



Interactions between the calanoid copepod *Acartia hongii* and the bloom-forming dinoflagellates *Karenia bicuneiformis* and *K. selliformis*

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Abstract

Copepods are a major component of metazooplankton and important prey for fish and invertebrates such as crabs, shrimps, and flatworms. Certain bloom-forming dinoflagellates can kill copepods, but there is little research on the interactions between copepods and the bloom-forming dinoflagellates *Karenia bicuneiformis* and *K. selliformis*. In this study, the survival and ingestion rates of the calanoid copepod *Acartia hongii* feeding on *K. bicuneiformis* and *K. selliformis* were determined as a function of prey concentration. On day 2, the survival of *A. hongii* incubated with *K. bicuneiformis* was 90–100% at all the tested prey concentrations, while that with *K. selliformis* was 0–20% at ≥ 582 ng C mL⁻¹. Compared to other harmful dinoflagellates from the literature, *K. bicuneiformis* caused low mortality of *Acartia*; however, *K. selliformis* caused almost the highest mortality at similar dinoflagellate concentrations. With increasing mean prey concentration, the ingestion rates of *A. hongii* feeding on *K. bicuneiformis* increased on day 1, but those on *K. selliformis* did not increase. *Acartia hongii* stopped feeding on *K. bicuneiformis* at mean prey concentrations of ≥ 341 ng C mL⁻¹ and *K. selliformis* at all prey concentrations on day 2. At the prey concentration of 1000 ng C mL⁻¹, the ingestion rate of *A. hongii* feeding on *K. bicuneiformis* was moderate among the rates of *Acartia* spp. feeding on harmful dinoflagellates; however, that on *K. selliformis* was the lowest. These results indicate that *K. bicuneiformis* and *K. selliformis* differentially affect the survival and ingestion rates of *A. hongii*.

Keywords Grazing · Kareniaceae · Marine · Mesozooplankton · Mortality · Survival · Toxin

Introduction

Copepods are ubiquitous in the world's oceans and play a crucial role as major components of marine ecosystems (Turner 2004; Rombouts et al. 2009). They feed on bacteria, protists, including autotrophic, heterotrophic, and mixotrophic species, and larval fish (Mullin 1963; Turner et al. 1985; Gifford 1991; Turner and Tester 1992; Jeong 1994; Jeong et al. 2001; Calbet et al. 2007; Besiktepe and Dam 2020; Lee et al. 2023) and are suitable prey for diverse metazoans (Turner 1984; Barroso et al. 2013). As a key link in marine food webs, the population dynamics of copepods can thus affect those of diverse marine organisms (Turner 2004; Kim et al. 2013; Zeldis and Décima 2020). Therefore, elucidating the interactions between copepods and other marine organisms is crucial to understanding marine ecosystem balance and fishery production.

Phototrophic dinoflagellates are a core component of marine ecosystems and play diverse ecological roles as

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primary producers and prey for various predators such as mixotrophic and heterotrophic dinoflagellates, ciliates, copepods, and invertebrate larvae (Kleppel 1993; Taylor et al. 2008; Jeong et al. 2010, 2021; Ok et al. 2023a; You et al. 2023; Kang et al. 2023). However, certain phototrophic dinoflagellate species are harmful to diverse marine organisms; they often cause large-scale mortality in fish, shellfish, and mammals (Landsberg 2002; Tillmann and John 2002; Burkholder et al. 2006; Holmes et al. 2014; Turner 2014; Ok et al. 2017; Basti et al. 2018; Broadwater et al. 2018). Thus, although some dinoflagellates can be prey for copepods, they can kill or adversely affect copepods by decreasing their ingestion rates, fecundity, and egg-hatching success (Shin et al. 2003; Turner 2006, 2014). Therefore, the interactions between harmful dinoflagellates and copepods warrant closer inspection. Furthermore, some phototrophic dinoflagellates often cause harmful algal blooms (HABs), and their abundance varies largely in marine environments (e.g., Band-Schmidt et al. 2010; Karlson et al. 2021; Gu et al. 2022). To identify the critical abundances at which bloom-forming dinoflagellate species affect copepods, copepod survival and ingestion rates should be explored as a function of prey concentrations.

Many species of the dinoflagellate genus *Karenia* produce neurotoxins (Hort et al. 2021). There are 10 described species of *Karenia* (Guiry and Guiry 2023), all of which cause blooms (e.g., Chang 1999; Yang et al. 2001; Botes et al. 2003; Davidson et al. 2009; Steidinger 2009; Yamaguchi et al. 2016; Iwataki et al. 2022; Orlova et al. 2022). Several studies exist on the interactions between the common calanoid copepod *Acartia* and two *Karenia* species, *K. brevis* and *K. mikimotoi*, which have been shown to inhibit copepod survival, ingestion, and fecundity (Uye and Takamatsu 1990; Turner et al. 1998; Speckmann et al. 2006; Breier and Buskey 2007; Cohen et al. 2007; Waggett et al. 2012). However, no studies have reported the interactions between other bloom-forming *Karenia* species and *Acartia*.

Karenia selliformis is known to produce brevetoxins, gymnodimines, or brevenal (Miles et al. 2000, 2003; Haywood et al. 2004; Mardones et al. 2020), and its blooms have been associated with the death of vertebrates, including fish and sea birds, and invertebrates, including abalone, chitons, sea anemones, sea urchins, sea stars, snails, limpets, and

octopuses, in many countries (Arzul et al. 1995; Mackenzie et al. 1996; Clément et al. 2001; Heil et al. 2001; Uribe and Ruiz 2001; Orlova et al. 2022). Although the lethality of *K. selliformis* has been reported in diverse marine organisms, such as phytoplankton, manila clams, oyster larvae, and juvenile kelp sporophytes (Hégaret et al. 2007; Natsuike et al. 2023; Ok et al. 2023b), no harmful effects have been reported for copepods. Moreover, *Karenia bicuneiformis* (= *K. bidigitata*) contains brevetoxins (Haywood et al. 2004); however, no harmful effects from its blooms have been reported to date when blooms by this species occurred in Gordon Bay, South Africa, and Benguelan current system (Botes et al. 2003; Trainer et al. 2010). Furthermore, the interactions between *K. bicuneiformis* and copepods have not yet been explored.

In this study, the survival and ingestion rates of *Acartia hongii*, feeding on *K. bicuneiformis* CAWD81 and *K. selliformis* NIES-4541, were determined as a function of prey concentration. These observed survival and ingestion rates were compared with those of *Acartia* spp. on other harmful dinoflagellate species based on literature data. Our results provide a basis for understanding the interactions between copepods and harmful *Karenia* species and HAB dynamics.

