



OPEN

Snails associated with the coral-killing sponge *Terpios hoshinota* in Okinawa Island, Japan

Hideyuki Yamashiro¹✉, Hiroaki Fukumori^{1,2}, Siti Nurul Aini³ & Yurika Hirose^{3,4}

Terpios hoshinota is a thin encrusting sponge that overgrows live scleractinian corals and it is linked to coral loss in many reefs. However, our knowledge of the species associated with this sponge species is poor. During a periodical survey of *T. hoshinota* in 2020, we found tiny snails crawling on the sponge in the subtropical waters around Okinawa Island, Japan. We observed egg capsules inside the sponge tissue and veliger larvae released from the egg capsules. Molecular analyses of both the snails and veliger larvae (cytochrome oxidase I, COI) showed that they were identical and belonged to *Joculator* sp. (family Cerithiopsidae). There was no direct observation of predation on the sponge by this snail; however, to the best of our knowledge, this is the first report on a close association between a snail and the sponge *T. hoshinota*.

Coral reefs are valuable ecosystems that supply numerous services to humans, and they are home to numerous coral-associated organisms, which are linked to the high levels of biodiversity observed in these ecosystems. However, coral reefs are threatened and degraded by repeated bleaching events, owing to increasing water temperatures, ocean acidification, coral predators, infectious diseases, and physical/chemical disturbances caused by human activities^{1–6}. However, sponges are predicted to be ‘winners’ in future coral reefs and, together with macroalgae, they could replace corals under a changing environment⁷. It is becoming increasingly likely that some sponges could replace corals to create sponge-dominated reefs. Changes from coral- to sponge-dominated reefs are reported in Caribbean, Atlantic, Indo-Pacific, and Pacific reefs. In Wakatobi Marine National Park, Sulawesi, Indonesia, coral coverage decreases with increasing sponges⁸. Sponge-eating organisms (spongivores) include a variety of marine species, including vertebrates such as fish and turtles; mollusks such as opisthobranchs/snails; echinoderms such as asteroids; crustaceans such as crabs, and shrimps^{9,10}.

The coral-killing sponge, *Terpios hoshinota* Rützler & Muzik, 1993, is prevalent in many areas, including Guam^{11,12}, Japan^{13–16}, Taiwan^{17,18}, the Great Barrier Reef, Australia¹⁹, Yongxing Island, China²⁰, Malaysia²¹, Indonesia^{22,23}, Maldives²⁴, and Mauritius²⁵.

Terpios hoshinota is a thin (<1 mm thick), encrusting demosponge with numerous symbiotic cyanobacteria in its tissues. It grows rapidly on live coral at the rate of 1 mm per day (linear progression rate of 11.5–23.0 mm month⁻¹) in tropical sites^{11,12,25,26}. Information about its prevalence is accumulating; however, the information on the relationships between the sponge and associated species is poor. The aim of this study was to describe the snails found for the first time on *T. hoshinota*, their sites of occurrence on sponges, and to examine their relationship with *T. hoshinota*, and identify the snail using molecular DNA barcoding techniques (Figs. 1, 2, 3, 4).

Results

In this study, snails crawling on the *T. hoshinota* sponge, which were overgrowing the branching coral *Montipora digitata*, were obtained from two sites around Okinawa Island (Nakijin and Odo). The material was inspected in the marine laboratory and live veliger larvae were collected from a sponge from Nakijin together with sponge larvae using a cup with nylon net (mesh size: 100 µm). Direct observation in the field and sampling were attempted in Odo, Sesoko, Nakijin, and Ogimi (> 40 branches in each site), but we failed to collect snails. This could be

¹Sesoko Station, Tropical Biosphere Research Center, University of the Ryukyus, Sesoko 3422, Motobu-cho, Okinawa 905-0227, Japan. ²Atmosphere and Ocean Research Institute, The University of Tokyo, Kashiwa, Chiba, Japan. ³Graduate School of Engineering and Science, University of the Ryukyus, Nishihara, Okinawa, Japan. ⁴Environmental Partnership Council, Tokyo, Japan. ✉email: hyama@lab.u-ryukyu.ac.jp

