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# A decade of coral biobanking science in Australia - transitioning into applied reef restoration

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Active restoration or intervention programs will be required in the future to support the resilience and adaptation of coral reef ecosystems in the face of climate change. Selective propagation of corals *ex situ* can help conserve keystone species and the ecosystems they underpin; cross-disciplinary research and communication between science and industry are essential to this success. Zoos and aquaria have a long history of managing *ex situ* breed-for-release programs and have led the establishment of wildlife biobanks (collections of cryopreserved living cells) along with the development of associated reproductive technologies for their application to wildlife conservation. Taronga Conservation Society Australia's CryoDiversity Bank includes cryopreserved coral sperm from the Great Barrier Reef, which represents the largest repository from any reef system around the globe. This paper presents results from an inventory review of the current collection. The review highlighted the skew toward five *Acropora* species and the necessity to increase the taxonomic diversity of the collection. It also highlighted the need to increase geographic representation, even for the most well represented species. The inventory data will inform Taronga's future research focus and sampling strategy to maximize genetic variation and biodiversity within the biobank and provide a test case for other practitioners implementing biobanking strategies for coral conservation around the world. Through co-investment and collaboration with research partners over the next decade, Taronga will prioritize and resource critical applied research and expand biobanking efforts to assist interventions for reef recovery and restoration.

## KEYWORDS

reef restoration, cryopreservation, biobanking, coral spawning, coral

## Introduction

The effects of climate change on our oceans imperil the health and resilience of coral reef systems around the globe. Addressing drivers of global climate change, while concurrently implementing restoration interventions at a large scale, provides the best chance for the continued existence of these ecosystems and their essential function and services (Anthony et al., 2017; Hoegh-Guldberg et al., 2019; Knowlton et al., 2021; Vardi et al., 2021). In Australia, the Great Barrier Reef (GBR) has experienced an increase in the frequency of climate-change-driven mass bleaching events over the last decade, including in its previously untouched southern range (Hughes et al., 2018; Australian Institute of Marine Science, 2021; Great Barrier Reef Marine Park Authority et al., 2022). These unprecedented heat stress events have caused a decline in coral cover and have reduced reproductive output (Hagedorn et al., 2016; Howells et al., 2016; Daly et al., 2022), consequently depressing larval supply (Cheung et al., 2021) and recruitment (Hughes et al., 2019).

In addition to action on climate change and improved local management practices, active reef restoration approaches to repair and re-seed reefs will likely be required for many sites to maintain or restore biodiversity and support or accelerate adaptation (Anthony et al., 2017; van Oppen et al., 2017; Suggett and van Oppen, 2022). Current restoration efforts primarily use asexual propagation methods (colony fragmentation) to generate new material for out-planting (Young et al., 2012; Boström-Einarsson et al., 2020); however, this approach has limitations for upscaling to a system as large as the GBR and it only generates clonal material. There is increasing interest in the sexual production of coral juveniles for out-planting (e.g., Randall et al., 2020), which is more amenable to upscaling, has the added advantage of helping to maintain population genetic diversity, and has the ability to strengthen or introduce adaptive traits to populations through selective breeding (Quigley et al., 2020; Hagedorn et al., 2021).

The sexual reproduction of corals *ex situ* is commonly achieved either by allowing individual colonies to naturally spawn in the same system or by controlling pairwise matches by selectively mixing freshly washed oocytes with optimized concentrations of fresh sperm (commonly termed ‘*in-vitro* fertilization’, or IVF). The relatively recent development of sperm cryopreservation technologies for corals (Hagedorn et al., 2006) enables selective breeding to also occur using sperm samples that are spatially and temporally removed from fresh oocyte supply. Cryopreservation refers to the controlled cooling of living cells and tissues to ultra-low temperatures using techniques that maintain the structure and function of cells when samples are brought back to physiological temperatures, allowing them to be stored for decades, possibly centuries, if maintained correctly. Biobanking of cryopreserved samples, when combined with assisted reproductive technologies such as IVF, can support species

conservation by preserving genetically distinct specimens over long time periods. The potential application of biobanking to wildlife conservation was recognized in the 1970s (Watson, 1978; Benirschke, 1984) and has progressed over ensuing decades with zoos emerging as global leaders in the establishment of wildlife biobanks (Comizzoli and Wildt, 2017; Hagedorn et al., 2019; Hobbs et al., 2019; Hagedorn et al., 2021).