Materials and methods

Preparation of experimental organisms

A culture of *K. bicuneiformis* CAWD81, initially collected from Fouveaux Strait, New Zealand, and a culture of *K. selliformis* NIES-4541, initially collected from Katsurakoi Port, Japan, were obtained from the Cawthron Institute Culture Collection of Microalgae and the Microbial Culture Collection at the National Institute for Environmental Studies, Japan, respectively (Table 1). All *Karenia* cultures were maintained in 250-mL flat culture flasks containing L1 medium at 20 °C under a 14:10 h light–dark cycle (50 $\mu\text{mol photons m}^{-2} \text{s}^{-1}$ produced by a cool-white fluorescent light). Every 3 weeks, they were transferred into identical flasks containing fresh L1 medium (Guillard and Hargraves 1993). The cell volumes of *K. bicuneiformis* (ellipsoid) and *K. selliformis* (prolate spheroid) were calculated following

Table 1 Information on the experimental organisms used in this study

Organism	Strain	Collection site	Time	T	S	References
<i>Karenia bicuneiformis</i> (= <i>Karenia bidigitata</i>)	CAWD81	Fouveaux Strait, New Zealand	NA	NA	NA	
<i>Karenia selliformis</i>	NIES-4541	Katsurakoi Port, Japan	Sep 24, 2021	16.5	28.2	Iwataki et al. (2022)
<i>Acartia hongii</i>		Shiwha Bay, Korea	Jun 01, 2023	18.2	32.1	This study

T water temperature (°C), S salinity, NA not available

Hillebrand et al. (1999). The carbon content of each culture was estimated using the measured cell volumes following Menden-Deuer and Lessard (2000).

Using a 303- μm mesh net, copepods were collected from Shiwha Bay, Korea, in June 2023, when the water temperature and salinity were 18.2 °C and 32.1, respectively. They were transported to the laboratory within a few hours (Table 1). The copepods were acclimatized in a temperature-controlled room at 20 °C. They were fed with the dinoflagellate *Prorocentrum cordatum* (approximately 200 cells mL^{-1}) as prey for 1 day for the *K. selliformis* experiment and additional 2 days for the *K. bicuneiformis* experiment. Adult female copepods collected during the same sampling event were used throughout this study.

Survival and ingestion rates

In Experiments 1–2, the survival and ingestion rates of *A. hongii*, feeding on either *K. bicuneiformis* (Experiment 1) or *K. selliformis* (Experiment 2), were measured as a function of the prey concentration (Table 2). The initial concentrations of *K. bicuneiformis* (or *K. selliformis*) were set up using autopipettes, and the initial copepod densities (10 female copepods per 500-mL bottle) were established by transferring the copepods individually using a Pasteur pipette (Table 2). Triplicate 500-mL polycarbonate experimental (predator plus prey) and prey control bottles (prey only) were established at each predator–prey density combination. Moreover, triplicate 500-mL predator control bottles (predator only) containing 10 female copepod individuals without prey were also established. Subsequently, a 10-mL aliquot was removed from each experimental and prey control bottle, transferred to a scintillation vial, and fixed with 2% Lugol's solution at the beginning of the experiment to determine the actual initial prey concentrations. The bottles were then filled with freshly filtered seawater, capped, placed on plankton wheels rotating at 0.9 rpm (0.00017 g), and incubated at 20 °C under 20 $\mu\text{mol photons m}^{-2} \text{ s}^{-1}$ illumination with a 14:10 h light–dark cycle. After incubation for 24 h, 10-mL aliquots were extracted from each experimental and prey control bottle and fixed with 2% Lugol's solution. The number of living copepods was enumerated by careful observation through the sides of the capped bottles. The

bottles were then refilled to capacity with freshly filtered seawater, capped, placed on rotating plankton wheels, and incubated under the same conditions mentioned above. After the bottles were incubated for 48 h, 10-mL aliquots were extracted from each experimental and prey control bottle and fixed with 2% Lugol's solution. The water in the experimental bottles was filtered through a 100- μm mesh, and the retained copepods were quickly placed into 50-mL beakers. All living and dead copepods were then enumerated under a dissecting microscope (SZX2-ILLB, Olympus, Tokyo, Japan). The prey concentration was measured by counting all the cells, or > 200 cells in 1-mL Sedgewick Rafter chambers under a compound microscope (BX53; Olympus). When Experiment 2 was conducted, the survival and ingestion rates of *A. hongii* feeding on *Prorocentrum cordatum* as a non-toxic prey control were measured at a single high prey concentration (10,000 cells mL^{-1}) in the same manner as described above.

Using the equations described by Frost (1972), the ingestion and clearance rates of the copepods for each target *Karenia* species at day 1 (from day 0 to day 1) and day 2 (from day 1 to day 2) were calculated. The dilution of the cultures associated with refilling the bottles on days 0 and 1 was considered in calculating the ingestion rates. The data on the *K. bicuneiformis* ingestion rates were fitted to a Michaelis–Menten equation:

$$IR = \frac{I_{\max}(x)}{K_{IR} + (x)} \quad (1)$$

where x = mean prey concentration (cells mL^{-1} or ng C mL^{-1}); I_{\max} = the maximum ingestion rate (cells predator $^{-1}$ day $^{-1}$ or ng C predator $^{-1}$ day $^{-1}$); and K_{IR} = the prey concentration sustaining $\frac{1}{2} I_{\max}$ (cells mL^{-1} or ng C mL^{-1}).

Molecular identification of *Acartia* species used in this study

To identify the copepods used in this study, each of the 10 copepods in the predator control bottle in Experiment 1 (with *K. bicuneiformis*) was transferred to an individual 1.5-mL tube after the experiment had ended. The genomic

Table 2 Experimental design and the actual initial densities of prey and predator species

Expt No	Prey		Predator	
	Species	Actual initial density (cells mL^{-1})	Species	Actual initial density (individuals in 500 mL)
1	<i>Karenia bicuneiformis</i>	0, 20, 97, 188, 464, 986, 1960	<i>Acartia hongii</i>	10
2	<i>Karenia selliformis</i>	0, 19, 72, 145, 390, 793, 1555	<i>A. hongii</i>	10

DNA of the copepods was extracted using the AccuPrep® genomic DNA extraction kit (Bioneer, Daejeon, Korea). For the amplification of small subunit ribosomal DNA (SSU rDNA), polymerase chain reaction (PCR) was conducted using F-Star Taq DNA Polymerase (BIOFACT Co., Daejeon, Korea) in a total volume of 50 μL containing 5 μL of $10\times$ Taq buffer, 1 μL of 10 mM dNTP mix, 0.25 μL of DNA polymerase, 5 μL of $5\times$ Band Helper™, 2 μL each of the forward and reverse primers, and 1 μL of the DNA template. The forward and reverse primers used in this study were 18s1075H (5'-CGA AGG CGM TCA GAT ACC GCC CTA G-3') and 18s1871Hr (5'-CAC CTA CGG AAA CCT TGT TAC GAC-3'), respectively (Easton and Thistle 2014). The thermocycler (AllInOneCycler™, Bioneer) was run under the following conditions: 95 °C for 2 min, followed by 40 cycles of 95 °C for 20 s, 59 °C for 30 s, and 72 °C for 1 min, followed by a final extension at 72 °C for 5 min. The PCR products were purified using the Wizard® SV Gel and PCR Clean-Up System (Promega, Madison, WI, USA) following the manufacturer's instructions. They were then sequenced using an ABI PRISM® 3700 DNA Analyzer (Applied Biosystems, Foster City, CA, USA). The obtained sequences were assembled using ContigExpress software (Infomax, Frederick, MD, USA) and compared with the SSU rDNA sequences of *Acartia* species in the NCBI GenBank database based on the BLASTn algorithm. Sequences with > 90% query coverage are shown in the sequence difference comparisons in this study.