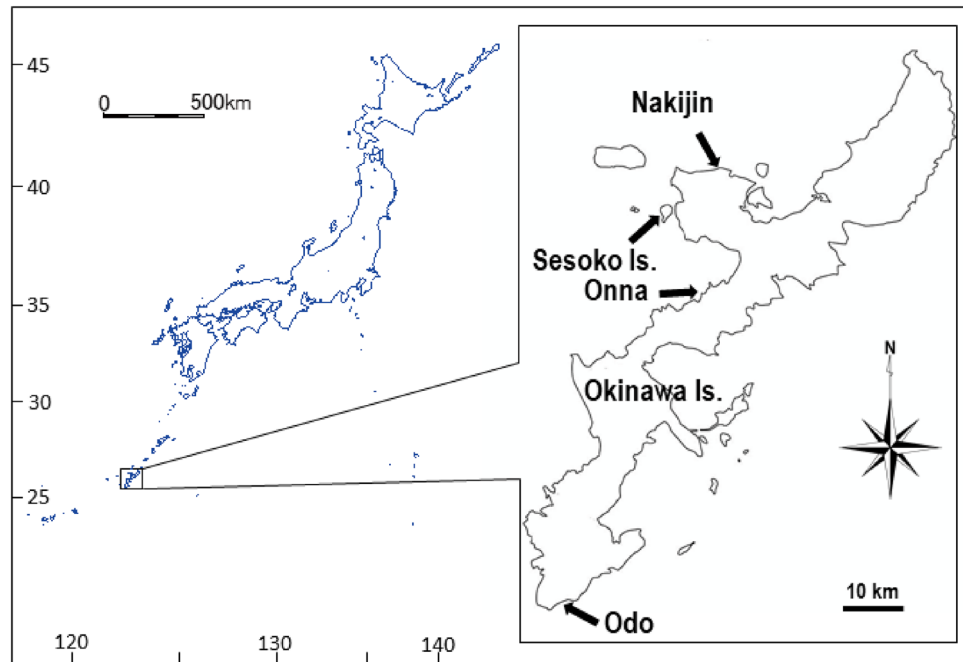


Figure 1. Map showing study sites. The marine station (Sesoko Station) of the University of the Ryukyus is located on Sesoko Island. The software used to create the map was HiMapMeister ver. 1.1.1, Teikoku-Shoin Co., Ltd, https://www.teikokushoin.co.jp/support/index_01.html.

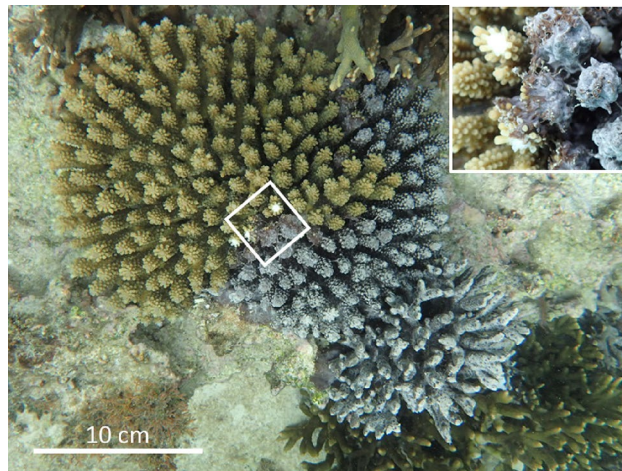


Figure 2. *Terpios hoshinota* covering coral colonies of *Acropora tenuis* and *Montipora digitata* on the Nakijin reef. Inset shows the sponge extending over branches of *Acropora* with thread-like tissues.

attributed to their small size (<2.5 mm in shell height, Fig. 3a–c) and dark coloration. Egg capsules with veliger larvae were found in the histological sections of the specimens from Sesoko Island (January 22, July 4, 2020) and from Onna (August 4, 2020). Cerithiopsidae (Fig. 3a–c) and Triphoridae (d: *Coriophora fusca*, e: *Euthymella elegans*) snails on *T. hoshinotae* were collected from August to November 2020. The mating behavior of the two snails was observed twice in September (Suppl. Movie 1), from the snails on the Nakijin sponge's surface and it continued even when the snails were moved to a Petri dish (Fig. 5a).

Live egg capsules were found for the first time on July 24, 2020, from the Nakijin sample. Egg capsules at the stage of nearly releasing veliger larvae were visible as swollen bumps near the sponge surface (Fig. 6a, suppl. movie 2), and their size was similar to that of sponge larvae. The position of egg capsules was consistent with that of the coral calice. On the day of hatching, the egg capsules became swollen, and larvae became visible through the capsule membrane with decreasing density of sand particles trapped by the sponge. The larvae swam actively inside the capsule and then hatched, swimming out of the capsule (Fig. 6b–f). The exact time of release was observed only once in the aquarium around 8 pm on December 10 (Fig. 6e). The mean number of

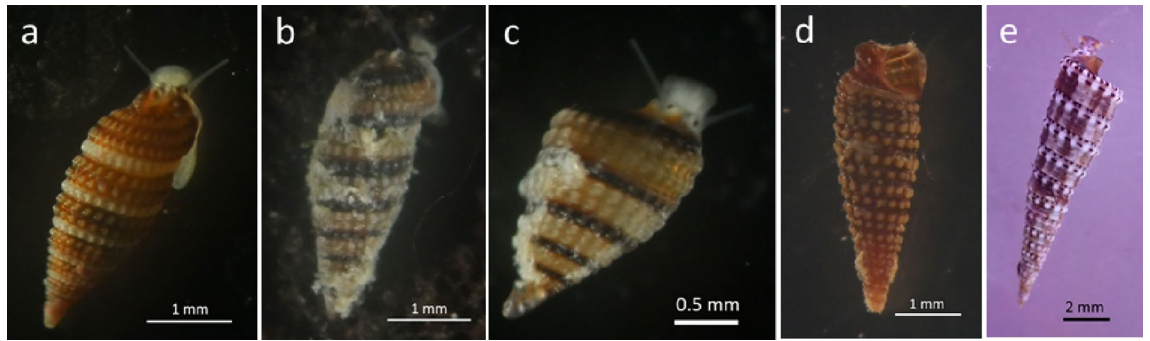


Figure 3. Snails collected from *Terpios hoshinota* sponge. The number of each snail collected during the study period were: (a) (6), (b,c) (3; c is a juvenile), (d) (1, preserved in ethanol), (e) (1).