In 2011, scientists from the Taronga Conservation Society and Zoo (Sydney, Australia) and the Smithsonian Institution (United States of America) initiated a program with collaborators from the Australian Institute of Marine Science (AIMS) to apply coral sperm cryopreservation methods to species from the Great Barrier Reef. Cryopreservation techniques proved directly transferable to Australian *Acropora* species (*Acropora tenuis*, *A. millepora*), leading to the establishment of Australia’s first frozen coral biorepository (Hagedorn et al., 2012a; Hagedorn et al., 2012b). Moderate upscaling of IVF methods for the generation of live offspring (larvae and juveniles) using cryopreserved material was successfully demonstrated over the first three years of the program (Hagedorn et al., 2017). Subsequent improvements to the specifically designed cryopreservation apparatus (Zuchowicz et al., 2021) have aided in the up-scaling of time-sensitive sperm sample processing. Biobanking of gametes from GBR corals by the Taronga and Smithsonian teams has continued annually during spawning, utilizing wild coral colonies held temporarily at the AIMS National Sea Simulator (Cape Ferguson, Queensland, Australia). These achievements have resulted in the integration of coral biobanking *via* cryopreservation as an enabling technology within the largest research and development program for coral reefs globally, the Reef Restoration and Adaptation Program (RRAP<sup>1</sup>). A decade from its conception, we conducted a review and quality assurance check of Taronga’s CryoDiversity Bank coral inventory. This review aimed to generate census data on the existing collection to understand the quantity and scope of sampling from bioregions and species, and to identify gaps in quality assessments and metadata that need to be addressed. As the Taronga CryoDiversity Bank transitions from a relatively small-scale opportunistic sperm sample collection to a more targeted approach, we will use this information to direct future resource investment, inform collection management and ensure that the collection remains relevant to the needs of the coral conservation and restoration communities. It is hoped that the information and experiences of the Taronga CryoDiversity Bank will be useful to other coral biorepositories around the world that will likely face similar considerations as they seek to support coral conservation and restoration efforts.

1 <https://gbrrestoration.org/program/cryopreservation/>.

## Methods

Data associated with Taronga's CryoDiversity Bank, in Sydney on Cammeragal Country and in Dubbo on Wiradjuri Country, are managed using Microsoft Excel and stored on a user-restricted server. Prior to conducting the inventory review, a quality assurance check of the metadata was performed. An example of the metrics and metadata collected from each colony, and the evolution of these data fields over the 10+ year program, are included in Supplementary Material (Figure S1). New metadata fields were added as new technologies (e.g. computer assisted sperm analysis; CASA) were incorporated into best practices and metadata needs for future restoration were refined by collaborators (e.g. GIS data).

An inventory review was conducted of the data associated with physical sperm samples collected from corals of the GBR and cryopreserved between November 2011 and February 2022.

To obtain estimates of colony numbers and quantity of material from species, reefs, and regions, data were handled as follows:

- Colonies collected from the Far North and Cairns/Cooktown management areas (MAs) were combined into one group, referred to throughout as North MA.
- Database entries for 30 colonies (Central MA) were missing data on specific reef of origin. These entries were allocated to a general grouping "Unidentified Location Central".
- During each spawning period, a colony may be sampled as an individual genetic unit and/or combined with samples from other individuals of the same species and reef to form a multi-colony pooled sperm sample. Pooled samples were created when individual sperm samples were either low in volume, or when batch spawning of corals within one container was conducted. Sperm samples were only pooled within species and source MA. When determining the number of colonies from each species, colonies were:
  - a. only counted once per year even if the colony was sampled on more than one spawning night each season;
  - b. counted towards the cumulative total of colonies sampled as individuals regardless of whether they also contributed to a pooled sperm sample that year, and;
  - c. counted towards the cumulative total of colonies contributing to pooled sperm samples if they were identified by colony name and not also sampled as individuals.
- If multiple pooled sperm samples were collected from the same source colonies across more than one night in the same year, but individual colonies contributing to that pool were not specifically identified, only the pool with

the highest number of individuals was counted towards the pooled colony total for each species (Table 1).