For the phylogenetic tree, SSU rDNA sequences of the *Acartia* species were obtained from GenBank and were aligned using the MEGA v4 program (Tamura et al. 2007). A phylogenetic tree based on the Bayesian and maximum likelihood analyses was established following Kang et al. (2010).

Swimming speed of *Karenia*

To explore whether the ingestion rates of *Acartia* species feeding on dinoflagellates were affected by the swimming speeds of the dinoflagellates, the swimming speeds of *K. bicuneiformis* CAWD81 and *K. selliformis* NIES-4541 were measured. A dense clonal culture of *K. bicuneiformis* (or *K. selliformis*) in a 50-mL cell culture flask, incubated at 20 °C under 20 $\mu\text{mol photons m}^{-2} \text{s}^{-1}$, was placed under a dissecting microscope (SZX10, Olympus). The field in which the flask was placed was leveled using a magnetic torpedo level. After 30-min acclimation, the swimming *Karenia* species were recorded in each flask using a video analyzing system (SRD-1673DN; Samsung Techwin, Seongnam, Korea) at a magnification of 20 \times . The swimming speeds of *K. bicuneiformis* (or *K. selliformis*) were calculated by measuring the travel distance and time of 20 swimming *Karenia* cells showing straight, linear paths.

Statistical analysis

The non-parametric Kruskal–Wallis test was used to examine the effect of *Karenia* concentration on *A. hongii* survival. Pearson's correlations were conducted to examine the relationships between the ingestion rates of *A. hongii* feeding on dinoflagellates, the mortality (%) of *A. hongii* incubated with the dinoflagellates, and the dinoflagellate cell sizes and swimming speeds. The software SPSS 26.0 (IBM-SPSS Inc., NY, USA) was used for statistical analyses. The criterion for statistical significance was set at a P value of 0.05.

Results

Molecular identification of *A. hongii*

The obtained SSU rDNA sequences of the 10 copepod individuals used in this study were 100% identical to each other and to those reported for *A. hongii* (Accession numbers GU969195 and MZ413971; Fig. 1). The *A. hongii* sequences obtained in this study were 13.2–15.8% different from those of other *Acartia* species such as *A. omorii*, *A. clausii*, *A. tonsa*, *A. bifilosa*, *A. negligens*, *A. danae*, *A. steueri*, *A. pacifica*, and *A. erythraea*. The SSU rDNA sequence of *A. hongii* used in this study was deposited in GenBank under accession number OR356145.

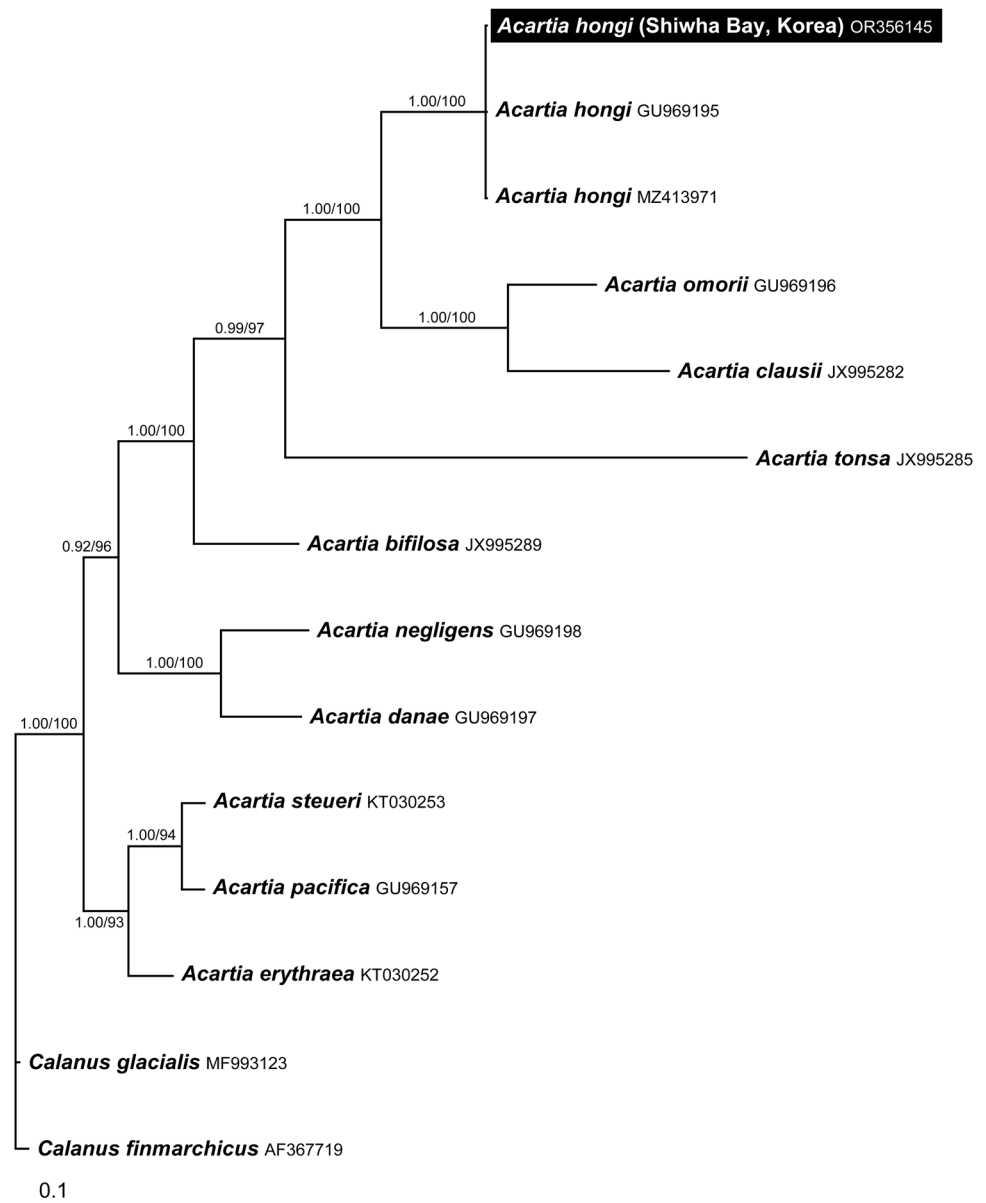
Survival of *A. hongii*

On day 1, the survival of *A. hongii* incubated with *K. bicuneiformis* at mean prey concentrations of 0–1352 ng C mL⁻¹ (0–1988 cells mL⁻¹) was 87–100% (Fig. 2a). On day 2, they were 90–100% at mean prey concentrations of 0–1394 ng C mL⁻¹ (0–2050 cells mL⁻¹) (Fig. 2b). The survival was not significantly affected by *K. bicuneiformis* concentration (Kruskal–Wallis test; $H_6 = 8.97$, $P = 0.175$ for day 1; $H_6 = 9.12$, $P = 0.167$ for day 2).

