Reproduction in deep-sea vent shrimps is shaped by diet and phylogeny, with rhythms unlinked to surface production

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Running head: Reproductive strategies in *Rimicaris* vent shrimps

Abstract

Variations in offspring production according to feeding strategies or food supply have been recognized in many animals from various ecosystems. Despite an unusual trophic structure based on non-photosynthetic primary production, these relationships remain largely under-studied in chemosynthetic ecosystems. Here, we use *Rimicaris* shrimps from deep-sea hydrothermal vents as a study case to explore relations between reproduction, diets and food supply in these environments. For that, we compared reproductive outputs of three congeneric shrimps differing by their diets. They inhabit vents located under oligotrophic waters of tropical gyres with opposed latitudes, allowing us to also examine the prevalence of phylogenetic vs environmental drivers in their reproductive rhythms. For this we used both our original data and a compilation of published observations on the presence of ovigerous females covering various seasons over the past 35 years. We report distinct egg production trends between

Rimicaris species relying solely on chemosymbiosis – *R. exoculata* and *R. kairei* – and those
relying on mixotrophy – *R. chacei* – where *R. chacei* produces broods with higher numbers of
smaller eggs. Besides, our data and historical records suggest a reproductive period with
substantial proportions of brooding females mostly between January and early April for all
examined species whatever the region. Intriguingly, this periodicity does not correspond to
seasonal variations in surface production with presence of brooding females during either
boreal winter or austral summer. These observations contrast with the long-standing paradigm
in deep-sea species for which periodic reproductive patterns have always been attributed to
seasonal variations of photosynthetic production sinking from surface. Our results suggest the
presence of intrinsic basis for biological rhythms in the deep sea, and bring to light the
importance of having year-round observations in order to understand life history of vent
animals.

Keywords

Biological rhythms, crustacean, deep sea, hydrothermal vent, reproduction, seasonality, trophic
ecology

44   **Introduction**

Relationships between feeding strategies and offspring's production have been
extensively studied for a number of animal phyla in various ecosystems (Pierotti and Annett
1990, Qian and Chia 1991, Broderick et al. 2003, Meiri et al. 2012, Sibly et al. 2012). Where no
clear trends between trophic regime and reproductive outputs could be observed in lizards
(Meiri et al. 2012), large variations of fecundity or reproductive outputs according to diets have
been highlighted in others including birds (Pierotti and Annett 1990, Sibly et al. 2012), sea
turtles (Broderick et al. 2003), annelids (Qian and Chia 1991), insects (Gutiérrez et al. 2020), or
crustaceans (Lin and Shi 2002, Tziouveli et al. 2011, Almeida et al. 2018). All these studies are
based on ecosystems where photosynthetic primary production is the sole basis of food webs.
A number of ecosystems on Earth, such as deep-sea hydrothermal vents, cold seeps and
organic falls, rely on another primary organic source: the chemosynthesis (German et al. 2011).
In chemosynthetic-based ecosystems lush and dense faunal communities are supported by

microbial communities using chemical energy arising from geofluids to produce their organic
matter, and animals there often form intricate symbiotic associations with these
chemosynthetic bacteria for nutrition (Dubilier et al. 2008). From total reliance on symbiosis to
bacterivory to mixotrophy to carnivorous, these particular ecosystems display a variety of diets
in networks often differing strikingly from those found elsewhere (Govenar 2012). In this
context, these environments are natural laboratories to study relationships between
reproduction and trophic regimes. However, no study has yet explored these relationships
extensively in chemosynthetic-based ecosystems, with variations in the reproductive output
among or within animal groups being mostly attributed to phylogenetic constraints (Tyler and
Young 1999) or variations in female body size (Ramirez-Llodra et al. 2000, Copley and Young
2006, Nye et al. 2013, Marsh et al. 2015, Marticorena et al. 2020).