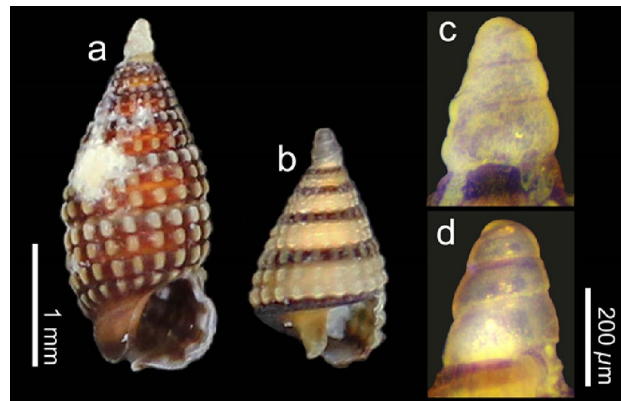


Figure 4. Sequenced specimens of cerithiopsid snails examined in this study. (a,b) shell. (c,d) protoconch. (a,c) *Joculator* sp. (DNA accession no. LC598716); (b,d) *Joculator* sp. (LC598717). The specimens preserved in ethanol were photographed using a microscope (LW-820T, Wraymer Inc., Japan) equipped with a digitalized camera (WRAYCAM-NOA630B, Wraymer Inc., Japan). Scale bars indicate 1 mm for (a) and (b); and 200 μm for (c) and (d).

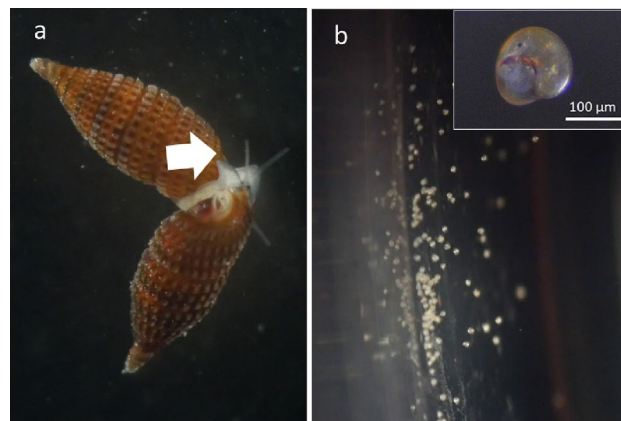


Figure 5. (a) Two snails showing mating behavior, separated from sponge, in a Petri dish. (b) Swimming veliger larvae toward the left (bright side) in a Petri dish. Inset shows the magnified image of a veliger larva on a slide glass.

veliger larvae per egg capsule was 111.7 ± 17.3 (mean \pm SD; range 83–132, $n = 6$), calculated using ethanol-fixed egg capsules. The shell length of veliger was 138.6 ± 6.0 μm (mean \pm SD; range 127.3–151.5 μm , $n = 51$). After hatching from the egg capsule, veliger larvae started to swim and showed strong positive phototaxis toward light (Fig. 5b, suppl. movie 3). We attempted to culture the larvae in a Petri dish with filtered seawater (< 0.45 μm), but they survived only a few days.