- Raw sample motility data were calculated as the average of percent total motility, whether data were collected by manual counting (2011-2017), or CASA ( $\geq 2018$ ; Table 1). From 2019 onwards, raw motility data were derived from a sub-sample activated with a standardized concentration of caffeine and bovine serum albumen (refer to methods in Daly et al., 2022). Thus, any motility values of  $\lesssim 50\%$  observed for samples collected prior to 2019 may be underestimated due to absent or incomplete activation.

## Results

As of March 2022, Taronga's CryoDiversity Bank contains 2614 cryovials of cryopreserved sperm sourced from:

- 381 coral colonies. Of these colonies, 230 (60%) are represented by single genotypes (i.e., sampled as individuals); the remaining 151 colonies only contributed to pooled sperm samples, comprising two or more colonies.
- 30 identified species of hard coral and one additional species of *Acropora* of unknown taxonomy (referred to throughout as *Acropora* sp.).
- Four MAs of the GBR Marine Park; North MA (combining Far North and Cairns/Cooktown MAs), Central MA (Townsville and Whitsunday) and South MA (Mackay/Capricorn).

Between 2011 and 2014 a total of 446 cryovials from 7 species were cryopreserved and banked (Figure 1A) consisting of pooled sperm samples only (Figure 1B). In 2016, the first medium-scale cryopreservation of coral sperm specifically for conservation biobanking was undertaken at the National Sea Simulator (AIMS) adjacent to the central GBR. This single collection event increased the total number of cryovials preserved, the number of colonies from which sperm were sampled, and the number of species represented by approximately 100%, to a cumulative total of 1088 cryovials from 123 colonies across 11 species. This was also the first year that sperm samples were collected and cryopreserved from single colonies, creating samples of individual genetic units (Figure 1B). Between 2016 and 2020, an average of 415 cryovials of cryopreserved sperm from 5 species and 60 new colonies were added to the bank each year.

Eleven of the 31 species represented in Taronga's CryoDiversity Bank had sperm samples collected and cryopreserved across more than one year (Figure 2), five of which were sampled across six or more years. Eighteen species have been sampled only on one occasion.

TABLE 1 Summary of sample quantity and type (pooled or individual), sample quality (motility and concentration), and the number and distribution of coral colonies across local management areas (origin) represented in Taronga's CryoDiversity Bank.

Species	Total # colonies per MA sum (pooled/individually sampled)			Median # tubes per colony individually sampled median [range]	Raw sperm concentration (x10 <sup>9</sup> sperm /ml) mean ± S.D.	Raw sperm % total motility mean ± S.D.*
	North	Central	South			
<i>Acropora aspera</i>			4 (1/3)	9.5 [8-14]	1.61 ± 0.30	57 ± 15%
<i>Acropora austera</i>		5 (5/0)			0.36	12%
<i>Acropora cytherea</i>		5 (0/5)		8.5 [7-19]	3.32 ± 1.25	68 ± 13%
<i>Acropora digitifera</i>		7(7/0)			1.8	15%
<i>Acropora divaricata</i>			1 (0/1)		0.58	77%
<i>Acropora donei</i>			1 (0/1)		1.88	92%
<i>Acropora florida</i>		11 (2/9)		8 [3-11]	2.15 ± 1.01	70 ± 24%
<i>Acropora glauca</i>			4 (2/2)	6 [2-9]	2.19 ± 1.41	67 ± 7%
<i>Acropora hyacinthus</i>	7 (0/7)	21 (9/12)	6 (3/3)	8 [1-11]	2.21 ± 1.43	74 ± 21%
<i>Acropora loripes</i>		46 (19/27)		8 [3-18]	2.69 ± 1.55	67 ± 24%
<i>Acropora millepora</i>		35 (14/21)	11 (5/6)	5 [3-17]	1.40 ± 1.22	69 ± 27%
<i>Acropora muricata</i>		10 (10/0)			0.45	0.51
<i>Acropora nobilis</i>		4 (0/4)	3 (2/1)	10 [9-10]	2.08 ± 1.08	57 ± 18%
<i>Acropora sarmentosa</i>		21 (4/17)		9 [3-15]	2.47 ± 1.72	79 ± 16%
<i>Acropora spathulata</i>			4 (0/4)	14 [6-16]	1.82 ± 1.00	49 ± 3%
<i>Acropora sp.</i>			8 (0/8)	8 [3-9]	2.24 ± 1.01	50 ± 11%
<i>Acropora tenuis</i>	30 (4/26)	58 (28/30)	6 (0/6)	5 [1-14]	2.16 ± 2.27	54 ± 29%
<i>Acropora valida</i>			3 (2/1)		3.70	87%
<i>Acropora vauughani</i>		1 (0/1)			0.90	66%
<i>Astrea curta</i>			3 (3/0)		2.56	0
<i>Cyphastrea microphthalma</i>			1 (0/1)		0.43	45%
<i>Echinopora lamellosa</i>		5 (5/0)			0.41 ± 0.32	61 ± 7%
<i>Disastraea matthaii</i>		3 (3/0)			1.40	53%
<i>Fungia fungites</i>		1 (**/1)			0.02	32%
<i>Galaxea fascicularis (aspera)</i>		2 (0/2)		7 [4-10]	4.22 ± 3.94	94 ± 2%
<i>Goniastrea aspera</i>		6 (6/0)			5.44 ± 2.91	81 ± 2%
<i>Goniastrea retiformis</i>	13 (6/7)	8 (0/8)		2 [1-10]	1.64 ± 1.31	88 ± 12%
<i>Montipora aequituberculata</i>			4 (4/0)		0.68	89%
<i>Mycodinium elephantotus</i>		4 (0/4)		6 [3-8]	1.17 ± 0.55	89 ± 9%
<i>Platygyra daedalea</i>		16 (8/8)	4 (0/4)	8 [2-10]	1.91 ± 1.23	78 ± 21%
<i>Platygyra lamellina</i>		2 (2/0)			0.40	N/A