On day 1, the survival of *A. hongii* incubated with *K. selliformis* at mean prey concentrations of 0–2400 ng C mL⁻¹ (0–1690 cells mL⁻¹) was 87–97% (Fig. 3a). On day 2, the survival decreased as a function of mean prey concentrations (Fig. 3b); all the copepods had died at a mean prey concentration of 2682 ng C mL⁻¹ (1889 cells mL⁻¹). The survival was not significantly affected by *K. selliformis* concentrations on day 1 (Kruskal–Wallis test, $H_6 = 4.03$, $P = 0.673$); however, they were significantly affected on day 2 (Kruskal–Wallis test, $H_6 = 18.23$, $P = 0.006$).

The survival of *A. hongii* incubated with *P. cordatum* was 95% at a single mean prey concentration of 1452 ng C mL⁻¹ (9680 cells mL⁻¹) for 2 days.

Fig. 1 Bayesian tree showing genetic relationships among the calanoid copepod *Acartia* based on the small subunit ribosomal DNA region (707 bp), using a GTR+G+I model. Numbers near branches represent the Bayesian posterior probability (left) and maximum likelihood bootstrap values (right). A black box represents *A. hongii*, which was used in this study. The assumed empirical nucleotide frequencies of SSU rDNA comprised a substitution rate matrix with A–C substitutions = 0.1152, A–G = 0.2406, A–T = 0.1173, C–G = 0.1114, C–T = 0.2874, and G–T = 0.1281. Rates were assumed to follow a gamma distribution with a shape parameter of 0.3661 for variable sites. The proportion of sites assumed to be invariable was 0.2416



Ingestion rate of *A. hongii*

On day 1, the ingestion rates of *A. hongii* feeding on *K. bicuneiformis* continuously increased with increasing mean prey concentrations (Fig. 4a). The highest ingestion rate of *A. hongii* on *K. bicuneiformis*, 3813 ng C predator⁻¹ day⁻¹ (5608 cells predator⁻¹ day⁻¹), was achieved at a mean prey concentration of 1352 ng C mL⁻¹ (1988 cells mL⁻¹). When Eq. (1) was used, the I_{\max} of *A. hongii* feeding on *K. bicuneiformis* was 9286 ng C predator⁻¹ day⁻¹ (13,656 cells predator⁻¹ day⁻¹), and the K_{IR} was 1710 ng C mL⁻¹ (2515 cells mL⁻¹). The maximum clearance rate of *A. hongii* feeding on *K. bicuneiformis* was 338 μ L predator⁻¹ h⁻¹ at a mean prey concentration of 13 ng C mL⁻¹ (19 cells mL⁻¹). On day 2, the ingestion

rates of *A. hongii* feeding on *K. bicuneiformis* increased at mean prey concentrations ≤ 116 ng C mL⁻¹ (171 cells mL⁻¹); however, they decreased to zero at higher prey concentrations, indicating that ingestion was inhibited at high prey concentrations (Fig. 4b).

On day 1, the ingestion rates of *A. hongii* feeding on *K. selliformis* were positive at mean prey concentrations ≤ 1159 ng C mL⁻¹ (816 cells mL⁻¹). However, the rate decreased to zero at a higher prey concentration (Fig. 5a). The highest ingestion rate of *A. hongii* on *K. selliformis*, 1373 ng C predator⁻¹ day⁻¹ (967 cells predator⁻¹ day⁻¹), was achieved at a mean prey concentration of 1159 ng C mL⁻¹ (816 cells mL⁻¹). The maximum clearance rate of *A. hongii* feeding on *K. selliformis* was 1612 μ L predator⁻¹ h⁻¹ at a mean prey concentration of 18 ng C mL⁻¹ (13 cells mL⁻¹). On day 2, the ingestion rates of *A.*

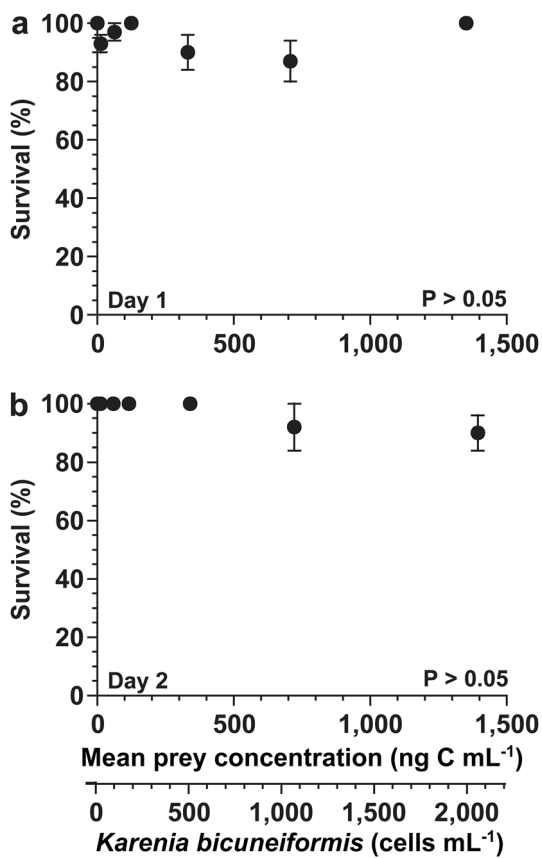


Fig. 2 Survival (%) of *Acartia hongii* as a function of mean *Karenia bicuneiformis* concentration on day 1 (a) and day 2 (b). Symbols represent treatment means \pm standard error

hongii feeding on *K. selliformis* were zero at all prey concentrations (Fig. 5b).

The ingestion rate of *A. hongii* feeding on *P. cordatum* was 4541 ng C predator⁻¹ day⁻¹ (30,273 cells predator⁻¹ day⁻¹) at a single mean prey concentration of 1452 ng C mL⁻¹ (9680 cells mL⁻¹) for 2 days.

Swimming speed of *Karenia*

Karenia bicuneiformis and *K. selliformis* showed linear and helical swimming behaviors; however, jumping behaviors were not observed. The average (\pm standard error) and maximum swimming speeds of *K. bicuneiformis* (n=20) were 257 (\pm 17) and 480 $\mu\text{m s}^{-1}$, respectively. The average and maximum swimming speeds of *K. selliformis* (n=20) were 232 (\pm 13) and 340 $\mu\text{m s}^{-1}$, respectively.