Although most species from deep-sea chemosynthetic-based ecosystems exhibit
continuous reproductive patterns (Jollivet et al. 2000, Faure et al. 2007, Tyler et al. 2008, Hilário
et al. 2009, Marticorena et al. 2020), seasonal cycles following variations in surface primary
production have been observed in many taxa such as bythograeid crabs, alvinocaridid shrimps,
and bathymodiolin mussels (Perovich et al. 2003, Dixon et al. 2006, Copley and Young 2006,
Tyler et al. 2007). In some groups, the surface production pattern is clearly linked to
reproductive patterns in the deep. For example, some vent crab species exhibit continuous
gametogenesis under the highly oligotrophic waters of the South Pacific Gyre, while others
living in more productive areas north of this gyre have seasonal reproduction (Perovich et al.
2003, Hilário et al. 2009). Such examples have led to the paradigm that rhythms of sinking
photosynthetic matter is the key time-synchronising cue for deep-water animals with seasonal
reproduction, signalling individuals to spawn at the same period of the year.

The symbiotic and sister-species shrimps *Rimicaris exoculata* and *R. kairei* are among the
most iconic hot-vent animals, with unique adaptations such as hosting symbiotic bacteria in
their enlarged head cavity and the fused "dorsal eye", a novel sensory system representing a
pinnacle of adaptation to vents (Zbinden and Cambon-Bonavita 2020). The entire family they
belong to, Alvinocarididae, is endemic to vents and seeps with different taxa exhibiting various
trophic regimes and various levels of reliance on symbiosis (Rieley et al. 1999, Gebruk et al.

2000, Apremont et al. 2018), constituting an ideal case for elucidating relationships between
reproduction and diet in chemosynthesis-based ecosystems. Species with different trophic
regimes often co-occur on same vent fields, for example *R. exoculata* which rely on symbiosis
and the mixotrophic *R. chacei*, allowing for a straight-forward comparison of species. The
symbiotrophic species such as *R. exoculata* and *R. kairei* exhibit greatly inflated cephalothorax
and live in large aggregations around the hot fluid flows (Rieley et al. 1999, Gebruk et al. 2000,
Van Dover 2002, Streit et al. 2015, Methou et al. 2020), whereas mixotrophic species such as *R.*
chacei (Gebruk et al. 2000, Apremont et al. 2018, Methou et al. 2020) lacks the inflated
cephalothorax and lives in small groups (Methou et al. 2021).

Attempts to understand the reproduction of *Rimicaris* shrimps have so far faced
contradicting evidences. While continuous reproduction was initially suggested from
examination of their reproductive tissues (Ramirez-Llodra et al. 2000, Copley et al. 2007) hardly
any ovigerous individuals could be collected in over 35 years of seagoing expeditions, despite a
relatively focused and repeated sampling effort (Williams and Rona 1986, Vereshchaka 1997,
Shank et al. 1998, Ramirez-Llodra et al. 2000, Copley et al. 2007, Komai and Segonzac 2008).
Brooding females were only found in large numbers in 2014 (Methou et al. 2019, Hernández-
Ávila et al. 2021) and to a lesser extent in 2007 (Guri et al. 2012), around a restricted period
between January and March. These females were living within dense aggregations close to vent
orifices (Hernández-Ávila et al. 2021), refuting previous hypothesis that brooding females
would migrate to vent periphery to protect their eggs from vent fluid, as observed for
bythograeid crabs (Perovich et al. 2003) as well as *Kiwa* squat lobsters (Marsh et al. 2015).
Taken together, these recent findings suggest a seasonal reproductive cycle possibly linked to
surface primary production, as shown in vent crabs, bathymodiolin mussels and some other
alvinocaridid shrimps (Dixon et al. 2006, Copley and Young 2006, Tyler et al. 2007). However,
the low primary productivity of oligotrophic surface waters and the overall low export to the
deep-sea compartment within the region where these shrimps have been collected (-3500 m
depth) (Pabortsava et al. 2017, Harms et al. 2021) challenges the possibility of such a link.

Here, we compare reproductive outputs – i.e., fecundities and egg volumes – of three
congeneric vent shrimps with contrasting diets, using two species with a full nutritional

dependence on their chemosymbiosis (i.e., *R. exoculata* and *R. kairei*) and one with a
mixotrophic diet combining symbiosis, bacterivory, and scavenging (i.e., *R. chacei*) to find out if
links between diets and reproduction are also present in chemosynthetic based ecosystems.
Distribution of these species in two different hemispheres (Northern Atlantic and Southern
Indian Ocean) with similar levels of, but seasonally opposed, surface production allows us to
evaluate if their brooding periods are in synchrony with patterns of surface primary
productivity.