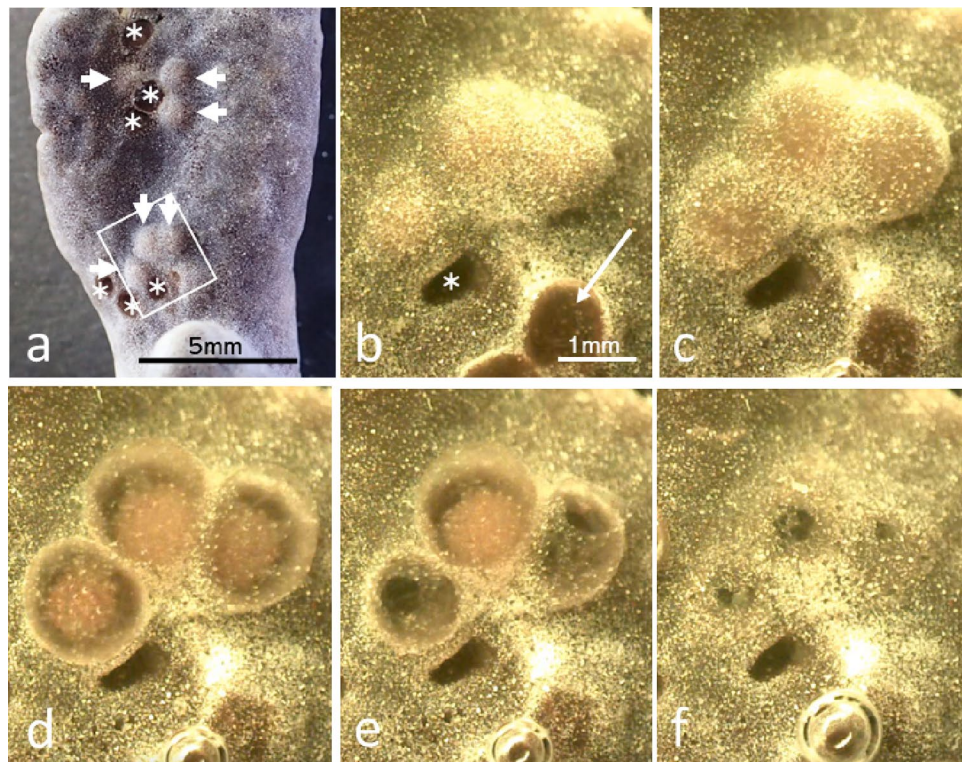


Figure 6. Egg capsules in the tissue of *Terpios hoshinota*. (a) Sponge covering the coral skeleton of *Montipora digitata*. Arrow heads: egg capsules, asterisk: holes after releasing veliger larvae. Photograph was taken with a dissecting microscope (SMZ-1000, Nikon Co., Japan). (b–f) Sequential pictures of egg capsules. Times of (b) to (f) (13:30, 17:00, 19:50, 20:00, 24:00 on December 10, 2020). The arrow in (b) shows the scar of a bubble released from sponge tissue. Each panel was captured from a time-lapse video using a digital microscope (Dino-Lite Premier, AnMo Elec. Co.).

Egg capsules were found in the histological sections initially prepared for observing sponge reproduction. The sponges containing egg capsules were observed in the samples obtained from Sesoko Is. on July 4 and from Onna on August 4, 2020. Figure 7 shows many egg capsules laid deep into the tissue of *Terpios hoshinota*, and the sizes of the egg capsules (1.2 mm in diameter) were close to that of sponge larvae (Fig. 7b).

Molecular analyses based on COI gene sequences indicated that each snail of *a* and *b* types (Fig. 3) and the veliger larvae are the same species, *Joculator* sp. In the phylogenetic tree (Fig. 8), veliger larvae were included in a monophyletic clade with *Joculator* sp. supported by high bootstrap values (100%). In addition, low levels of genetic divergence, ranging from 0.6 to 1.5%, were observed between the two snail specimens identified as *Joculator* sp. and the veliger larvae. These values for the COI sequences of *Joculator* sp. were similar to the range of intraspecific divergences for the other cerithiopsid species (0.0 to 2.8%⁴⁰).

Discussion

In this study, we examined the snails associated with the sponge *Terpios hoshinota* for the first time. The number of snails observed in this study was small (<6 individuals per species, Fig. 3); however, sponge-associated snails may be distributed widely, because snails, egg capsules, and veliger larvae were found at four Okinawa Island sites. *Terpios*-affected islands are abundant along the Ryukyu Archipelago¹⁵. Egg capsules and veliger larvae were observed between July and December in the present study, indicating that their reproductive season lasts for at least six months, from summer to fall.

Spongivores (sponge-eating organisms) include various animals, such as nudibranchs, snails, echinoids, fish, and turtles^{9,10,26,28}. Relatively large (5–20 cm in length) dorid nudibranchs consume *Terpios* sp. in the northeastern Pacific^{27,29}. *Terpios hoshinota* is a spiculate demosponge¹³ and has a cytotoxic compound³⁰, therefore, this sponge is not palatable for predators. In addition, the sponge spicules (ca. 200 μ m long) and particles on the surface of *T. hoshinota* tissues act as barriers to predators. However, this sponge armored with spicules, particles, and toxic substances would be a relatively safe place for snail larvae to lay their egg capsules. This study did not determine the direct evidence of the snails feeding on sponge tissues; however, there is a possibility that, like other cerithiopsids, these snails use sponges as a food source via excavation of soft tissue using their proboscises^{31,32}.

In this study, we collected three different snail species from the surface of *Terpios hoshinota*. The number of snails was small; however, more intensive and quantitative surveys could find more sponge-associated snails, from the widely distributed *Terpios* in southern Japan. Therefore, survey of areas containing sponge-affected reefs along the Ryukyu Archipelago is required. It is possible that even if the sponge-associated snails consume

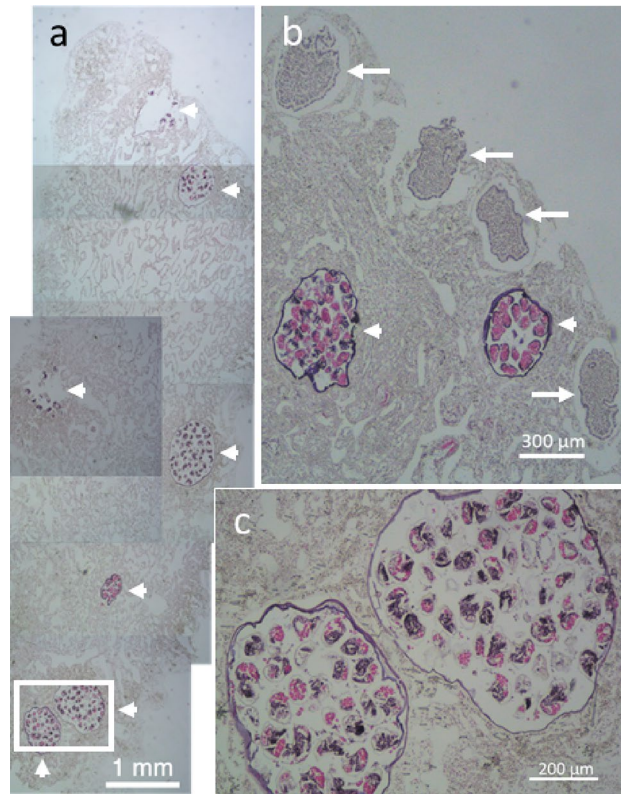


Figure 7. Combined histological pictures of *Terpios hoshinota* tissue. (a) cross section of *Montipora digitata* branch covered by *T. hoshinota* (collected from Onna, August 4, 2020). Arrow heads show egg capsules of the snail. (b) Sponge tissue from the specimen from Sesoko Is. (collected July 4, 2020). Arrows show sponge larvae.