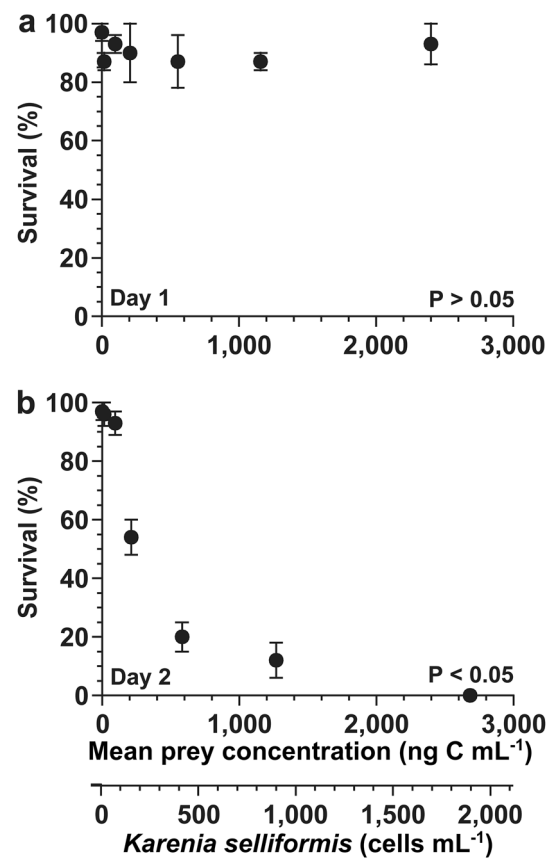


Fig. 3 Survival (%) of *Acartia hongii* as a function of mean *Karenia selliformis* concentration on day 1 (a) and day 2 (b). Symbols represent treatment means \pm standard error

Discussions

The present study reports, for the first time, the interactions between *Acartia* species and the dinoflagellates *K. bicuneiformis* CAWD81 and *K. selliformis* NIES-4541. Previous studies on the interactions between *Acartia* species and *K. brevis*, *Acartia* species and *K. mikimotoi*, and *Temora longicornis* and *K. selliformis* K-1319 have been explored (Uye and Takamatsu 1990; Turner et al. 1998; Breier and Buskey 2007; Cohen et al. 2007; Waggett et al. 2012; Xu and Kiørboe 2018).

This study highlights that *A. hongii* feeds on *K. bicuneiformis* and *K. selliformis*; however, it stops feeding on them depending on the mean prey concentration and incubation time. The highest ingestion rate of *A. hongii* when feeding on *K. bicuneiformis* was higher than that when feeding on *K. selliformis*. Furthermore, *A. hongii* died when exposed to *K. selliformis* but not when exposed to *K. bicuneiformis*. Therefore, *K. bicuneiformis* and *K. selliformis* interact with *A. hongii* differently.

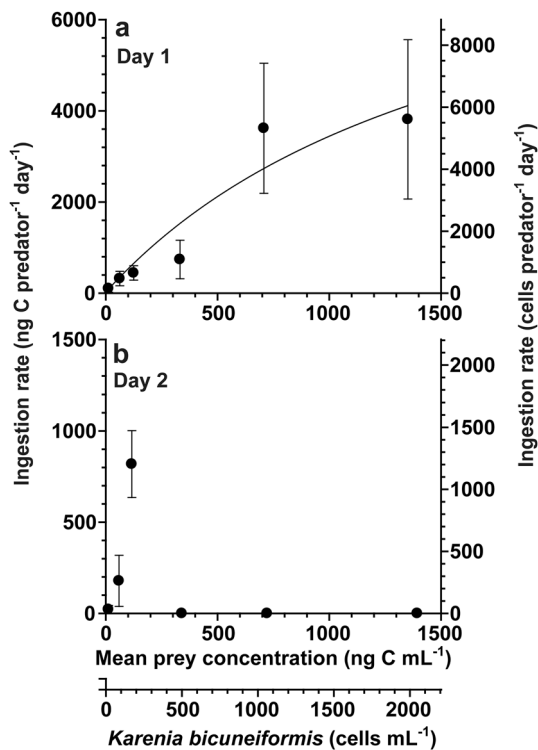


Fig. 4 Ingestion rates of *Acartia hongii* feeding on *Karenia bicuneiformis* as a function of mean prey concentration (x) on day 1 (a) and day 2 (b). The curve in (a) is fitted using the Michaelis–Menten equation. Ingestion rate ($\text{ng C predator}^{-1} \text{ day}^{-1}$) = $9286 [x/(1710+x)]$, $r^2=0.525$. Symbols represent treatment means \pm standard error

Effects of two *Karenia* species on the survival of *A. hongii*

Acartia hongii was killed by *K. selliformis* NIES-4541 but not by *K. bicuneiformis* CAWD81. Before this study, in the family Kareniaceae, *K. brevis* CCMP2228, the Nagasaki University strain of *K. mikimotoi*, *Karlodinium armiger* K-0668, and the Alfacs Bay strain of *Karlodinium corsicum* were known to kill *Acartia* spp., whereas *Karlodinium veneficum* K-1385, K-1635, K-1640, and K-1386 does not kill them (Uye and Takamatsu 1990; Delgado and Alcaraz 1999; Waggett et al. 2012; Berge et al. 2012). At similar Kareniacean dinoflagellate concentrations (1100–1500 ng C mL^{-1}) after 2 days of incubation, the mortality (the inversion of the survival) of *A. hongii* incubated with *K. selliformis* NIES-4541 was slightly lower than those of *Acartia* spp. incubated with *Kl. armiger* K-0668 and the Alfacs Bay strain of *Kl. corsicum*; however, much higher than those of *Acartia* spp. incubated with *K. brevis* CCMP2228, *K. bicuneiformis* CAWD81, and *Kl. corsicum* GCORS1 (Table 3; Fig. 6). These comparisons indicate that *K. selliformis* is more harmful to *A. hongii* than *K. bicuneiformis*. To date, there have been no reports on the mortality of marine animals during *K. bicuneiformis* blooms (e.g., Botes et al. 2003; Trainer

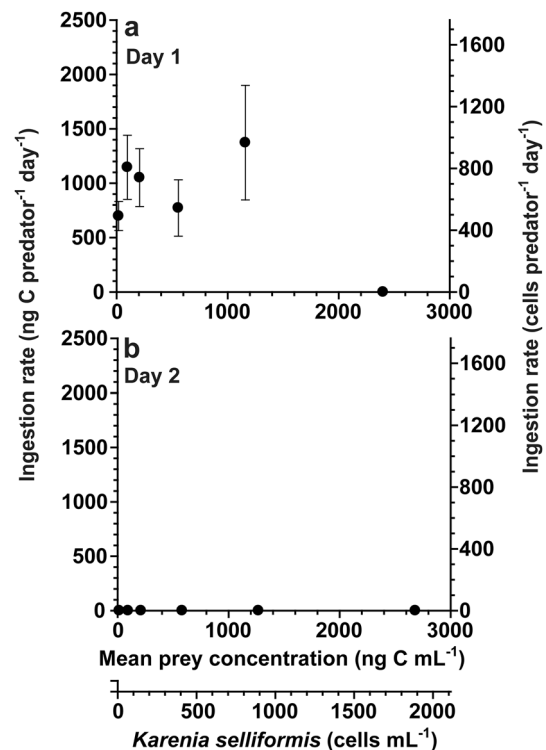


Fig. 5 Ingestion rates of *Acartia hongii* feeding on *Karenia selliformis* as a function of mean prey concentration on day 1 (a) and day 2 (b). Symbols represent treatment means \pm standard error

et al. 2010). However, reports from various countries, such as Chile, Kuwait, New Zealand, and Russia, have detailed the mortality of marine animals during *K. selliformis* blooms (Mackenzie et al. 1996; Clément et al. 2001; Heil et al. 2001; Uribe and Ruiz 2001; Orlova et al. 2022). Thus, the field reports on the relationships between marine animal mortality and *K. bicuneiformis* and *K. selliformis* blooms are similar to the results of this study, which shows that *K. bicuneiformis* and *K. selliformis* have different effects on the survival of *A. hongii*.