Materials & Methods

Field sampling

Shrimps were collected from three hydrothermal vent fields (figure 1): TAG
(26°08′12.12′′N, 44°49′36.12′′W, 3620 m depth) and Snake Pit (23°22′5.88′′N 44°57′0′′W, 3470
m depth) on the Mid Atlantic Ridge (MAR) between March and April in 2017 (HERMINE
(doi:10.17600 /17000200)) and between February and March in 2018 (BICOSE2
(doi:10.17600/18000004)) for *R. exoculata* and *R. chacei*, and Kairei (25°19′10.2′′S, 70°2′24′′E,
2415 m depth) on the Central Indian Ridge (CIR) between February and March both in 2002 and
2016, as well as in November 2009 for *R. kairei*. Shrimps were sampled using suction samplers
of the human-occupied vehicle (HOV) *Nautile* or the HOV *Shinkai 6500*. Upon recovery on-
board, ovigerous females of *R. exoculata* and *R. chacei* were identified and individually sorted
from the rest of the population. An additional published dataset for *Rimicaris exoculata*
collected at TAG and Snake Pit in January-February 2014 is also included in our analyses and
available in the Ifremer SEANOE (SEA scieNtific Open data Edition) database at:
https://doi.org/10.17882/84112.

Ovigerous females include both brooding females and females that have just released
their larvae but still retained modified pleopods, characteristic of a recent brooding status,
following Nye at al. 2013 (Nye et al. 2013). For *R. kairei*, ovigerous females were identified and
sorted from the rest of the population in the laboratory after the cruise. Egg stages in broods
were identified on-board during the BICOSE2 expedition before being fixed in 10% neutral
buffered formalin for 24h and then rinsed and stored in 80% EtOH. For other expeditions,

ovigerous females were fixed and stored in 10% formalin on-board and identification of egg
stages was conducted in shore-based laboratory.

Shrimps and Eggs Measurements

Carapace length (CL) of each female was measured with Vernier callipers from the
posterior margin of the ocular shield (or eye socket) to mid-point of the posterior margin of the
carapace, with an estimated precision of 0.1 mm. For each brood, total number of eggs was
manually counted. Ten eggs were selected randomly to measure maximum and minimum
diameters. Volume of embryos was estimated following the same method as (Hernández-Ávila
et al. 2021), considering a spheroid volume $v = (4/3).\pi.r1.r2^2$, where r1 and r2 are half of
maximum and minimum axis of each egg, respectively. Eggs in broods were classified in three
developmental stages (early, mid, and late), following the classification used in (Methou et al.
2019). Some ovigerous females harboured damaged or aborted broods and were thus
discarded from our fecundity analysis. Furthermore, we also compiled data for the presence of
ovigerous female shrimps from the literature and using dive videos from cruises over the last 35
years, in order to assess the reproductive seasonality. Proportions of ovigerous females within
samples were calculated against the number of sexually mature females (See supplementary
material). Dataset compiling size, reproductive status, fecundities, egg developmental stages,
and egg diameters of each shrimp individual is available in the Ifremer SEANOE (SEA scieNtific
Open data Edition) database at: https://doi.org/10.17882/84195.

Statistical analysis

Brooding females were grouped according to sampling year, species and vent field.
Visual examination of our dataset and Shapiro-Wilk normality tests revealed that size, egg
number, and egg volume of *Rimicaris* brooding females deviated from normal distribution.
Therefore, non-parametric tests were used for intergroup comparison, with a Kruskal-Wallis
test followed by post-hoc Dunn tests when three or more groups were compared. Spearman
rank-order correlation were used to assess relationship between $\log_e$-transformed realized
fecundity and $\log_e$-transformed carapace length of the different shrimp groups. Egg stage

proportions between groups were compared using χ^2 test with Yate's correction. All analyses
were performed in R v. 4.0.3 (R Core Team 2020).

