Denmark</td>
<td>35.0 ± 5.4</td>
<td>67</td>
</tr>
<tr>
<td><em>Karenia armiger</em></td>
<td>K-1345</td>
<td>S5</td>
<td>Limfjorden, Denmark</td>
<td>31.8 ± 4.8</td>
<td>88</td>
</tr>
<tr>
<td><em>Karenia selliformis</em></td>
<td>K-0668</td>
<td>S6</td>
<td>Alfacs Bay, Spain</td>
<td>15.3 ± 3.1</td>
<td>595</td>
</tr>
<tr>
<td><em>Karenia brevis</em></td>
<td>K-1319</td>
<td>S7</td>
<td>Gulf of Gabes, Tunisia</td>
<td>20.8 ± 4.2</td>
<td>349</td>
</tr>
<tr>
<td></td>
<td>CCMP 2229</td>
<td>S8</td>
<td>Offshore of Manasota Key, USA</td>
<td>23.0 ± 5.2</td>
<td>286</td>
</tr>
<tr>
<td></td>
<td>CCMP 718</td>
<td>S9</td>
<td>Florida, USA</td>
<td>25.5 ± 4.2</td>
<td>168</td>
</tr>
<tr>
<td><em>Karenia mikimotoi</em></td>
<td>CCMP 429</td>
<td>S10</td>
<td>Sutton Harbour, United Kingdom</td>
<td>26.4 ± 5.1</td>
<td>195</td>
</tr>
<tr>
<td></td>
<td>K-0260</td>
<td>S11</td>
<td>Oslofjorden, Norway</td>
<td>23.3 ± 5.1</td>
<td>245</td>
</tr>
<tr>
<td><em>Dinophysis acuminata</em></td>
<td>S12</td>
<td></td>
<td>Denmark†</td>
<td>33.5 ± 11.3</td>
<td>89</td>
</tr>
<tr>
<td><em>Prototheca reticulata</em></td>
<td>CCMP 1889</td>
<td>S13</td>
<td>Friday Harbor, USA</td>
<td>32.0 ± 2.3</td>
<td>84</td>
</tr>
</tbody>
</table>

**Notes:** Equivalent spherical diameters (ESD; mean ± SD) are based on measurements of linear dimensions 20 cells of each species and by assuming spherical or spheroid shapes (Hillebrand et al. 1999). The initial cell concentration is that used in the video observation. Cell cultures were obtained from Alfred Wegener Institute, Germany; National Center for Marine Algae and Microbiota; Scandinavian Culture Collection for Algae and Protozoa; and Marine Biological Section, University of Copenhagen.

†Cultures were originally isolated from seven different locations in Denmark (Nielsen et al. 2012).
Table 2. Toxin profiles and contents of algae.

<table>
<thead>
<tr>
<th>Algae</th>
<th>Strain</th>
<th>No.</th>
<th>PSP†</th>
<th>GDA</th>
<th>PbTx</th>
<th>OA</th>
<th>DTX</th>
<th>PTX2</th>
<th>YTX</th>
</tr>
</thead>
<tbody>
<tr>
<td>Alexandrium tamarense</td>
<td>Alex5</td>
<td>S1</td>
<td>44.1</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
</tr>
<tr>
<td></td>
<td>K-1471</td>
<td>S2</td>
<td>ND</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
</tr>
<tr>
<td></td>
<td>CCPM 1598</td>
<td>S3</td>
<td>ND</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
</tr>
<tr>
<td>Alexandrium pseudogonyaulax</td>
<td>K-1344</td>
<td>S4</td>
<td>-</td>
<td>15.6</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
</tr>
<tr>
<td></td>
<td>K-1345</td>
<td>S5</td>
<td>-</td>
<td>19.8</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
</tr>
<tr>
<td>Karlodinium armiger</td>
<td>K-0668</td>
<td>S6</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
</tr>
<tr>
<td>Karenia brevis</td>
<td>CCMP 2229</td>
<td>S8</td>
<td>-</td>
<td>ND</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
</tr>
<tr>
<td></td>
<td>CCMP 718</td>
<td>S9</td>
<td>-</td>
<td>ND</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
</tr>
<tr>
<td>Karenia mikimotoi</td>
<td>CCMP 429</td>
<td>S10</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
</tr>
<tr>
<td></td>
<td>K-0260</td>
<td>S11</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
</tr>
<tr>
<td>Dinophysis acuminata</td>
<td>S12</td>
<td>-</td>
<td>-</td>
<td>ND</td>
<td>ND</td>
<td>-</td>
<td>19.1</td>
<td>-</td>
<td>-</td>
</tr>
<tr>
<td>Protoceratium reticulatum</td>
<td>CCPM 1889</td>
<td>S13</td>
<td>ND</td>
<td>ND</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
</tr>
</tbody>
</table>