Some strains in the genus *Karenia* have diverse toxin compositions such as brevetoxins, brevenal, gymnodimines, and gymnocins (Baden 1989; Seki et al. 1995; Miles et al. 2000, 2003; Satake et al. 2002, 2005; Bourdelais et al. 2005; Rasmussen et al. 2017; Mardones et al. 2020). Therefore, it is difficult to compare the lethality of these Kareniacean dinoflagellate species and strains using the amount of toxins, and a different method of comparing their lethality is needed. The results of this study suggest that comparing the mortality of *Acartia* at similar concentrations of Kareniacean dinoflagellates can be a useful method for understanding the relative harmfulness of these organisms. There may be intraspecific variability in the amount of secondary metabolites produced by each dinoflagellate species (e.g., Burkholder and Glibert 2006), and this amount may also be

Table 3 Mortality (%) of *Acartia* incubated with harmful dinoflagellates at the prey concentrations of 1100–1500 ng C mL⁻¹ for 2 days, calculated ingestion rate (ng C predator⁻¹ day⁻¹) of *Acartia* species feeding on harmful dinoflagellates at 1000 ng C mL⁻¹, and the cell size (ESD, equivalent spherical diameter, μm) and maximum swimming speed (MSS, μm s⁻¹) of the dinoflagellates

Copepod	Dinoflagellate prey (strain or origin)	Mortality	Ingestion rate	ESD	MSS	References
<i>A. tonsa</i>	<i>Karlodinium armiger</i> (K-0668)	100 ^a		14.9 ^f		Berge et al. (2008, 2012)
<i>A. grani</i>	<i>Karlodinium corsicum</i> (from Alfacs Bay)	100		12.6		Delgado and Alcaraz (1999) da Costa and Fernández (2002)
<i>A. hongii</i>	<i>Karenia selliformis</i> (NIES-4541)	90	1241 ^c	29.7	340	This study
<i>A. tonsa</i>	<i>Margalefidinium polykrikoides</i> (CP1)	55 ^b	2393 ^{bc}	28.2	1449 ^h	Jeong et al. (1999) Jiang et al. (2009)
<i>A. tonsa</i>	<i>Karenia brevis</i> (CCMP2228)	42 ^b		22.4 ^g	417 ^h	McKay et al. (2006) Waggett et al. (2012)
<i>A. grani</i>	<i>Gymnodinium catenatum</i> (GC19V)	20	1247 ^c	33.9	440 ^h	Fraga et al. (1989) Jeong et al. (2004) da Costa et al. (2012)
<i>A. hongii</i>	<i>Karenia bicuneiformis</i> (CAWD81)	10	3427 ^d	21.0	480	This study
<i>A. grani</i>	<i>Karlodinium corsicum</i> (GCORS1)	6 ^b		12.6		da Costa and Fernández (2002) da Costa et al. (2005)
<i>A. tonsa</i>	<i>Karenia brevis</i> (Wilson)		3469 ^{ce}	20.4	417 ^h	McKay et al. (2006) Cohen et al. (2007)
<i>A. clausi</i>	<i>Alexandrium minutum</i> (A1 IV)		3688 ^d	22.0	474 ^h	Guisande et al. (2002) Lewis et al. (2006)
<i>A. clausi</i>	<i>Alexandrium minutum</i> (BAH-ME 91)		5136 ^c	19.4	474 ^h	Dutz (1998) Lewis et al. (2006)
<i>A. tonsa</i>	<i>Alexandrium catenella</i> (1119/27)		5703 ^{bc}	25.8 ^{fg}	185 ^h	Karp-Boss et al. (2000) Abdulhussain et al. (2020)
<i>A. hudsonica</i>	<i>Alexandrium catenella</i> (CB-301)		7658 ^{de}	19.2	185 ^h	Karp-Boss et al. (2000) Colin and Dam (2007)

^aThe mortality at 45 h^bData estimated from the figure^cData calculated by linear interpolation using two adjacent data points^dData calculated using the equation provided in the reference^eThe highest ingestion rate of *Acartia* spp. feeding on the same strains in several experiments^fThe median value of the ranges provided in the reference^gData calculated using the equation in Menden-Deuer and Lessard (2000)^hValues of a strain different from the listed strain

affected by the growth phase and culture conditions. Thus, with careful consideration of intraspecific variability and the growth phase of the culture, the relative harm to strains or species should be compared.

Among the dinoflagellates causing mortality of *Acartia* spp., the mortality of *Acartia* spp. incubated with cells of *Kl. armiger* K-0668, the Alfacs Bay strain of *Kl. corsicum*, *K. selliformis* NIES-4541, *Margalefidinium polykrikoides* CP1, *K. brevis* CCMP2228, and *Gymnodinium catenatum* GC19V at the dinoflagellate concentrations of 1100–1500 ng C mL⁻¹ for 2 d was significantly higher than that in controls such as being incubated without the dinoflagellate species, with their lowest cell density, or with non-toxic prey species (Fig. 6). Physical contact and chemicals (toxins or inhibitory substances) can cause mortality of *Acartia* spp. incubated with these dinoflagellates. Delgado and Alcaraz (1999) reported

that the death of *A. grani* was caused by direct physical contact with the Alfacs Bay strain of *Kl. corsicum*. Berge et al. (2012) also reported that *Kl. armiger* K-0668 attacked, immobilized, and ingested *A. tonsa*. In addition, *Kl. armiger* K-0668, *K. brevis* CCMP2228, and *G. catenatum* GC19V contain karmitoxin, brevetoxins, and gonyautoxins and saxitoxin, respectively, which cause mortality of *Acartia* spp. (da Costa et al. 2012; Waggett et al. 2012; Rasmussen et al. 2017). Moreover, *M. polykrikoides* CP1 has harmful effects on copepods and contains reactive oxygen species-like chemicals (Jiang et al. 2009; Tang and Gobler 2009). Therefore, toxins or inhibitory substances of *Kl. armiger* K-0668, *M. polykrikoides* CP1, *K. brevis* CCMP2228, and *G. catenatum* GC19V are likely to cause mortality in *Acartia* spp. Some strains of *K. selliformis* are known to contain toxins such as gymnodimines or brevenal (Miles et al. 2000,