Results

Interannual variation of *R. exoculata* reproduction

Realized fecundity of *Rimicaris exoculata* correlated positively with carapace length in
both 2014 (Spearman correlation r= 0.67, $p < 0.0001$) and 2018 (Spearman correlation r= 0.73,
$p < 0.0001$) (figure 2a). Moreover, slopes of the relationship between $\log_e$-transformed realized
fecundity and $\log_e$-transformed carapace length did not differ significantly between these two
sampling years (F= 1.175, $p = 0.28$) or between TAG and Snake Pit in 2018 (F= 0.004, p= 0.95).
Although no correlation was found in 2017 (Spearman correlation r= 0.38, p= 0.16), probably
due to the lower sampling effort, realized fecundity in 2017 still fell within the range of the
other sampling years (figure 2a).

Overall, size-specific fecundity ranged from 19 to 95 eggs mm^{-1} in 2014, from 15 to 80
eggs mm^{-1} in 2017 and from 10 to 134 eggs mm^{-1} in 2018 (figure 2b). Limited differences in size-
specific fecundities were observed between different sampling years (Kruskal-Wallis, H= 65.025
$p < 0.05$) with significant variations between 2014 and 2018 at TAG only (2014: 37.3 ± 11.4;
2018: 49.4 ± 12.9; Dunn's Multiple Comparison Test, $p < 0.05$). Differences in size-specific
fecundity were also observed between the two vent fields both in 2014 and in 2018, (Dunn's
Multiple Comparison Tests, $p < 0.05$). Carapace length of brooding females from TAG were
indeed larger in 2018 than in 2014 (Dunn's Multiple Comparison Test, $p < 0.05$) suggesting that
size-specific fecundity increase was attributable to this larger body size in 2018. Similarly,
carapace lengths of brooding females were always larger at Snake Pit compared to TAG both in
2014 and in 2018 (Dunn's Multiple Comparison Tests, $p < 0.05$).

Variations were also observed in egg volumes between different sampling years
(Kruskal-Wallis, H= 87.434, $p < 0.05$) (figure 2c). Larger egg volumes were found in 2014
compared to other years of sampling for both vent fields and across all developmental stages
(Dunn's Multiple Comparison Tests, $p < 0.05$). We note that the 2014 dataset was produced

separately in a previous study (See supplementary material) and may be biased by an observer
effect.

<u>Variations in reproductive patterns between *Rimicaris* species</u>

Like *R. exoculata*, the realized fecundity of *R. kairei* and *R. chacei* correlated positively
with carapace length (Spearman's correlation: *R. kairei*: r= 0.40, p= 0.0044; *R. chacei*: r= 0.70, p=
0.00066) (figure 3a). Moreover, the slopes of the relationship between $\log_e$-transformed
realized fecundity and $\log_e$-transformed carapace length did not significantly differ between *R.*
exoculata and its sister species *R. kairei* (*F*= 1.6168, p= 0.2054). However, they significantly
differed between *R. exoculata* and its co-occurring congener *R. chacei* (*F*= 6.6376, $p < 0.05$).

Overall, size-specific fecundity ranged from 23 to 112 eggs mm^{-1} for *R. kairei* and from
26 to 255 eggs mm^{-1} for *R. chacei* (figure 3b). Large differences in size-specific fecundity were
observed between *R. exoculata* and other *Rimicaris* species (Kruskal-Wallis *H*= 68.836, $p < 0.05$).
Size-specific fecundity of *R. chacei* was always significantly greater compared to *R. exoculata*
both at TAG (*R. exoculata*: 49 ± 13; *R. chacei*: 129 ± 59; Dunn's Multiple Comparison Test, $p <$
0.05) and Snake Pit (*R. exoculata*: 69 ± 31; *R. chacei*: 148 ± 87; Dunn's Multiple Comparison
Test, $p < 0.05$). These differences were independent of size variations between the two species
with *R. chacei* brooding females being either of similar size at TAG (Dunn's Multiple Comparison
Test, $p > 0.05$) or even smaller at Snake Pit (Dunn's Multiple Comparison Test, $p < 0.05$). Less
differences in size-specific fecundities were observed between *R. exoculata* and *R. kairei* with
only a higher size-specific fecundity for *R. kairei* (72 ± 20) from Kairei compared to *R. exoculata*
from TAG (49 ± 13; Dunn's Multiple Comparison Test, $p < 0.05$). Such variations may be linked
to differences in body size of brooding females, as *R. exoculata* from TAG were smaller in
average than *R. kairei* from Kairei (Dunn's Multiple Comparison Test, $p < 0.05$).