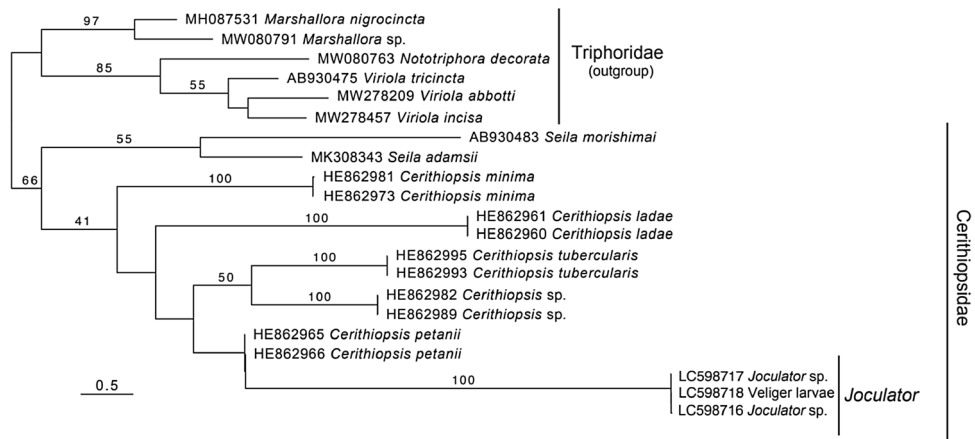


Figure 8. Maximum-likelihood phylogenetic tree of the family Cerithiopsidae reconstructed using the COI sequences (621 bp). Snails and veliger larvae were obtained from the sponge *Terpios hoshinota*. Bootstrap probability values for each node below 40% are not shown. Scale bar represents branch length (substitutions/site).

sponges, they are unable to alter the growth of the sponge significantly, owing to their small size. Therefore, the snails are unlikely to be candidate biological control agents for inhibiting the spread of the coral-killing sponge *Terpios hoshinota*. However, studies on the species composition, geological distribution, and abundance of associates, including snails, would reveal a new view of the coral-killing sponge *Terpios hoshinota* as a host organism.

Materials and methods

Study sites. The study sites where the sponge-associated snails or the veliger larvae were observed include Sesoko Island (26°39'07.82" N, 127° 51'23.26" E), Nakijin (26°42'30.9" N, 127°56'59.2" E), Onna (26°31'39.52" N, 127° 55'14.99" E) and Odo (26°05'20.10" N, 127°42'31.07" E), all around Okinawa Island, Japan (Fig. 1). At all sites, dense aggregations of branching *Montipora* corals had developed in a shallow moat (maximum depth 2 m) together with massive *Porites* spp., foliose *M. aequituberculata*, leafy *Pavona frondifera*, corymbose *Acropora* spp., and other scleractinians. Some of these hard corals were fully or partly covered by *Terpios hoshinota* (Fig. 2). We collected the snails during the regular monthly sampling in Sesoko Island and Nakijin, during reproductive studies of the sponge, as well as from other sites where snail or veliger larvae were observed.

Collection of snails and veliger larvae. The small size and dark color of the snails made it difficult to find them on the black sponge in the field. Most snails were found during close observation using a dissecting microscope. Veliger larvae released from the sponge were trapped in a filter cup (100 µm nylon mesh filter, cell strainer, BD Biosciences Discovery Labware) together with sponge larvae. The histological observations for the reproductive studies in sponge tissues were conducted as follows: the tissues were fixed with 10% formalin solution, dehydrated with a graded series of ethanol, embedded in paraffin, and stained with hematoxylin/eosin dyes. Presence/absence of snail egg capsules in the sponge tissue were recorded.

The snails, egg capsules, and veliger larvae were observed using a light microscope (Eclipse Ci, Nikon Co.), a dissecting microscope (SMZ-1000, Nikon Co.), and a digital microscope (Dino-Lite Premier, AnMo Elec. Co.) to obtain time-lapse images. The snails were observed in the field on the collection day; other observations and culture experiments were performed in the marine laboratory at Sesoko Station, Tropical Biosphere Research Center, University of the Ryukyus.