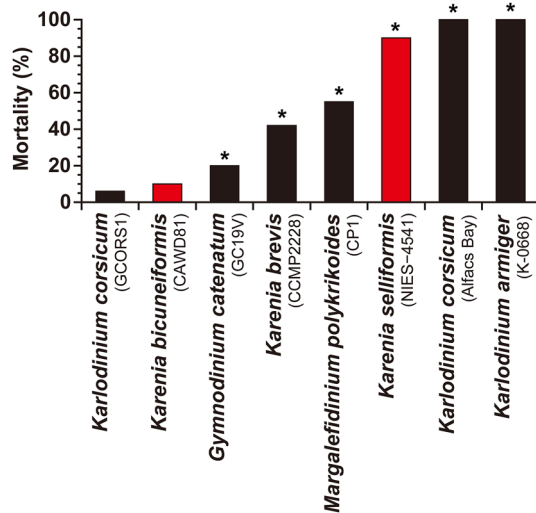


Fig. 6 Mortality (%) of *Acartia* species incubated with harmful dinoflagellates at 1100–1500 ng C mL⁻¹ concentrations for 2 days. Red bars indicate the mortality measured in this study. An asterisk on each bar indicates that the value in this figure was significantly or ≥20% different from the mortality of *Acartia* species without the target dinoflagellate species (i.e., predator-only control), with the lowest cell density, or with non-toxic prey species in the reference. See Table 3 for details

2003; Mardones et al. 2020), and thus *K. selliformis* NIES-4541 may potentially possess toxins leading to mortality of *A. hongii*. However, the presence of toxins of *K. selliformis* NIES-4541 has not been reported. The Alfacs Bay strain of *Kl. corsicum* at 1100–1500 cells mL⁻¹ for 2 days of incubation resulted in high mortality of *A. grani*, while *Kl. corsicum* GCORS1 did not cause significant mortality of *A. grani* at similar *Kl. corsicum* concentrations, implying intraspecific variability (Fig. 6). Therefore, considering potential intraspecific variability, further examination is needed to confirm the presence of toxins of *K. selliformis* NIES-4541.

Effects of two *Karenia* species on the ingestion rate of *A. hongii*

The calculated ingestion rates of *Acartia* spp. on seven reported harmful dinoflagellate species (nine strains) at 1000 ng C mL⁻¹ varied, ranging from 1241 to 7658 ng C predator⁻¹ day⁻¹ (Table 3; Fig. 7a). The calculated ingestion rate of *A. hongii* feeding on *K. bicuneiformis* at 1000 ng C mL⁻¹ was lower than those of *Acartia* spp. feeding on *Alexandrium catenella*, *Alexandrium minutum*, and *K. brevis*; however, it was higher than those of *Acartia* spp. feeding on *M. polykrikoides*, *G. catenatum*, and *K. selliformis*. Thus, the ingestion rate of *A. hongii* feeding on *K. bicuneiformis* was close to the median of those of *Acartia* spp. feeding on harmful dinoflagellates. The calculated ingestion rate of *A. hongii* feeding on *K. selliformis* was lower than that of

Acartia spp. feeding on the other six harmful dinoflagellates (eight strains) at a prey concentration of 1000 ng C mL⁻¹. The mortality of *A. hongii* incubated with *K. selliformis* for 2 days was higher than those of *Acartia* spp. incubated with *M. polykrikoides*, *G. catenatum*, and *K. bicuneiformis* (Fig. 7b). Thus, this high *A. hongii* mortality may be partially responsible for the low ingestion rate of *A. hongii* feeding on *K. selliformis*.

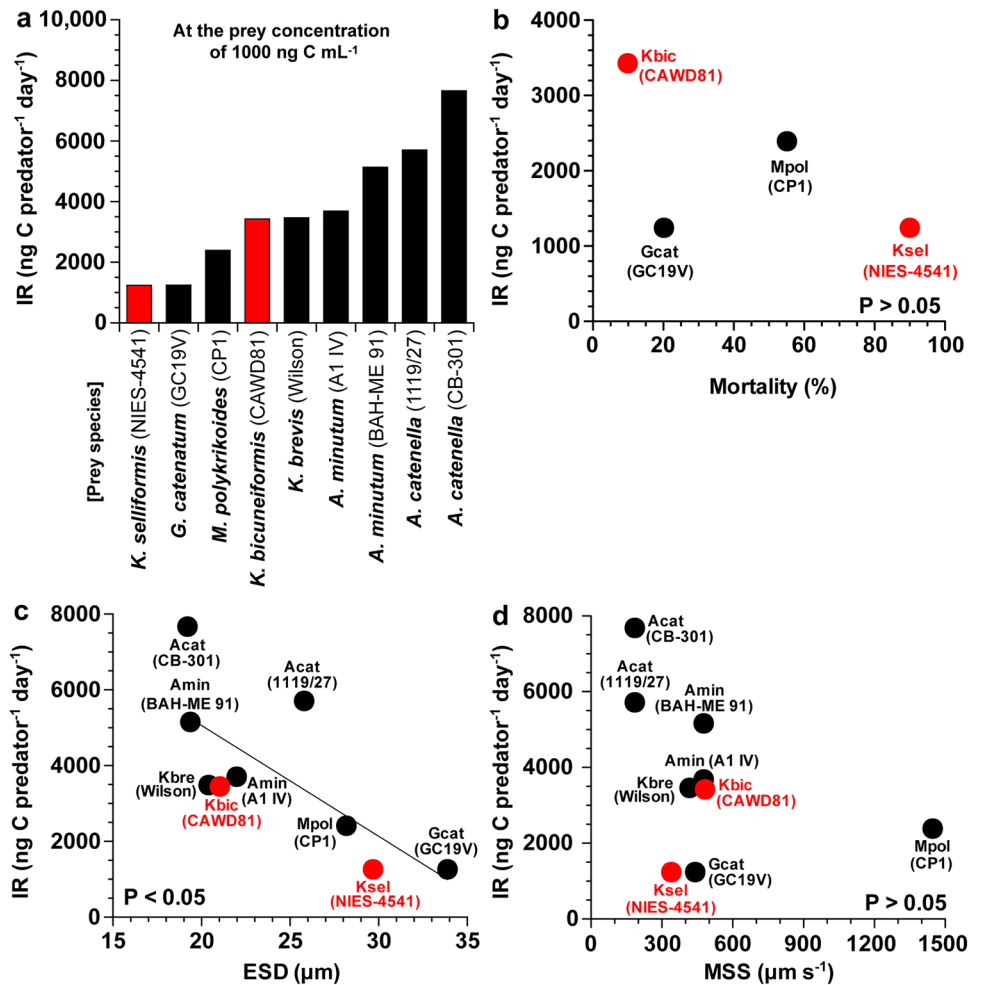
The calculated ingestion rates of *Acartia* spp. feeding on the nine harmful dinoflagellate strains at a prey concentration of 1000 ng C mL⁻¹ showed a negative correlation with the equivalent spherical diameters of the harmful dinoflagellates (19.2–33.9 μm; Pearson's correlation; $r = -0.727$, $P = 0.026$; Fig. 7c). The calculated ingestion rate of *A. hongii* feeding on *K. selliformis* at 1000 ng C mL⁻¹ was the lowest, while its equivalent spherical diameter was the second largest. Nival and Nival (1976) reported that modified filtration efficiency could decrease when particle sizes are ≥ 26.2 μm. Thus, the large size of *K. selliformis* may be partially responsible for the relatively low ingestion rate of *A. hongii* feeding on *K. selliformis*.