\*Raw data from both pooled and individually sampled corals are assessed together. The majority of samples were assessed for raw motility without activation using a refined protocol implemented from 2019 onwards (caffeine + BSA). Therefore motility may be an underestimate for samples <50%.

\*\*pooled sample of unknown number of contributing colonies.

The number of colonies represented as "pooled" or "individual" are identified as these samples may have different use strategies for genetic management and research. N/A, not assessed.

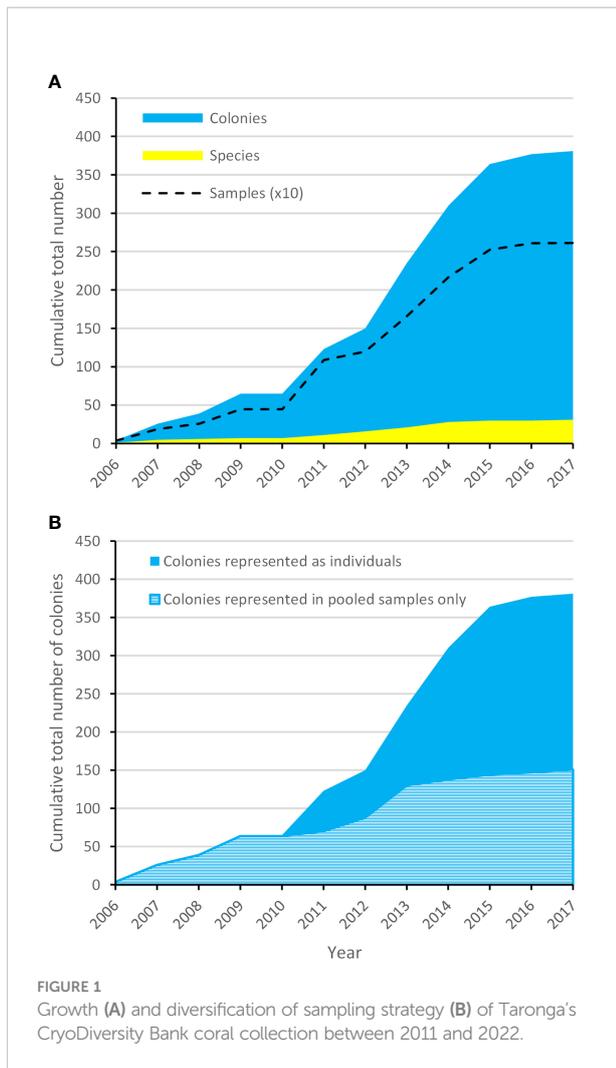
## Geographic diversity

Between 2011 and 2022, sperm samples were collected and cryopreserved from corals originating from 22 reefs across the GBR Marine Park (Figure 3).