Notes: PSP, Paralytic shellfish poisoning toxins; GDA, Goniodomin A; PbTx, Brevetoxin; OA, Okadaic acid; DTX, Dinophysistoxin; PTX2, Pecteno toxin-2; YTX, Yessotoxin; -, means toxins were not measured; ND, means toxins were not detected.
†Values were calculated as saxitoxin equivalents (pg STXeq/cell), including saxitoxin (STX) and four derivatives (NEO, GTX2/3, GTX1/4, and C1/2).

Copepods

The feeding-current feeding copepod, Temora longicornis, was collected in Øresund, Denmark (56°04′ N; 12°38′ E), and kept in continuous cultures in darkness at 15°C, a salinity of 32, and fed with a mixed diet of Akashiwo sanguinea, Heterocapsa triqueta, Prorocentrum micans, Thalassiosira weissflogii, and Rhodomonas salina. Animals for experiments were of F3 + generation.

For each treatment, approximately 15 adult females were tethered to their dorsal surface with a short length of human hair by a small drop of cyanoacrylate glue. Tethered copepods were placed in filtered sea water and starved overnight at the same temperature and salinity as the cultures. Tethered copepods can live for many days, feeding, defecating, and producing eggs, apparently unaffected by the tether.

Experimental design and setup

Tethered copepods in good condition were attached to a plastic pipette by the other end of the hair and connected to a micromanipulator. Each individual was submerged in a 10 × 10 × 10 cm³ aquarium with 900 mL filtered sea water and a single prey species. The initial volume concentration of algae was the same in all experiments, approximately 1.4 × 10⁶ µm³/mL, resulting in different cell number concentrations (Table 1). The water was gently stirred to keep cells in suspension. The aquarium was placed between an infrared light source and the camera equipped with lenses to yield a field of view of approximately 2.5 × 1.6 mm². Feeding behaviors of copepods were recorded by a high-speed camera (Phantom V210; Vision Research, Wayne, New Jersey, USA) that was connected to both a computer for saving short high-speed sequences (2,200 frames/s, resolution: 720 × 567 pixels) and a tape recorder for saving of longer low-speed videos (25 frames/s, resolution: 1,280 × 800 pixels) and one to three high-speed sequences (2,200 frames/s, resolution: 720 × 576 pixels). The experiments were conducted in a dark, constant temperature room at 16°C.

The experiment started when prey cells were added to the aquarium. During the subsequent 4 h, 5-min low-speed video sequences were recorded every 30 min and one to three high-speed sequences of up to 2.5 s duration were recorded every two hours. After 24 h, three low-speed videos and two high-speed videos were recorded within 1 h. Each treatment had three replicates.

Capture, ingestion, and rejection events were enumerated from every five-minute low-speed video, as described earlier (Xu et al. 2017). The fraction of time that the copepod was beating its feeding appendages was calculated from the number of frames in which the appendages were beating during the entire five-minute period. From high-speed videos, the beating frequency of the feeding appendages was estimated and details of feeding behaviors were observed. Clearance rate was calculated as the capture rate divided by prey concentration. Images were analyzed with ImageJ (version 1.51 h; National Institutes of Health, Bethesda, Maryland, USA) and Phantom Cine Viewer (version 2.6; Vision Research).

Statistical differences over time and between prey species in clearance rate, ingestion rate (converted to biomass ingested), beating frequency, and fraction of time beating were tested with one-way or two-way ANOVAs followed by
Holm–Sidak post hoc test. The assumption of normality was tested. The difference in fraction of cells rejected after being captured between treatments was compared by the 95% Wilson score interval.