The calculated ingestion rates of *Acartia* spp. feeding on harmful dinoflagellates at a prey concentration of 1000 ng C mL⁻¹ were not significantly correlated with the maximum swimming speeds of the nine harmful dinoflagellate strains (Pearson's correlation; $r = -0.407$, $P = 0.278$; Fig. 7d). The calculated ingestion rate of *A. hongii* feeding on *K. bicuneiformis* was higher than that of *A. hongii* feeding on *K. selliformis*, and the maximum swimming speed of *K. bicuneiformis* was higher than that of *K. selliformis*. Contrary to this trend, the calculated ingestion rate of *A. tonsa* feeding on *M. polykrikoides* was relatively low, but *M. polykrikoides* has the highest maximum swimming speed among the harmful dinoflagellates (Jiang et al. 2009; Jeong et al. 2015). Thus, the fast swimming speed of *M. polykrikoides* may lower the ingestion rate of *Acartia* feeding on it; however, the maximum swimming speed of *K. selliformis* may not be high enough to lower the ingestion rate of *Acartia*.

Difference in the survival and ingestion rates on days 1 and 2

The survival and ingestion rates of *A. hongii* feeding on *K. selliformis* NIES-4541 on day 2 were much lower than those on day 1. Furthermore, the ingestion rates of *A. hongii* feeding on *K. bicuneiformis* CAWD81 at high mean prey concentrations on day 2 were much lower than those on day 1, although the survival on day 2 was not different from those on day 1. Therefore, differences in the survival or ingestion rates of *A. hongii* feeding on *K. bicuneiformis* and *K. selliformis* were observed as the incubation time increased. Thus, if the target dinoflagellate species are harmful, it is

Fig. 7 Calculated ingestion rates ($\text{ng C predator}^{-1} \text{ day}^{-1}$) of *Acartia* species feeding on harmful dinoflagellates at the prey concentration of $1000 \text{ ng C mL}^{-1}$ (IR; **a**). Correlations of IR with mortality (%) of *Acartia* species incubated with the dinoflagellates at $1100\text{--}1500 \text{ ng C mL}^{-1}$ (**b**) for 2 days, equivalent spherical diameters (ESD) of the dinoflagellates (**c**), and maximum swimming speeds (MSS) of the dinoflagellates (**d**). Red bars or circles indicate the results of this study. See Table 3 for details. Linear equation in (**c**): $\text{IR} = -292 (\text{ESD}) + 10,900$, $r^2 = 0.529$



reasonable to measure the survival and ingestion rates of *Acartia* on days 1 and 2, with subsampling on days 0, 1, and 2.

Ecological implications

Blooms of *K. selliformis* have been reported in many countries, such as Chile, Japan, Kuwait, New Zealand, Russia, and Tunisia (Mackenzie et al. 1996; Clément et al. 2001; Heil et al. 2001; Elleuch et al. 2021; Iwataki et al. 2022; Orlova et al. 2022; Boudriga et al. 2023). During these blooms, the reported *K. selliformis* abundance ranges from 43 to 60,000 cells mL^{-1} ; however, it is usually ≥ 140 cells mL^{-1} (Table 4). In the present study, half of *A. hongii* were killed at a *K. selliformis* concentration of 149 cells mL^{-1} on day 2. Moreover, the surviving *A. hongii* did not feed on *K. selliformis* on day 2. Therefore, during *K. selliformis* blooms, *Acartia* spp. may be killed by the inhibiting substances from *K. selliformis* or starvation. Exploring the absence of *Acartia* spp. during *K. selliformis* blooms will be beneficial.

Karenia selliformis is known to lyse several phytoplankton, such as the prasinophyte *Pyramimonas* sp. and the cryptophytes *Teleaulax amphioxiea* and *Storeatula major* (Ok et al. 2023b). Therefore, *K. selliformis* may eliminate alternative prey for the copepods. During *K. selliformis* blooms, *Acartia* spp. may have difficulty maintaining their populations due to the direct (i.e., production of inhibiting substances) and indirect effects of the dinoflagellate (i.e., removal of alternative prey).

The results of this study show that *K. bicuneiformis* was predated by *A. hongii* at prey concentrations of < 500 cells mL^{-1} but not at higher prey concentrations on day 2. Thus, *K. bicuneiformis* cells may be eaten by *A. hongii* at the initial stage of a *K. bicuneiformis* bloom but not during the main bloom.

The interactions between copepods and six of the ten described *Karenia* species (i.e., *K. asterichroma*, *K. brevisulcata*, *K. concordia*, *K. cristata*, *K. longicanalis*, and *K. papilionacea*) have not yet been explored. These *Karenia* species are known to cause HABs (Chang 1999; Yang et al. 2001; Botes et al. 2003; de Salas et al. 2004; Chang

Table 4 Locations and times of *Karenia selliformis* blooms and the highest abundance (cells mL⁻¹) during each bloom

Location	Time	Abundance	References
Magellanic fjords, Chile	Apr, 1999	43	Uribe and Ruiz (2001)
Spaseniya Bay, Russia	Oct, 2020	66	Orlova et al. (2022)
Foveaux Strait, New Zealand	Feb, 1994	140	MacKenzie et al. (1996)
First Kuril Strait, Russia	Oct, 2020	162	Orlova et al. (2022)
Utashud Island, Russia	Oct, 2020	254	Orlova et al. (2022)
Three Sisters Bay, Russia	Oct, 2020	482	Orlova et al. (2022)
Sarannaya Bay, Russia	Oct, 2020	622	Orlova et al. (2022)
Timaru Harbor, New Zealand	Apr, 1994	> 900	MacKenzie et al. (1996)
Ellouza, Tunisia	Jun–Jul, 2020	1604 ^a	Elleuch et al. (2021)
Kuwait Bay, Kuwait	Oct, 1999	> 6000	Heil et al. (2001)
Chiloé Archipelago, Chile	Mar–Apr, 1999	8000	Clément et al. (2001)
Hiroo, Japan	Oct, 2021	9600	Iwataki et al. (2022)
Toyokoro, Japan	Oct, 2021	10,560	Iwataki et al. (2022)
Sfax, Tunisia	Sep–Oct, 2019	27,000	Boudriga et al. (2023)
Sfax, Tunisia	Oct, 1994	60,000	Arzul et al. (1995)

^aqPCR measurement

and Mullan 2012; Yamaguchi et al. 2016). Therefore, to better understand the interactions between copepods and harmful dinoflagellates in marine ecosystems, the survival and ingestion rates of copepods feeding on the six *Karenia* species as a function of the *Karenia* concentration should be explored.

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Author contributions JHO and HJJ designed the study conception and drafted the manuscript. MJL collected copepods. JHO, JHY, SAP, HCK, SHE, and JR conducted experiments including feeding experiments and swimming speed measurement. JHO and HJJ conducted data analyses. All authors discussed the results.

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Data availability Data collected and analyzed during this study are available from the corresponding author on reasonable request.

Declarations

Conflict of interest The authors declare that they have no competing interests.

Ethical approval All applicable international, national, and/or institutional guidelines for the care and use of organisms were followed.

Consent to participate Not applicable.

Consent to publish All authors consent to the publication of this manuscript.

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