The MA with the highest diversity of species represented in the bank is the Central MA surrounding Townsville, with 21 species sourced from at least seven reefs (Figure 3). Almost as diverse is representation from the South MA, where sperm have been cryopreserved from 15 species across five reefs in the Keppel Islands and around Heron Island. Ten of the 15

species from this MA have not had sperm samples preserved from any other region, and only 20% of all colonies represented within the bank are from this MA. Only three species from eight reefs of the North MAs are represented in the CryoDiversity Bank (*A. tenuis*, *A. hyacinthus* and *Goniastrea retiformis*).

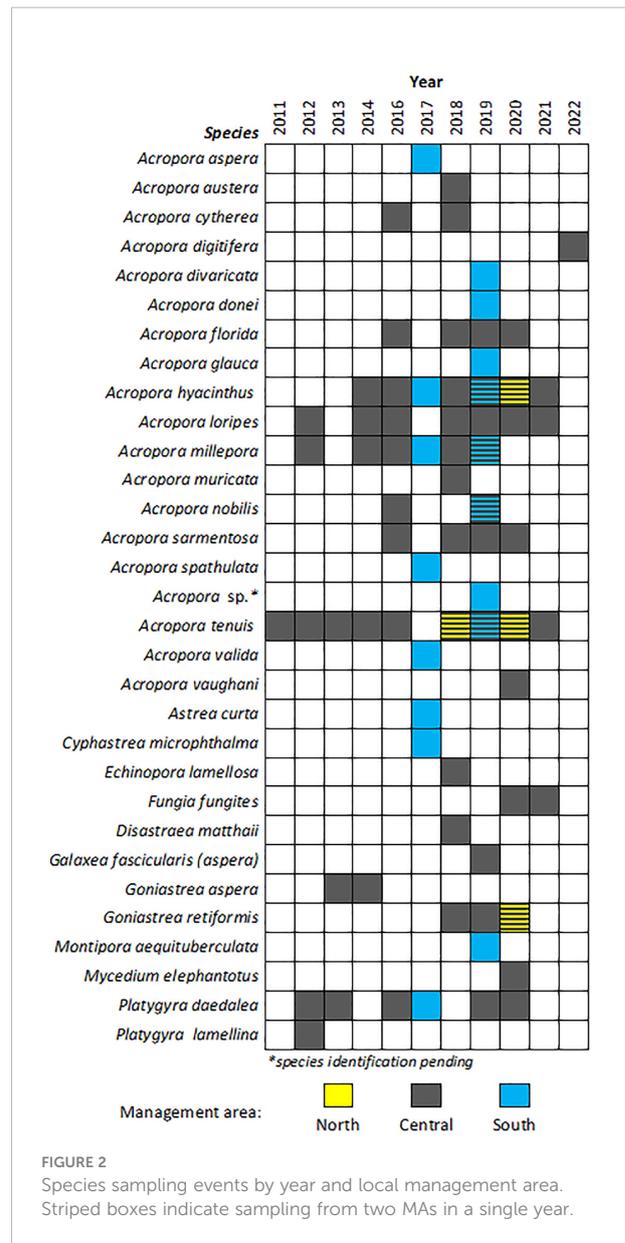
Six species from which sperm have been sampled originated from more than one MA. *Acropora tenuis* and *A. hyacinthus* are the only species with sperm samples cryopreserved across all MAs, originating from 14 and 11 reefs, respectively. Eight species were sampled from between two and six reefs and the remaining 21 species have been sampled from one reef only.