Molecular identification of snails and veliger. We could collect multiple samples of only two morphological types of snails. The shell of *a* type was brown (Fig. 3a), and that of *b* type was sandy-yellow with a dark red-brown suture (Fig. 3b,c). Two cerithiopsid snails (one specimen each of *a* and *b* types; Fig. 4a,b for suture, c and d for protoconch, respectively), collected from Odo in October 2020, and approximately 60 individuals of unidentified veliger larvae were fixed and preserved in pure ethanol for morphological and molecular identification. Cerithiopsid snails were identified to the genus level based on shell morphology, as described previously^{31,33–36}. In addition to the visual morphological identification, molecular identification was performed using cytochrome *c* oxidase subunit I (COI) sequences. The total DNA of snails and veliger larvae was extracted from foot tissue and 20 whole veligers, respectively, using the DNeasy Tissue Extraction Kit (Qiagen). The mitochondrial COI sequences (658 bp) were amplified through polymerase chain reaction (PCR) using the primer pairs LCO1490 and HCO2198³⁷, following the conditions described earlier³⁸. The PCR products were visualized through electrophoresis on a 1.5% Tris–Borate–EDTA agarose gel and purified with ExoSAP-IT (Thermo Fisher Scientific). The purified products were Sanger sequenced in both directions using an ABI 3730xl Genetic Analyzer (Applied Biosystems) at Eurofins Genomics (Tokyo, Japan). The COI sequences were manually aligned using Mesquite version 3.61^{38,39} and compared with previously reported sequences of cerithiopsid species^{40,41}. Genetic divergences among the sequences were quantified using the Kimura 2-Parameter (K2P) distance model⁴² using MEGA X version 10.1.7⁴³. Phylogenetic relationships of cerithiopsid species were reconstructed from COI sequences (621 bp) using the maximum-likelihood (ML) methods. The ML tree reconstruction was performed under GTR+G model in RAxML v.7.4.2⁴⁴ with a bootstrap analysis of 1,000 pseudoreplicates. Nucleotide sequences were deposited in the DNA Data Bank of Japan (DDBJ) under the accession numbers LC598716–LC598717 for snails and LC598718 for veliger larvae. The sequenced specimens were deposited as a voucher (specimen number: 20210831-HF010-12) in the Atmosphere and Ocean Research Institute (AORI), The University of Tokyo (<https://www.aori.u-tokyo.ac.jp>, contact person: Hiroaki Fukumori, fukumori@aori.u-tokyo.ac.jp).

Sampling and field studies. All necessary permits for sampling and observational field studies were obtained from the concerned authorities. Coral sampling was performed with approval from the authorities of Okinawa Prefecture, Japan.

Data availability

The datasets generated or analyzed during the current study are available from the corresponding author upon reasonable request.

Received: 10 February 2021; Accepted: 7 October 2021

Published online: 20 October 2021

References

- Hughes, T. P. *et al.* Climate change, human impacts, and the resilience of coral reefs. *Science* **301**, 929–933. <https://doi.org/10.1126/science.1085046> (2003).
- Hoegh-Guldberg, O. *et al.* Coral reefs under rapid climate change and ocean acidification. *Science* **318**, 1737. <https://doi.org/10.1126/science.1152509> (2007).
- Sokolow, S. Effects of a changing climate on the dynamics of coral infectious disease: A review of the evidence. *Dis. Aquat. Org.* **87**, 5–18. <https://doi.org/10.3354/dao02099> (2009).
- De'ath, G., Fabricius, K. E., Sweatman, H. & Puotinen, M. The 27-year decline of coral cover on the Great Barrier Reef and its causes. *Proc. Natl. Acad. Sci. USA* **109**, 17995–17999. <https://doi.org/10.1073/pnas.1208909109> (2012).