**RESULTS**

**Ingestion rate**

Upon adding phytoplankton to the copepods, their ingestion rates decreased rapidly during the first two hours of observation for some of the strains (Appendix S1: Fig. S1 and Appendix S2: Fig. S1) but subsequently did not vary with time (two-way ANOVA, \( P = 0.407 \)). In the following, we thus compare averages for the observations during the last 22 h. Ingestion rates varied by a factor of up to 20 between prey species (Fig. 1, two-way ANOVA, \( P < 0.001 \)). The nontoxic control prey, *P. reticulatum* (S13), was ingested at the highest rate and *A. pseudogonyaulax* strains (S4 and S5) and *K. brevis* strains (S8 and S9) (Table 2) at the lowest rates, with the other strains in between. Variation in ingestion rate between prey can be due to different mechanisms, that is, to differences in capture rate, prey rejection rate, and feeding activity, which we examine in the following.

**Prey capture and clearance rate**

Capture rate (normalized by cell concentration = clearance rate) increased with prey size (within the range of 15–35 \( \mu m \)) at the start of the experiment (first five minutes), but this dependency rapidly disappeared (Fig. 2a) and clearance rate was subsequently controlled mainly by other factors. The dependency of ingestion and rejection behaviors on prey size similarly disappeared (Fig. 2b,c).

Average clearance rate (after 2 h) varied by a factor of about 40 between prey species and strains (Fig. 3a, two-way ANOVA, \( P < 0.001 \)). Compared to the control prey, *P. reticulatum* (S13), copepods had significantly higher clearance rates on *K. mikimotoi* (K-0260, S11) and *A. tamarense* (CCMP 1598, S3) and significantly low clearance rates on *A. tamarense* (Alex5, S1), *K. brevis* (CCMP 2229, S8), and *K. brevis* (CCMP 718, S9) (one-way ANOVA with Holm–Sidak post hoc test, \( P < 0.05 \)). Clearance rates were in two cases similar between strains of the same species (*A. pseudogonyaulax* S4-5, *K. brevis* S8-9), and in other one case differed significantly and substantially between strains (*A. tamarense* S1-3) (one-way ANOVA with Holm–Sidak post hoc test, \( P > 0.05 \) and \( P < 0.05 \), respectively).

**Fraction of cells rejected**

Captured cells may be rejected leading to reduced consumption rate. The copepod captures the cell, examines it during a few ms, and subsequently rejects it (Video S1 and S2). High rejection rate was the most consistent mechanism leading to low consumption rate (Fig. 3b and Appendix S3: Fig. S1). Around 20% of captured nontoxic control cells were rejected by the copepods, but with one exception (*A. tamarense*, Alex5, S1), all other prey were rejected at a similar (*A. tamarense* K-1471, S2) or higher to much higher rates. Especially, noteworthy is the very high rejection rates of *A. pseudogonyaulax* strains (S4 and S5) and also of *K. mikimotoi* (K-0260, S11) and *D. acuminata* (S12). This is the main reason that these cells are consumed at a low rate (Fig. 1). The low rejection rate of *A. tamarense* (Alex5, S1) and the high rejection rate of *K. brevis* (CCMP 718, S9) are based on very few observations (\( n = 72 \) and 24, respectively) due to low feeding activity and, thus, probably not representative.

**Feeding appendage activity**

A reduction in frequency and time fraction that *T. longicornis* beats its appendages to produce a feeding current

![Fig. 1. Average biomass of algae ingested by Temora longicornis from 3rd to 24th h. Error bars showed standard errors (\( n = 9 \)).](image-url)
may similarly account for low consumption rate of prey cells. In most cases, the feeding appendages beat near 100% of the time at a frequency of about 30 Hz (Fig. 3c and Appendix S4: Fig. S1), but in a few cases, the activity was significantly reduced and/or variable over time (Fig. 3d and Fig. 4). Thus, upon introduction of A. tamarense (Alex5, S1) the copepod rapidly almost ceased beating its appendages, and when beating, it was at a significantly lower frequency (compared to control prey, two-way ANOVA with Holm–Sidak post hoc test, \( P < 0.05 \), Fig. 3c). When offered K. brevis strains (S8 and S9), the beating frequency was reduced, although not significantly (compared to control prey, two-way ANOVA with Holm–Sidak post hoc test, \( P > 0.05 \), respectively), and the fraction of time beating also rapidly dropped to very low levels, although subsequently increased again somewhat after a couple of hours (Fig. 4).

