ABSTRACT: Sponges are an ecologically important component of modern Caribbean coral reefs. However, little is known about the structure of sponge communities prior to the large-scale degradation of Caribbean reef ecosystems. Here we explore changes in the sponge community over the past millennium by analyzing the composition of sponge spicules from a sediment core collected from a lagoonal reef within the archipelago of Bocas del Toro, Caribbean Panama. The analysis reveals a change in spicule composition that began approximately 400 yr ago. During this time, the share of monaxial spicules, belonging mostly to haplosclerid and axinellid sponges, decreased while the relative number of spherical spicules, found typically in *Placospongia*, *Geodia*, and some chondrillids, increased. These results were compared with previously published data on parrotfish, corals, and reef accretion rates obtained from the same core. The increased share of spherical spicules did not correlate with contemporaneous declines in the abundance of parrotfish (determined from fish teeth) or with trends in the relative abundance of dominant coral species (determined from coral skeletal remains) but was weakly correlated with reef accretion rates (determined from sediment accumulation rates). Spicule morphogroup diversity and evenness increased over the past ~400 yr, suggesting community changes were not due to reef environments becoming less habitable for reef sponges. Although not tested directly, the increase in spherical spicules may be due to declines in the abundance of sea turtles that preferentially feed on sponges that contain these spicule types.

KEY WORDS: Sponge spicules · Hawksbill turtles · *Placospongia* · *Geodia* · Bocas del Toro

INTRODUCTION


As the abundance of Caribbean reef-building corals has been reduced by 50% over the past few decades (Jackson et al. 2014), sponges have become the dominant benthic organisms on many Caribbean reefs (McClenahan et al. 1999, Loh et al. 2015). Fluctuations in sea temperature and chemistry, diseases, and water pollution are thought to be among the main factors responsible for the rise in sponge dominance (D’Croz et al. 2005, Easson et al. 2015). Although detailed moni-
MATERIALS AND METHODS

Study area

The Bocas del Toro archipelago is a geographically complex area composed of numerous islands surrounded by extensive mangrove systems and fringing and patch reefs (Fig. 1A; D’Croz et al. 2005, Cramer 2013). Modern reef ecosystems began to accrete around 8000 yr ago following postglacial sea level rise (Coates et al. 2005) and started to experience anthropogenic impact as early as 250 yr ago (Cramer et al. 2017). More recent changes were first noted by surveys conducted in the 1980s (Lessios et al. 1984), with accelerated changes documented in subsequent decades (Guzmán & Guevara 1998a, b, 1999, 2001, Seemann et al. 2014). Although the biomass and diversity of sponges often exceed that of corals in Bocas del Toro (see supplementary material in Loh & Pawlik 2014), there are currently no data on trends in sponge communities for this region.

Reconstructing sponge community composition

To track long-term change in sponge communities, we analyzed the composition of sponge spicule assemblages within a 305 cm long sediment core obtained from the Punta Donato patch reef (9.3573° N, 82.3680° W; Fig. 1A). The reef core was composed primarily of corals, mollusc shells, and other carbonates in a sandy–muddy matrix. Details of the extraction and subsampling of this core are in Cramer et al. (2017). To provide a moderate-resolution record of change in spicule composition, we analyzed spicule assemblages from 23 samples of 5 cm thickness along the length of the core (Fig. 1B). Although the larger study of millennial-scale, ecosystem-level change in Bocas del Toro involved the collection of 2 replicate cores from each of 3 reef sites in Bahía del Almirante, the Punta Donato site provided a high-resolution record from the historical period to the present (Cramer et al. 2017). To ensure that sufficient core sediments were preserved for analysis of other fossil groups, we focused on a single core from Punta Donato for our spicule analysis. Because of dif-
ferring sampling protocols across fossil groups, only 9 of the 23 samples analyzed for spicules also have data for corals and reef fish teeth (published in Cramer et al. 2017).

Sample maceration

Samples of approximately 50 g of reef matrix sediment were macerated using dilute hydrogen peroxide to remove organic matter and separate individual sponge spicules. Samples were heated on a hotplate to accelerate reactions, rinsed with water several times to remove all reaction by-products, and dried at 70°C. Next, the samples were sieved to remove all sediments >5 mm which did not contain sponge spicules. The residue was digested with 30% acetic acid to dissolve all calcium carbonate, resulting in the exclusion of 2 calcareous sponge species from our analyses (i.e. Clathrina primordialis and Leucetta floridana). Following rinsing, the remaining residue, consisting mostly of siliceous sponge spicules and inorganic silica particles, was dried. The spicule residue was laid out on a counting plate with a brush and counted under a stereomicroscope from randomly chosen 2 × 2 mm square areas. Each spicule encountered was assigned to 1 of 10 morphological groups (Tables S1 & S2 in the Supplement at www.int-res.com/articles/suppl/m601p127_supp.pdf, Figs. 2–4) until 1000 spicules were counted per sample. For further details, see Łukowiak et al. (2013), Łukowiak (2016), and the literature cited therein.