5. Hughes, T. P. *et al.* Global warming and recurrent mass bleaching of corals. *Nature* **543**, 373–377. <https://doi.org/10.1038/nature21707> (2017).
6. May, L. A. *et al.* Effect of Louisiana sweet crude oil on a Pacific coral, *Pocillopora damicornis*. *Aquat. Toxicol.* **28**, 105454. <https://doi.org/10.1016/j.aquatox.2020.105454> (2020).
7. Bell, J. J., Davy, S. K., Jones, T., Taylor, M. W. & Webster, N. S. Could some coral reefs become sponge reefs as our climate changes?. *Glob. Change. Biol.* **19**, 2613–2624. <https://doi.org/10.1111/gcb.12212> (2013).
8. Bell, J. J. & Smith, D. Ecology of sponge assemblages (Porifera) in the Wakatobi region, south-east Sulawesi, Indonesia: Richness and abundance. *J. Mar. Biol. Assoc. UK* **84**, 581–591. <https://doi.org/10.1017/S0025315404009580h> (2004).
9. Wulff, J. L. Ecological interactions of marine sponges. *Can. J. Zool.* **84**, 146–166. <https://doi.org/10.1139/z06-019> (2006).
10. Wooster, M. K., Marty, M. J. & Pawlik, J. R. Defense by association: Sponge-eating fishes alter the small-scale distribution of Caribbean reef sponges. *Mar. Ecol.* **38**, e12410. <https://doi.org/10.1111/maec.12410> (2017).
11. Bryan, P. G. Growth rate, toxicity, and distribution of the encrusting sponge *Terpios* sp. (Hadromerida: Suberitidae) in Guam, Mariana Islands. *Micronesica* **9**, 237–242 (1973).
12. Plucer-Rosario, G. The effect of substratum on the growth of *Terpios*, an encrusting sponge which kills corals. *Coral Reefs* **5**, 197–200. <https://doi.org/10.1007/BF00300963> (1987).
13. Rützler, K. & Muzik, K. *Terpios hoshinota*, a new cyanobacteriosponge threatening Pacific reefs. *Sci. Mar.* **57**, 395–403. e0120853 (1993).
14. Reimer, J. D., Nozawa, Y. & Hirose, E. Domination and disappearance of the black sponge: A quarter century after the initial *Terpios* outbreak in Southern Japan. *Zool. Stud.* **50**, 394 (2010).
15. Reimer, J. D., Mizuyama, M., Nakano, M., Fujii, T. & Hirose, E. Current status of the distribution of the coral-encrusting cyanobacteriosponge *Terpios hoshinota* in southern Japan. *Galaxea J. Coral Reef Stud.* **13**, 35–44. <https://doi.org/10.3755/galaxea.13.35> (2011).
16. Yomogida, M., Mizuyama, M., Kubomura, T. & Reimer, J. D. Disappearance and return of an outbreak of the coral-killing cyanobacteriosponge *Terpios hoshinota* in Southern Japan. *Zool. Stud.* **56**, 1–7. <https://doi.org/10.6620/ZS.2017.56-07> (2017).
17. Liao, M.-H. *et al.* The “black disease” of reef-building corals at Great Island, Taiwan outbreak of a cyanobacteriosponge *Terpios hoshinota* (Suberitidae; Hadromerida). *Zool. Stud.* **46**, 520 (2007).
18. Nozawa, Y., Huang, Y. S. & Hirose, E. Seasonality and lunar periodicity in the sexual reproduction of the coral-killing sponge, *Terpios hoshinota*. *Coral Reefs* **35**, 1071–1081. <https://doi.org/10.1007/s00338-016-1417-0> (2016).
19. Fujii, T. *et al.* Coral-killing cyanobacteriosponge (*Terpios hoshinota*) on the Great Barrier Reef. *Coral Reefs* **30**, 483. <https://doi.org/10.1007/s00338-011-0734-6> (2011).
20. Shi, Q., Liu, G. H., Yan, H. Q. & Zhang, H. L. Black disease (*Terpios hoshinota*): A probable cause for the rapid coral mortality at the northern reef of Yongxing Island in the South China Sea. *Ambio* **41**, 446–455. <https://doi.org/10.1007/s13280-011-0245-2> (2012).
21. Hoeksema, B. W., Waheed, Z. & de Voogd, N. J. Partial mortality in corals overgrown by the sponge *Terpios hoshinota* at Tioman Island, Peninsular Malaysia (South China Sea). *Bull. Mar. Sci.* **90**, 989–990. <https://doi.org/10.5343/bms.2014.1047> (2014).
22. Van der Ent, E., Hoeksema, B. W. & de Voogd, N. J. Abundance and genetic variation of the coral-killing cyanobacteriosponge *Terpios hoshinota* in the Spermonde Archipelago, SW Sulawesi, Indonesia. *J. Mar. Biol. Assoc. UK* **96**, 453–463. <https://doi.org/10.1017/S002531541500034X> (2015).
23. Madduppa, H., Schupp, P. J., Faisal, M. R., Sastrira, M. Y. & Thoms, C. Persistent outbreaks of the “black disease” sponge *Terpios hoshinota* in Indonesian coral reefs. *Mar. Biodivers.* **47**, 149–151. <https://doi.org/10.1007/s12526-015-0426-5> (2017).
24. Montano, S., Chou, W.-H., Chen, C. A., Galli, P. & Reimer, J. D. First record of the coral-killing sponge *Terpios hoshinota* in the Maldives and Indian Ocean. *Bull. Mar. Sci.* **91**, 97–98. <https://doi.org/10.5343/bms.2014.1054> (2015).
25. Elliott, J. B., Patterson, M., Vitry, E., Summers, N. & Miternique, C. Morphological plasticity allows coral to actively overgrow the aggressive sponge *Terpios hoshinota* (Mauritius, Southwestern Indian Ocean). *Mar. Biodivers.* **46**, 489–493. <https://doi.org/10.1007/s12526-015-0370-4> (2016).
26. Thinesh, T., Mathews, G., Raj, K. D. & Edward, J. K. P. Outbreaks of *Acropora* white syndrome and *Terpios* sponge overgrowth combined with coral mortality in Palk Bay, southeast coast of India. *Dis. Aquat. Org.* **126**, 63–70. <https://doi.org/10.3354/dao03155> (2017).
27. Birenheide, R., Amemiya, S. & Motokawa, T. Penetration and storage of sponge spicules in tissues and coelom of spongivorous echinoids. *Mar. Biol.* **115**, 677–683. <https://doi.org/10.1007/BF00349376> (1993).
28. Vicente, J., Osberg, A., Marty, M. J., Rice, K. & Toonen, R. J. Influence of palatability on the feeding preferences of the endemic Hawaiian tiger cowrie for indigenous and introduced sponges. *Mar. Ecol. Prog. Ser.* **647**, 109–122. <https://doi.org/10.3354/meps13418> (2020).
29. Penney, B. K. How specialized are the diets of northeastern Pacific sponge-eating dorid nudibranchs?. *J. Moll. Stud.* **79**, 64–73. <https://doi.org/10.1093/mollus/ey038> (2013).
30. Teruya, T. *et al.* Nakiterpiosin and nakiterpiosinone, novel cytotoxic C-nor-D-homosteroids from the Okinawan sponge *Terpios hoshinota*. *Tetrahedron* **60**, 6989–6993. <https://doi.org/10.1016/j.tet.2003.08.083> (2004).
31. Marshall, B. A. Cerithiopsidae (Mollusca: Gastropoda) of New Zealand, and a provisional classification of the family. *New Zeal. J. Zool.* **5**, 47–120. <https://doi.org/10.1080/03014223.1978.10423744> (1978).
32. Collin, R. Development of *Cerithiopsis gemmulosum* (Gastropoda: Cerithiopsidae) from Bocas del Toro, Panama. *Caribb. J. Sci.* **40**, 192–197 (2004).
33. Cecalupo, A. & Perugia, I. Cerithiopsidae and Newtoniellidae (Gastropoda: Triphoroidea) from New Caledonia, western Pacific. *Visaya Suppl.* **7**, 1–175 (2016).
34. Cecalupo, A. & Perugia, I. Cerithiopsidae. In *Philippine Marine Mollusks* Vol. V (ed. Poppe, G.) 1352–1375 (Conchbooks, 2017).
35. Cecalupo, A. & Perugia, I. New species of Cerithiopsidae (Gastropoda: Triphoroidea) from Papua New Guinea (Pacific Ocean). *Visaya Suppl.* **11**, 1–187 (2018).
36. Cecalupo, A. & Perugia, I. New species of Cerithiopsidae and Newtoniellidae from Okinawa (Japan-Pacific Ocean). *Visaya Suppl.* **12**, 1–84 (2019).
37. Folmer, O., Black, M., Hoeh, W., Lutz, R. & Vrijenhoek, R. DNA primers for amplification of mitochondrial cytochrome *c* oxidase subunit I from diverse metazoan invertebrates. *Mol. Mar. Biol. Biotech.* **3**, 294–299 (1994).
38. Kano, Y. & Fukumori, H. Predation on hardest molluscan eggs by confamilial snails (Neritidae) and its potential significance in egg-laying site selection. *J. Moll. Stud.* **76**, 360–366. <https://doi.org/10.1093/mollus/eyq018> (2010).
39. Maddison, W. P. & Maddison, D. R. Mesquite: a modular system for evolutionary analysis. Version 3.61. <http://www.mesquiteproject.org> (2019).
40. Modica, M. V., Mariottini, P., Prkić, J. & Oliverio, M. DNA-barcoding of sympatric species of ectoparasitic gastropods of the genus *Cerithiopsis* (Mollusca: Gastropoda: Cerithiopsidae) from Croatia. *J. Mar. Biol. Assoc. UK* **93**, 1059–1065. <https://doi.org/10.1017/S0025315412000926> (2012).
41. Takano, T. & Kano, Y. Molecular phylogenetic investigations of the relationships of the echinoderm-parasite family Eulimidae within Hypsogastropoda (Mollusca). *Mol. Phylogenet. Evol.* **79**, 258–269. <https://doi.org/10.1016/j.ympev.2014.06.021> (2014).
42. Kimura, M. A. Simple method for estimating evolutionary rate of base substitutions through comparative studies of nucleotide sequence. *J. Mol. Evol.* **16**, 111–120. <https://doi.org/10.1007/BF01731581> (1980).