**DISCUSSION**

After initial acclimation to the diet where prey clearance rate was governed mainly by prey size, all toxic algae were consumed at a lower or much lower rate than the nontoxic control prey, but the mechanisms by which feeding rates were reduced differed between algal species and strains. Significantly low consumption rates were in particular recorded for one strain of A. tamarense (Alex 5, S1), the two strains of A. pseudogonyaulax (S4 + 5), and the two strains of K. brevis (S8 + 9) and D. acuminata (S12). In one case, A. tamarense (Alex 5, S1), the algae were consumed at a very low rate simply because the copepod ceased feeding. However, the most consistent responses to the toxic algae were high rejection rate, which were observed in 10 out of 12 species/strains, and in some cases combined with reduced feeding activity. This response suggests that whatever substances the algae are producing, they function as true grazer deterrents and constitute an evolutionary stable defense strategy. The dominance of deterrent effects is a novel discovery and different from most of the very many earlier reports of grazers becoming intoxicated or dying in response to exposure to toxic algae or reports from black box experiments that have been unable to distinguish between toxic and deterrent effects (Turner 2014). Indeed, there are only very few cases where grazer-deterrent effects have been convincingly demonstrated for a small selection of freshwater cyanobacteria (DeMott and Moxter 1991) and marine dinoflagellates (Teegarden 1999, Selander et al. 2006, Schultz and Kiørboe 2009).

In the case of the two K. brevis strains (S8 + 9), the low clearance rates (Fig. 3a) cannot be explained solely by high rejection rates and reduced feeding activity, and must
depend on a precapture mechanism that may also constitute a response to a genuine grazer deterrent. Below we discuss the various mechanisms that lead to reduced cell consumption rates.

Prey perception and handling

Copepods perceive and capture prey cells individually (Koehl and Strickler 1981), and this allows the possibility that cells can be selected against based on their chemical or mechanical properties. The fact that cells of very similar shape and size, and even different strains of the same species, were rejected at significantly different rates strongly suggests that gustation and "taste" rather than the mechanical properties of the cells provide the cues for selection. It is, however, unclear what that chemical cue is since intracellular substances that are toxic to the grazer cannot necessarily be detected from the outside of the cell. Also, variations in consumption rates are typically not clearly related to the measured toxicity profiles of the different phytoplankton strains compared, neither in copepods (Teegarden et al. 2008, Xu et al. 2017), other organisms (Tillmann and John 2002), nor in our experiments (Table 2 and Fig. 1). This suggests that the intracellular substances we measure may not necessarily be the substances to which the copepods and other grazers respond behaviorally and physiologically. It will be beneficial for the chemically defended prey to use exuded compounds to signal its toxicity to grazers, like “warning coloration” (aposematism), and in fact, such taste signals are required to account for our observations. Many dinoflagellates are known to release dissolved molecules, some of which with documented allelopathic effects but often poorly characterized chemically (Tillmann and John 2002, Tillmann et al. 2009), and such substances may constitute signal molecules that are detected by grazers. Whether or not these molecules are themselves toxic to the grazer, they should reflect genuine toxicity to the grazer if the cell is consumed. If the signal molecule is different from the toxic molecules, this would allow for the evolution of Batesian mimicry, that is, cheaters that produce the signal but not the costly toxin. Bioluminescence triggered by prey handling may be another way to signal toxicity (Lindström et al. 2017).

For the two *K. brevis* strains, the low capture and consumption rates cannot be fully accounted for by low feeding activity and high rejection rates, and an additional mechanism may operate. We consider two possibilities: (1) the copepod can identify these cells at distance through olfaction (smell) rather than gustation (taste) and decide not to capture them or (2) subsequent to consumption of some cells, the copepods’ “motivation” to capture further cells of any kind declines, yet it keeps producing a feeding current to continuously sample for palatable prey. The former response would allow an evolutionary stable defense strategy in the phytoplankton while the latter would benefit competitors and cheaters equally well. It has earlier been suggested that organic molecules leaking from phytoplankton cells arriving in the sheared feeding current of a copepod

![Graphs](image-url)
would reach the copepod before the cell itself and allow the copepod to perceive, respond to, and capture the cell (Strickler 1982, Andrews 1983, Jiang et al. 2002). Subsequent observations and theoretical analyses have, however, suggested that this is unfeasible as a general prey perception mechanism simply because the leakage rates of sugars, amino acids, or other molecules common to phytoplankton cells appear insufficient to provide a signal that significantly exceeds the background (Tiselius et al. 2013, Gonçalves and Kiørboe 2015, Kiørboe et al. 2016). However, more specific molecules, such as dimethyl sulfide (DMS) that is produced by some phytoplankton, may elicit responses at very low concentrations in both microorganisms (Strom et al. 2003, Seymour et al. 2010) and copepods (Steinke et al. 2006). Similarly, molecules signaling toxicity may be detected at homeopathic concentrations, thus allowing remote discrimination of toxic cells. There is some experimental evidence for this model (Schultz and Kiørboe 2009).