Large differences were also observed in egg volumes between the three *Rimicaris*
species (Kruskal-Wallis *H*= 49.80, $p < 0.05$) (figure 3c,d). Thus, egg volumes of *R. chacei* were
significantly smaller than those of *R. exoculata* at any stage across all vents (Dunn's Multiple
Comparison Test, $p < 0.05$) (figure 3c). In contrast, differences in egg volumes were more
limited between *R. exoculata* and *R. kairei* with significantly larger eggs for *R. kairei* from Kairei

compared to *R. exoculata* from TAG for mid stages only (figure 3d). Surprisingly, we did not find gradual increase of egg volumes between successive developmental stages of *R. chacei* as was observed in the two other *Rimicaris* species, either at TAG (Kruskal-Wallis *H*= 2.57, *p* > 0.05) or at Snake Pit (Kruskal-Wallis *H*= 0.61, *p* > 0.05). Of note, we did not observe ready-to-hatch or hatched egg broods in *R. chacei* as was seen in the other shrimps.

Temporal variations in the presence of *Rimicaris* ovigerous females

A large number of ovigerous females of *R. exoculata* and *R. chacei* were retrieved in February 2018 at both MAR vent fields (figure 4a). Indeed, 55.4% and 18.2% of sexually mature *R. exoculata* females collected from TAG and Snake Pit respectively, were ovigerous females, while these account for 70.6% and 32.3% of sexually mature *R. chacei* females from TAG and Snake Pit, respectively. Ovigerous females of *R. exoculata* were also present between late March and April 2017 although in a relatively lower proportion, constituting 25.7% and 4% of sexually mature females collected at TAG and Snake Pit, respectively (figure 4a). The majority of *R. exoculata* ovigerous females were collected in dense aggregations close to vent orifices with only two ovigerous females taken outside these assemblages. On the other hand, ovigerous females of *R. chacei* were collected either in hidden aggregations where large populations of their adults have been retrieved (Methou et al. 2021) or within dense aggregations of *R. exoculata*. As highlighted previously (Hernández-Ávila et al. 2021), ovigerous females of *R. exoculata* were almost absent from samples collected outside this period of the year – i.e., January to April – despite more than 35 years of sampling efforts along the MAR (figure 4a). Similarly, ovigerous females of *R. chacei* were almost absent from other sampling reports with only two ovigerous females collected in January 2014 at Snake Pit and one in June 1994 at Lucky Strike (figure 4b)

Large numbers of ovigerous *R. kairei* were also retrieved in February 2002 and 2016 at Kairei on the CIR (figure 4c), constituting respectively 16.3% and 19.2% of sexually mature females collected at this vent field. Conversely, ovigerous females were completely absent from samples collected in November 2009 (figure 4c). Like *R. exoculata*, all of these ovigerous females were collected in dense aggregations of shrimps next to hot fluid emissions.

Discussion

Distinct reproductive strategies in _Rimicaris_ shrimps with evidence of diet influence

Rimicaris exoculata and *Rimicaris chacei* exhibit distinct reproductive traits unlikely to simply result from differences in mean body sizes. *Rimicaris kairei* is similar to its sister species *R. exoculata*, in terms of size-specific fecundity or mean egg size. Our results indicate that the two lineages (*R. exoculata* and *R. kairei*, vs *R. chacei*) are situated at two ends of the trade-off between size-specific fecundity and egg sizes currently observed within alvinocaridids (Ramirez-Llodra et al. 2000, Ramirez-Llodra and Segonzac 2006, Copley and Young 2006, Nye et al. 2013). On one hand, *R. chacei* brooding females display one of the highest size-specific fecundity and one of the smallest mean egg volumes among alvinocaridids, whereas *R. exoculata/R. kairei* brooding females present a rather low size-specific fecundity and a large mean egg volume for this family.