Assessing timing of change in sponge communities

Sample ages were estimated using uranium–thorium (U–Th) dating of coral skeletal fragments (Clark et al. 2012, Cramer et al. 2017). A highly constrained chronology was produced for 1 replicate core at Punta Donato, where 4 U–Th ages were obtained from the top 1 m of the core and 1 date from approximately every 0.5 m interval below that, yielding 6 ages total for the portion of the core that represents coral reef habitat (the bottom of the core represents soft-bottom habitat prior to reef formation; Cramer et al. 2017). From this core, accretion rates were estimated using linear interpolation between each pair
of ages. For the second Punta Donato replicate core from which spicules were analyzed, 2 U–Th ages were obtained (1 each from the top and bottom of the section representing reef habitat; Fig. 1B). Sedimentation (reef accretion) rates from the well-dated core were linked with the replicate core by using the top age to constrain the sedimentation rate of the latter to that of contemporaneous periods of the former (for details, see Cramer et al. 2017). Because reef accretion rates were available for each sample, all samples analyzed for spicules contain an accretion estimate.

Analysis of community change

Temporal patterns in sponge spicule assemblages were assessed by computing the relative abundance (proportion of total number of spicules counted) of each of the 10 spicule morphotypes within each sample and then plotting these values for the most common groups (comprising >1% of assemblage on average). Temporal patterns were assessed across the time span representing the full core length using non-parametric locally weighted regression (loess)-smoothed trendlines (Cleveland & Devlin 1988). To assess general patterns across the full time series while capturing non-unidirectional fluctuations in the data, 90% of the time series was used to fit each local regression (smoothing parameter = 0.9). Contemporary patterns in possible drivers of sponge composition—parrotfish abundance (proportion of fish teeth fossils positively identified as belonging to parrotfish) and reef accretion rates (mm of reef sediment accumulated per year, based on accretion rate estimates derived from U–Th dates as described in the previous subsection), both taken from Cramer et al. (2017)—were plotted alongside the relative abundance of dominant spicule type in assemblages. Temporal patterns in spicule morphogroup richness, diversity (measured as Shannon’s diversity index $H$, shown in Eq. (1), where $p_i =$ proportion of spicules made up of the $i^{th}$ morphogroup and $S =$ total number

$$H = -\sum_{i=1}^{S} p_i \log_2 p_i$$

Fig. 2. Spicule morphotypes present in reef sediments of the studied area belonging to the following morphogroups: oxeas (1–5: oxeas, 6: acanthoxea, 7: strongyle, 8–10: styles), tylostyles (11–14: tylostyles), others (15: acanthostyle?, 16, 17: tylotes, 18–20: acanthoxeas; 21: acanthostrongyle; 22: diactine). Scale bars: 1–9, 11–16, 18–21, 22: 50 μm; 10, 17: 100 μm.
RESULTS

Sponge spicule composition and preservation

Ten morphogroups were designated from the spicule assemblage within the Punta Donato core: amphitriaenes, aspidasters, calthrops, oxeas, sterrasters/selenasters, spherasters/anthasters, triaenes, triods, tylostyles, and an ‘others’ group which represented other, uncommon morphogroups (Figs. 2–4, Table S1). This assemblage captures the overall morphological variability of the siliceous spicule material observed in bulk sediment samples within Bahía del Almirante (e.g. Łukowiak et al. 2013). Oxeas (Fig. 2.1–2.10) and sterrasters/selenasters (Fig. 3.23–3.24) were consistently the most abundant morphogroups throughout the core, accounting for 87 and 9%, respectively, of all spicules across all samples (Table S1, Fig. 5). Spheraster/anthaster (Fig.

\[ H = \text{abs}\sum_{i=1}^{S} p_i \times \ln(p_i + 0.001) \]
3.28–3.34) and tylostyle (Fig. 2.11–2.14) morphogroups were subdominant (comprising >1% on average). Spicule preservation, as assessed by physical appearance, was uniform along the entire length of the core. There was no apparent difference (as visually estimated across all samples) in the frequency of broken spicules within samples over time and all spicule morphogroups were observed in all samples, suggesting little bias in preservation over time.