Diversity of grazer responses and the role of toxins as a defense and competition tools in phytoplankton

The responses of grazers to toxic algae appear to be highly diverse and variable both between and within grazer and phytoplankton species, for example, between different populations of the same grazer species and/or between different strains of the same phytoplankton species (Teegarden 1999, Colin and Dam 2002, Xu et al. 2017). It is striking from our studies and from many earlier reports that some cells with obvious toxic effects on copepods, such as cessation of feeding activity (e.g., the response of one strain of *A. tamarense*, S1, in the present experiments), other “sick” motor responses ([Lasley-Rasher et al. 2016, Xu et al. 2018]), regurgitation of consumed cells (Sykes and Huntley 1987, Xu et al. 2017), or even death of the grazer (Rasmussen et al. 2017), are readily consumed, at least until the toxic effect materializes. The observation that toxic algae often have toxic rather than deterrent effects appears inconsistent with the idea that toxin production serves a defensive purpose (Flynn 2008). It is also striking that a phytoplankton species or strain that is toxic to one species or population of copepods is harmless to others (Colin and Dam 2002, Cohen et al. 2007). In some cases, such variability can be traced back to rapid evolutionary adaptation over just a few generations as has been demonstrated for the copepod *Acartia tonsa* to *A. tamarense* (Colin and Dam 2007) and *Cochlodinium polykrikoides* (Jiang et al. 2011) (see also review by Dam [2013]). Finally, we note that the diversity of toxin profiles within species is very high (Ichimi et al. 2002, Tillmann et al. 2009). How can we understand this perplexing diversity in grazer–alga interactions and in toxin profiles if toxin production mainly serves a defensive purpose?

The high diversity of grazer–phytoplankton interactions, the rapid evolutionary adaptation of grazers, and the multiple strains with radically different toxicity profiles found in many dinoflagellates may simply reflect different stages of an ever-ongoing evolutionary “arms race” between predators and their prey (Smetsacek 2001) that results in a sort of “red queen” dynamics (Dieckmann et al. 1995). A grazer adapts to the toxin of a co-occurring toxic alga by becoming resistant to the toxin, and the alga may then evolve to produce new toxins with toxic or deterrent effects; the copepod subsequently evolves resistance that allows it to utilize an important food source, and so on. The constant mixing of water masses and populations in the ocean allows for a sustained high diversity of toxin profiles and behavioral and physiological grazer responses. An additional source of variability is that researchers often expose phytoplankton strains to grazer populations isolated from different geographical regions, and thus, organisms without a coevolutionary history. Finally, mimicry, as described above, may further add to the diversity of phytoplankton–grazer interactions. However, the dominance of deterrent over toxic grazer responses was revealed by our study.

Toxic substances released by some dinoflagellates may suppress the activity of their protozoan grazers (John et al. 2015) or the growth of their competitors (Jonsson et al. 2009) as a result of the collectively produced harmful concentration in the environment. Thus, toxic substances...
released by one strain of *A. tamarense* reduce the grazing from a predatory protist, but a nontoxic co-occurring strain enjoyed equal grazer protection (John et al. 2015). How do such “public goods” and other allelochemical substances that appear to work through cooperation evolve as defense or competition tools? Allelochemical effects typically materialize only at high densities of the defended or competing cells. Jonsson et al. (2009) argued that allelopathic effects may therefore be nonadaptive side effects of individual predator–prey interactions; thus, their evolution is not facilitated by their allelochemical effects. Allelopathic effects may emerge as the result of eco-evolutionary dynamics that depend on a trait that yield individual cells a selective advantage at low cell densities, that is, a “private good” (Driscoll et al. 2016). Such “private goods” may be offensive chemicals that kill or paralyze prey during direct cell-to-cell contact, as has been demonstrated in a few cases for mixotrophic dinoflagellates (Uchida et al. 1999, Adolf et al. 2007, Blossom et al. 2014), and with the collective effect only emerging at high cell densities. Alternatively, it may be toxic signal molecules that reduce the predation risk of individuals, as has been demonstrated here. While we have argued that toxic cells should produce signal molecules to prevent them from being eaten, collective allelochemical effects would only emerge in cases where the signal molecule itself is toxic.

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**Literature Cited**


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