These variations mirror differences observed in benthic post-settlement stages of these two lineages, with significantly smaller body size for settled juveniles, as well as much larger relative proportions of juveniles in populations of *R. chacei* (Methou et al. 2020, 2021). Higher egg numbers per brood in *R. chacei* might thus contribute to its high supply of settlers despite an apparently lower breeding population size than *R. exoculata* in northern MAR vent fields. Previous work has hypothesized distinct larval life histories between these two species based on distinct stable isotope ratios (Methou et al. 2020). Given we did not observe ready-to-hatch or hatched egg broods for *R. chacei*, it is unclear if observed differences in mean egg size between the two species would ultimately lead to distinct larval sizes at hatching. No size variation was found among first zoeal larval stages of four phylogenetically distinct alvinocaridids (Hernández-Ávila et al. 2015). Although *R. chacei* zoea have not been observed so far, it is unlikely that hatching size difference only, if it exists, explains the large size difference at recruitment between *R. exoculata* and *R. chacei*, which also probably results from differences in their pelagic lives during their dispersal phase.

The sister species of *R. chacei* is *R. hybisae* from vents on the Mid Cayman Spreading Centre, but interestingly size-specific fecundity of *R. hybisae* is more similar to *R. exoculata/R.*

kairei rather than *R. chacei* (Nye et al. 2013). *Rimicaris hybisae* share with *R. exoculata/R. kairei* both a comparable ecological context, living in dense aggregations close to vent fluids emissions, and a comparable gross morphology, with an enlarged cephalothorax (Zbinden and Cambon-Bonavita 2020), characters lacking in *R. chacei*. These are linked to distinct feeding strategies between the two groups of shrimps, with a mixotrophic diet for *R. chacei* (Gebruk et al. 2000, Apremont et al. 2018, Methou et al. 2020) and dependency on chemosynthetic symbionts for others including *R. hybisae* (Rieley et al. 1999, Gebruk et al. 2000, Van Dover 2002, Streit et al. 2015, Methou et al. 2020). Variations in fecundity according to feeding strategies have been widely recognized in marine animals such as between carnivorous and herbivorous sea turtles (Broderick et al. 2003), among seabirds with different feeding habits (Pierotti and Annett 1990) or between *Capitella* worms fed with different food sources in laboratory (Qian and Chia 1991). Reduced or increased fecundities have also been observed experimentally in several shrimp species according to their diets (Lin and Shi 2002, Tziouveli et al. 2011, Almeida et al. 2018). We therefore posit that the differences we observe between fecundities in our three *Rimicaris* shrimps should be linked to their feeding ecology, rather than being constrained by their phylogeny.

Despite these variations in average egg number, mean egg sizes of *R. hybisae* and *R. chacei* were in contrary comparable with similar maximum and minimum egg diameters (Nye et al. 2013). As *R. exoculata* and *R. kairei* also exhibit similar egg volumes, phylogenetic constraints are possibly exerted on mean egg sizes for these shrimps. Further work on more alvinocaridid shrimps with different feeding strategies should help to elucidate these questions.

The intriguing reproductive cycles of *Rimicaris* shrimps

We report here a much-increased presence of ovigerous *R. exoculata* in February 2018 and to a lower extent in late March-April 2017, compared to other year periods. Combined with previous data from the literature (summarized in figure 4A), these results support the existence of a brooding period for *R. exoculata* mostly during boreal winter, starting in January/February and ending around March/April. Moreover, we did not observe any variability in reproductive features of *R. exoculata*, either in terms of fecundity or proportions of reproductive females,

between different sampling years – indicating a relative inter-annual stability of these traits