### Biodiversity and representation

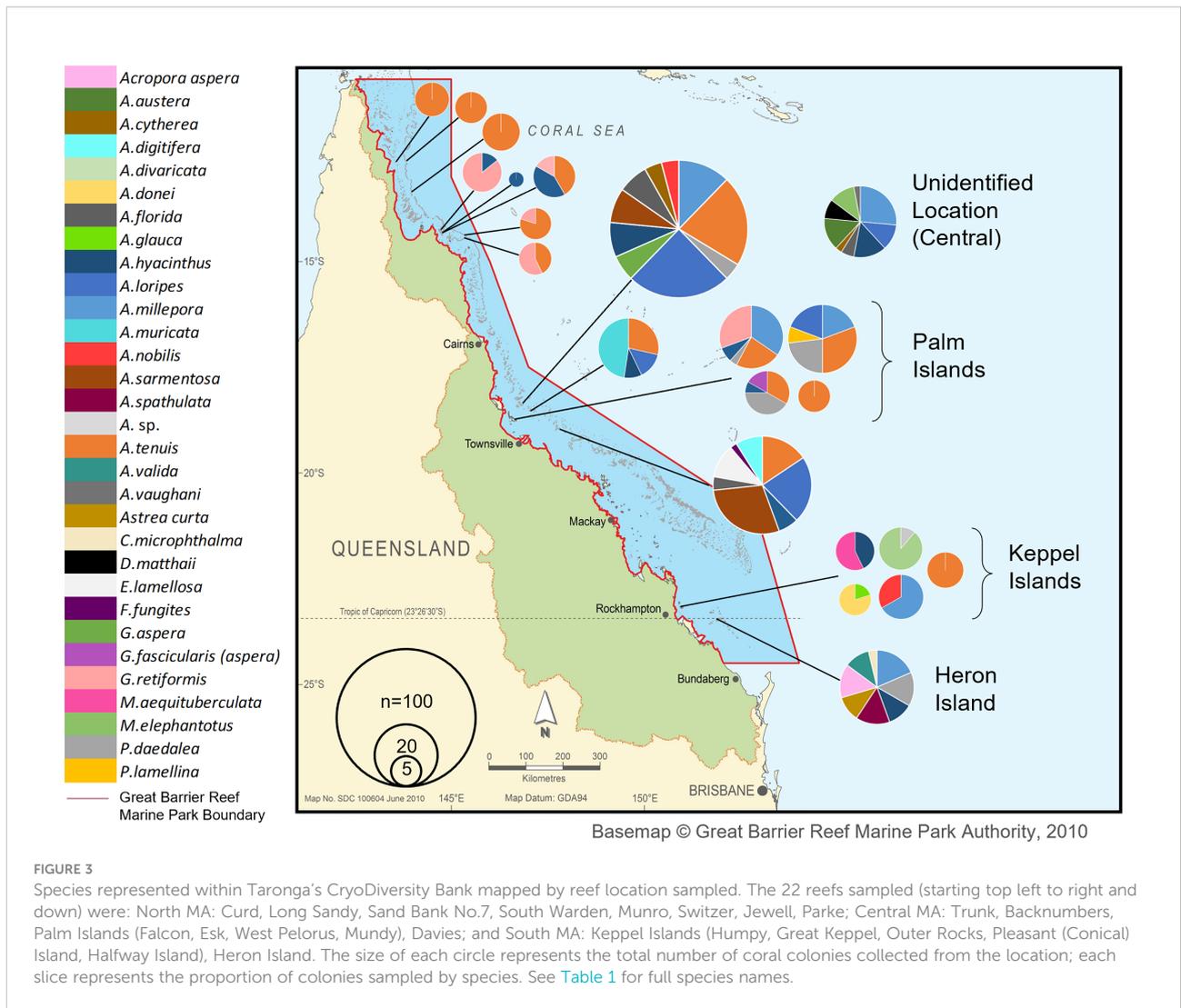
The most highly represented species in the collection, based on the total number of colonies sampled, are *A. tenuis* (25% of all colonies sampled), *A. loripes* (12%), *A. millepora* (12%), *A. hyacinthus* (9%), and *A. sarmentosa* (6%). Combined, these five species make up 63% of all colonies sampled and 71% of all cryovials in the bank. Four species (*A. divaricata*, *A. donei*, *A. vaughani*, *Cyphastrea microphthalma*) have had sperm samples cryopreserved from only one colony. Fourteen species have had two to five colonies sampled; five species have had between five and 19 colonies sampled; and seven species have had 20 or more colonies sampled (Table 1).

The most represented species in the collection is *A. tenuis* with 94 colonies represented, 62 of which have been sampled as individuals. On average, sperm samples from 6.7 (median 6.0; range 3-21) *A. tenuis* colonies have been cryopreserved from each of the 14 source reefs. *Acropora hyacinthus* is the second most



geographically diverse species in the collection (34 colonies), with an average of 3.1 (median 3.0; range 1-11) colonies sourced from 11 different reefs, although both *A. millepora* and *A. loripes* exceed it in number of colonies represented, at 46 colonies per species (Table 1).