Long-term change in sponge assemblages

The most notable change is the decrease in the relative abundance of monaxial spicules (oxeas) and increase in spherical spicules (sterrasters/selenasters) over time beginning approximately 400 yr ago (Fig. 5A). From 1600 to 1954 AD, oxeas declined from 94 to 67 % of spicules while sterrasters/selenasters increased from 3 to 29 % of spicules (Fig. 5A). Although parrotfish abundance and reef accretion rates also followed this general decline over the past 400 yr (Fig. 5C,D), the relative abundance of spherical spicules was not correlated with the relative abundance of parrotfish teeth and only weakly correlated with reef accretion rates (Fig. 6A,B). Relative abundance of spherical spicules was not correlated with the relative abundance of either of the most common coral species found in the Punta Donato core, low-relief branching *Madracis mirabilis* and *Porites* spp. (Fig. 6C,D).

Components of spicule morphogroup diversity—richness and evenness—showed contrasting patterns during the period of observed change in spicule assemblage encompassing the past 4 centuries. Richness per sample fluctuated from 4 to 9 morphogroups over
Fig. 5. Relative abundance of (A) dominant and (B) subdominant sponge spicule morphotypes, (C) parrotfish teeth, and (D) reef accretion rates in Punta Donato core. Percent parrotfish teeth computed from all teeth found in sediments >63 μm; accretion rates determined from linear interpolation of U–Th ages from well-dated Punta Donato replicate core (see Cramer et al. 2017). Trendlines from loess-smoothed estimates; smoothing parameter = 0.9

Fig. 6. Relative abundance of spherical spicules (sterrasters/selenasters) versus potential drivers of sponge spicule composition: (A) parrotfish teeth, (B) reef accretion rates, (C) Madracis mirabilis coral, (D) branching Porites coral. Trendline indicates significant (Pearson) correlation at p = 0.1 level.
the last millennium and did not show a clear pattern over the past 400 yr (Fig. 7), while evenness remained stable during the first 600 yr of the time series (from ~1000 to 1600 AD) and then gradually increased from 1600 to 1956 AD. The Shannon-Wiener diversity index followed a similar pattern as evenness.

**DISCUSSION**

**Assessing sponge community composition from spicules**

Because sponge species can include multiple spicule morphotypes, it is often impossible to assign spicule types to a single taxonomic group. However, spherical spicules belonging to the sterraster/seLENaster and spheraster/anthaster groups can be confidently assigned to a small number of taxa present in the Bocas del Toro region, assuming there has been no extinction in the last ~2000 yr. Sterrasters and selenasters dominate in *Geodia* and *Placobiongxia*, respectively, while spherasters and anthasters can be found in *Chondrilla*, *Placobiongxia*, *Tethya*, and *Diplastrella* (Table S1). In addition, the spicules falling within the oxea morphogroup (i.e. all monaxial spicule morphotypes except for tylostyles) are characteristic for poecilosclerids, astro phorids, halichon- drids, spirophorins, bubarids, and tethyids. However, they dominate only in haplosclerids (including *Placobiongxia, Niphates*, and *Oceanapia*) and axinellids. Tylostyles appear in clionoids, hadromerids, and some poecilosclerids (Table S1).

**Potential drivers of long-term sponge community change**

Although we are not able to directly track the full taxonomic composition of past sponge communities from spicule assemblages, we are able to gather insight into changes in sponge taxa that are preferentially eaten by various predators. Among sponges present in the studied area, those that are most palatable for fishes bear both spherical and monaxial spicules. Parrotfish preferentially feed on *Placobiongxia, Geodia*, and *Callyspongia* (Pawlik et al. 1995) and angelfish on *Callyspongia, Iotrochota*, and *Chondrilla* (Randall & Hartman 1968). Hawksbill turtles, in turn, feed mostly on sponges that bear almost exclusively spherical spicules of the sterraster/seLENaster and spheraster/anthaster morphogroups, i.e. *Placobiongxia, Geodia*, and *Chondrilla* (Pawlik et al. 2018). Thus, our spicule record provides a possible proxy for long-term change in sponge predator abundance, particularly in the absence of other explanations for change in the spicule composition of reef sediments.