(figure 2). Since gamete production can be sustained continuously in species depending on local

chemosynthesis for their diet, mechanisms driving periodicity in shrimp life cycles are more

likely to act on their planktotrophic larval phase through seasonality in availability of

photosynthetically derived food. The few previous studies looking at a limited number of

samples indeed reported asynchronous ovarian development for *R. exoculata* suggesting

continuous egg production (Ramirez-Llodra et al. 2000, Copley et al. 2007). However, the

oocyte maximum size appeared to be lower in individuals collected summer than in those

collected in autumn, regardless of the vent site origin [16, 34], which may also point at a

gradual accumulation of mature oocytes in ovaries, culminating in winter, prior to an increase

in females spawning activity. Additional observations of intraovarian development through the

year are clearly needed to better characterize rhythms in oocyte production, which may also

well be decorrelated from spawning activity as seen in some deep-sea corals from the MAR

(Rakka et al. 2021). Large proportions of *R. chacei* ovigerous females were also retrieved in

February 2018 at the two vent fields, in contrast to historical sampling at other periods along

the MAR (figure 4B). Hence, a similar periodicity in reproduction likely exists for both *Rimicaris*

shrimps at MAR. Ovigerous *R. chacei* females were however scarce in other expeditions both

during and outside this period (figure 4B), which could be due to difficulties in sampling this

rather fast swimming species, thus preventing so far, a full appreciation of its reproductive

activity.

We also found large proportions of *Rimicaris kairei* ovigerous females during austral

summer in February 2002 and 2016, while they were absent from collections in August 2000

and November 2009 (figure 4c). Spawning in *R. kairei* thus appears to follow the same timing as

R. exoculata and *R. chacei*. Such pattern was unexpected given their distribution in different

hemispheres, at vent fields with opposite latitudes (figure 1). Reproductive periodicity in deep-

sea animals has often been related to seasonal variations in surface photosynthetic production

[26]-[28], and *Rimicaris* shrimps from the two hemispheres are expected to experience

opposite trends in terms of surface primary production along the year (Pabortsava et al. 2017,

Harms et al. 2021), driving opposite reproductive timing. Although our dataset for *R. kairei* is

more limited than for *R. exoculata*, these sister species thus appear to share same periodic
brooding cycle, regardless of local seasonal variations in surface photosynthetic productivity.
Rimicaris species studied here provide a first example of reproductive periodicity in deep-sea
ecosystems apparently unlinked to seasonal variations of photosynthetic materials from the
surface.

Besides light and food supply, temperature has also been recognized as an important
external time-giving cue in marine organisms (Mat 2019). Compared to most of the deep sea
where temperature is relatively constant within regions, hydrothermal vents constitute heat
anomalies with steep and unpredictable gradients at small spatial scales, even within a same
animal aggregation (Schmidt et al. 2008). Thermal variability at an annual scale could perhaps
serve as a time-synchronising cue for these *Rimicaris* shrimps. So far, thermal variations in
these ecosystems at a year-round scale remain poorly documented. However, long-term
observatories deployed on two vent fields from the northern hemisphere have reported clear
tidal influence, although temperatures are relatively stable throughout the year (Barreyre et al.
2014, Cuvelier et al. 2017). Recently, it was revealed that bathymodiolin mussels inhabiting
vents on the MAR retain and express circadian clock genes following tidal cycles which may be
mediated by either tidal stimulus or internal clock (Mat et al. 2020). Possibly, *Rimicaris* shrimp
species studied herein also use a set of biological clocks to time their reproductive activity.
Though, not all alvinocaridid shrimps seem to exhibit periodicity in their reproductive patterns,
despite a gametogenesis phase being strongly constrained phylogenetically (Ramirez-Llodra et
al. 2000). For example, regular sampling of *Mirocaris fortunata* yielded ovigerous females
throughout the year (Ramirez-Llodra et al. 2000, Komai and Segonzac 2003, Methou 2019) and
similarly in *Rimicaris variabilis* (Komai and Tsuchida 2015, Komai et al. 2016). It therefore
appears unlikely that phylogenetic constraints alone have maintained a periodic brooding in
several alvinocaridid genera but not in others. Then again, other alvinocaridid species may also
indeed follow seasonality in surface production, like *Alvinocaris stactophila* with a brooding
period between November and March (Copley and Young 2006), and inhabiting relatively
shallow cold seeps under highly productive coastal waters. Hence, several synchronizing factors
may co-exist in vents and other deep-sea ecosystems, even within the same family of animals.