43. Kumar, S., Stecher, G., Li, M., Knyaz, C. & Tamura, K. MEGA X: Molecular evolutionary genetics analysis across computing platforms. *Mol. Biol. Evol.* **35**, 1547–1549. <https://doi.org/10.1093/molbev/msy096> (2018).
44. Stamatakis, A. RAxML-VI-HPC: Maximum likelihood-based phylogenetic analyses with thousands of taxa and mixed models. *Bioinformatics* **22**, 2688–2690. <https://doi.org/10.1093/bioinformatics/btl446> (2006).

Acknowledgements

We are grateful to Dr. T. Kurozumi of Chiba Prefectural Museum and Dr. T. Takano of Meguro Parasitological Museum for their valuable comments on the snails. We are also thankful to J.D. Reimer of University of the Ryukyus and an anonymous reviewer, for their invaluable comments and suggestions to improve the manuscript. We thank the staff of the Sesoko Station, Tropical Biosphere Research Center, University of the Ryukyus for their kind support.

Author contributions

All authors contributed to the conception and design of the study. Sampling was performed by Y.H., S.N.A., and H.Y. Y.H. found most snails and observed the behavior of the snails. Molecular analysis was performed using H.F., and S.N.A. prepared histological sections of the sponge. All authors read and approved the final manuscript.

Funding

This work was supported by a grant from the Japan Society for the Promotion of Science (JSPS) KAKENHI (Grant no. 19K06091), and Halekulani Okinawa.

Competing interests

The authors declare no competing interests.

Additional information

Supplementary Information The online version contains supplementary material available at <https://doi.org/10.1038/s41598-021-00185-x>.

Correspondence and requests for materials should be addressed to H.Y.

Reprints and permissions information is available at www.nature.com/reprints.

Publisher's note Springer Nature remains neutral with regard to jurisdictional claims in published maps and institutional affiliations.



Open Access This article is licensed under a Creative Commons Attribution 4.0 International License, which permits use, sharing, adaptation, distribution and reproduction in any medium or format, as long as you give appropriate credit to the original author(s) and the source, provide a link to the Creative Commons licence, and indicate if changes were made. The images or other third party material in this article are included in the article's Creative Commons licence, unless indicated otherwise in a credit line to the material. If material is not included in the article's Creative Commons licence and your intended use is not permitted by statutory regulation or exceeds the permitted use, you will need to obtain permission directly from the copyright holder. To view a copy of this licence, visit <http://creativecommons.org/licenses/by/4.0/>.

© The Author(s) 2021