We hypothesize that the observed decline in sponges bearing monaxial spicules (mostly haplo- sclerids and axinellids) and rise of sponges bearing spherical spicules (e.g. *Placobiongxia, Geodia*, and *Chondrilla*) since 1600 AD is best explained by the loss of a major sponge predator, the hawksbill turtle *Eretmochelys imbricata*, from historical overfishing. The selective feeding of hawksbill turtles on sponge taxa bearing spherical spicules has been shown to play an important role in structuring the reef sponge community and limiting the distribution of some Caribbean sponges (Meylan 1988, Dunlap & Pawlik 1996, Pawlik 1998, León & Bjorndal 2002, Pawlik et al. 2018). Hawksbill turtle populations have been greatly reduced in the Caribbean by overfishing since European contact, with their abundance reduced to between 1 and 10% of pre-Columbian levels (Parsons 1972, Meylan & Donnelly 1999, McClenachan et al. 2006). Archeological data from Bocas del Toro show that exploitation of sea turtles by indigenous populations predates European contact, intensifying between 1100 and 1400 AD (Wake et al. 2013). Turtle exploitation continued to increase after first contact with Europeans; old historical notes, dating back as far as 1684, suggest apparent attempts to use hawksbill turtles for their meat (McClenachan et al. 2006, Pawlik et al. 2018). However, for the next 2 centuries, their consumption was likely limited because their meat was generally considered toxic and not particularly edible (McClenachan et al. 2006). Nevertheless, the turtle meat industry had already been established at the time, and hawksbills were being hunted also for their fat, eggs, and shells (Wake et al. 2013).
In pre-Columbian and around first contact times, when hawksbills were presumably still abundant, the ratio of turtles to palatable sponges may have forced hawksbills to feed on both chemically undefended and defended sponge species, which could have affected the taste of their meat. Around 1900, however, the meat of hawksbill turtles was already considered edible and nourishing (McClenachan et al. 2006). This shift in the tastefulness of the turtle meat is suggested to be due to increased availability of palatable sponge species following declines in turtle predation.

Fishes, including parrotfish, feed on Caribbean sponges as well (Dunlap & Pawlik 1996). However, they are not considered major sponge predators in this region (Randall 1967). Although the number of parrotfish apparently began declining as early as 300 yr ago at the Punta Donato core (Cramer et al. 2017), we did not find a correlation between abundance of parrotfish teeth and spherical spicules. This may be because the sponge species preferred by parrotfish (i.e. Placosporgia, Geodia, and Callyspongea; see Pawlik et al. 1995) bear spicules falling within both the monaxial and spherical morphogroups. As such, our data cannot test the hypothesis that overfishing of parrotfish was a driver of observed changes in the sponge community (Pawlik et al. 2013, Loh & Pawlik 2014).

Sponge spicule morphogroup evenness and diversity increased over the past ~400 yr, suggesting that reef environments became more favourable for sponges with mineralized spicules at this reef. This timing coincides with a change in the dominant reef-building coral species at Punta Donato, from branching Porites spp. to Madracis mirabilis, that was also initiated approximately 500 yr ago (Cramer et al. 2017). Notably, the relative abundance of spherical spicules continued to increase over the past 200 yr despite resumption in dominance of branching Porites corals during this period. These trends suggest that the observed shift in dominant morphogroup, from less palatable sponges bearing oxeas to more palatable sponges with sterrasters/selenasters, is not primarily due to a change in reef coral community composition but instead relates to prey release following the overfishing of sponge predators such as hawksbill turtles.

**Potential biases**

Several taphonomic filters could have affected the composition of sponge spicule assemblages in our study. First, we were only able to analyze spicules from taxa that produce an opal mineral skeleton (170 of ~198 species, Table 1 in Łukowiak 2016). Although this represents only 86% of taxa, such groups represent over 40% of total living sponge biomass in Bocas del Toro today (Łukowiak et al. 2013). Second, small spicules less than 25 µm in size were not included in our analyses because of their poor preservation potential. For a more extensive discussion regarding caveats related to the assessment of loose sponge spicules and their relationship to the living sponge community, see Łukowiak et al. (2013). Third, monaxial spicules (oxeas, styles, and strongyles) are more likely to be physically damaged because of their thin and elongated morphology than more robust spherical types (selenasters, sterrasters, and spherasters) (Rützler & MacIntyre 1978). Thus, spherical spicules may be generally overrepresented in our samples. However, we observed that the proportion of broken spicules did not change significantly over time and the general preservation of spicules was continually excellent, even in the oldest samples, suggesting low levels of selective dissolution. Moreover, we found that the relative abundance of the more delicate monaxial spicules was greater in older not younger samples, suggesting that breakage of fragile spicules did not significantly bias our results (Fig. 6). Thus, our spicule assemblage accurately records long-term changes in reef sponge community composition in Bocas del Toro that may be related to historical overfishing of sea turtles. This study highlights the unique ability of the paleoecological record to detect alterations in reef ecosystems that occurred centuries before modern reef surveys, providing valuable insights into the anthropogenic mechanisms of the recent changes of these systems.

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