For example, our results here demonstrate that *R. exoculata* and *R. kairei* exhibit seasonality
decoupled from surface seasonality, but as the study areas have relatively low surface primary
productivity and export of particulate organic carbon (POC) to the deep (Pabortsava et al. 2017,
Harms et al. 2021) it is also possible that a higher POC export may begin to interfere with this
pattern. Taken together – primary surface production may be the key synchronising cue for
some deep-sea species as demonstrated in bathymodiolin mussels and vent crabs (Dixon et al.
2006, Tyler et al. 2007), but our results underscore that reproductive cycles and seasonality at
the deep is not as simple as it was thought to be.

A thorough understanding of reproductive ecology of the dominant species shaping the
ecosystem is key to assessing the resilience of deep-sea ecosystems, but little has been done so
far even for emblematic taxa like *Rimicaris*. With deep-sea mining for massive sulphides and
other resources being imminent (Van Dover et al. 2018), the habitat of these amazing animals
that can live nowhere else is being threatened. Elucidating the ecology of these inhabitants of
the deep is now more important than ever, and our data exemplifies the importance of time-
series studies. Research expeditions in a particular region tend to be conducted in similar
timings of the year due to seasonal variations in sea and weather conditions, but our results
show that communities need to be revisited at different times of the year.

Ethics

The animals used in this study were invertebrate carideans collected in the French contract
area for the exploration of polymetallic sulphur deposits (Mid-Atlantic Ridge) or International
Waters (Central Indian Ridge). Each individual was immediately preserved in ethanol after
recovery and no live experiments were carried out.

Author's contributions

P.M. collected and sorted specimens in 2018 on the MAR, carried out the identification and
measurement of specimens from all sampling cruises, lead the data analysis, participated in the
conception and design of the study and drafted the manuscript; C.C. carried out the collection
of specimens in 2016 on the CIR and helped draft the manuscript; H.K.W. carried out the
collection of specimens in 2009 and critically revised the manuscript; M-A.C-B. collected

specimens in 2017 on the MAR, coordinated the study and critically revised the manuscript; F.P. collected and sorted specimens in 2017 conceived and designed the study, coordinated the study and helped draft the manuscript. All authors gave final approval for publication and agreed to be held accountable for the work performed therein. The authors declare they have no competing interests.

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<u>Figure Captions</u>

● *Rimicaris chacei* ● *Rimicaris exoculata* ● *Rimicaris kairei*

Figure 1. Context of the study. **a.** Current geographic distribution of the three *Rimicaris* species studied. **b.** Dense aggregations of *R. kairei* shrimps in February 2016 at the Kairei hydrothermal vent field.

Figure 2. Interannual comparison of reproductive features of *Rimicaris exoculata* collected between 2014 and 2018. **n:** number of individuals for each condition. **a.** Variation of $\log_e$-transformed minimum realized fecundity with $\log_e$-transformed carapace length. **b.** Fecundity corrected for body size (carapace length) **c.** Egg volumes at different developmental stages.

Figure 3. Comparison of reproductive features between *Rimicaris* species from the MAR and the CIR. **n:** number of individuals for each condition. **a.** Variation of $\log_e$-transformed minimum realized fecundity with $\log_e$-transformed carapace length. **b.** Fecundity corrected for body size (carapace length). **c.** Egg volumes at different developmental stages of *Rimicaris* shrimps from the MAR. **d.** Egg volumes at different developmental stages of *R. exoculata* and *R. kairei*.

[1]Williams and Rona (1986), [2]Vereshchaka (1997), [3]Shank et al. (1998), [4]Ramirez-Llodra et al. (2000), [5]Copley et al. (2007), [6]Komai and Segonzac (2008), [7]Guri et al. (2012) [8]Lunina and Vereshchaka (2014), [9]Methou et al. (2019) & Hernàndez-Avila et al. (2021), [10]Cruise Report BICOSE (2014), [11]This work, [12]Watabe & Hashimoto (2002)

Figure 4. Diagrams summarizing occurrence of reproductive stages of the studied *Rimicaris* shrimp species over different samplings realized between 1985 and 2018 according to the year period **a.** for *R. exoculata.* **b.** for *R. chacei.* **c.** for *R. kairei.*

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