Approximately half the species (15 species: 8 Central, 7 South) represented in the bank have 20 or fewer cryovials banked in total. Sperm from nine species are represented as pooled samples only (*A. austera*, *A. digitifera*, *A. muricata*, *Astrea curta*, *Echinopora lamellosa*, *Dipsastraea matthaii*, *Goniastrea aspera*, *Montipora aequituberculata*, *Platygyra lamellina*). Nine species were sampled only as individual genotypes and the remaining 12 species were sampled in both formats (Table 1; Figure S2).



### Sample quality metadata

Assessment of sperm quality metrics is essential in biobanking, both to determine the suitability of material for banking and to select frozen samples for end-use. Fresh and post-thaw sperm metrics are collected to assess cryopreservation efficacy. Average sperm motility of fresh samples was highly variable both within and between species (Table 1) and across years. The average fresh sperm concentration in 22 of 31 species was above 1 billion sperm cells per mL. Samples from the gonochoric species *Fungia fungites* recorded the lowest sperm concentration, at 20 million cells/mL.

### Discussion

Zoos are global leaders in the establishment of wildlife biobanks both for fundamental research and as a tool to support genetic management of *ex situ* breeding programs (Comizzoli and Wildt,

2017; Hagedorn et al., 2019; Hobbs et al., 2019; Taylor-Holzer et al., 2019; Holt & Comizzoli, 2021). Alongside biobanking, development of reproductive technologies across a range of taxa (Comizzoli, 2015; Herrick, 2019) has seen these tools more frequently integrated into breed-for-release programs and species recovery plans in recent years (Swanson et al., 2007; Howard et al., 2016; O'Brien et al., 2016; Della Togna et al., 2020; Hagedorn et al., 2021; Hobbs et al., 2021). Biobanking has been championed by these organisations for decades as a viable conservation tool that does not detract from, and should not be to the exclusion of, other conservation strategies.

Through collaborative programs over the last decade, Taronga's CryoDiversity Bank coral collection has become the

2 <https://nationalzoo.si.edu/center-for-species-survival/coral-species-cryopreserved-global-collaborators>

3 <https://taronga.org.au/conservation-and-science/current-research/reef-recovery>

largest single biorepository of cryopreserved coral sperm<sup>2,3</sup>, in both number of species represented and quantity of material. Crucially, this collection includes sperm samples that pre-date the recent mass-bleaching events on the GBR (e.g., 2016, 2017, 2020 and 2022) along with samples from colonies sourced from targeted resilient populations that have the demonstrated ability to produce heat-tolerant offspring (Quigley and van Oppen, 2022). The CryoDiversity Bank therefore provides an important reservoir of genotypic and phenotypic biodiversity that could become a unique resource for reef restoration practices, especially those that incorporate adaptation and adaptive management into their strategies (Quigley et al., 2022; Shaver et al., 2022). Our recent review showed that the Bank contains sperm samples from over 380 individual colonies of 30 identified species of scleractinian corals collected from a wide geographic area of the GBR Marine Park. Over the lifetime of Taronga's Reef Recovery Project<sup>3</sup>, most sperm samples have been collected opportunistically from colonies prioritized and sourced by our collaborators conducting their own research on a small number of coral species year-to-year. Hence, the collection is dominated (70% of samples) by five species (*A. hyacinthus*, *A. loripes*, *A. millepora*, *A. sarmentosa* and *A. tenuis*) and by reefs in close proximity to the National Sea Simulator on the Central GBR.

Sampling strategies have been suggested to support *ex situ* propagation of genetically diverse coral populations (e.g., Shearer et al., 2009; Baums et al., 2019; Quigley et al., 2019), which are directly applicable to the sampling of coral donors for sperm biobanking. One such target is the sampling of  $\geq 30$  genets per species, which captures approximately 95% of allelic diversity of an average population of unknown genetic heritage (Shearer et al., 2009). It is important to undertake genotyping of all banked samples, particularly for those species with less resolved population genetics, to confirm genetic diversity prior to potential future use of cryopreserved sperm for conservation breeding and to inform ongoing collection targets. To this end, we recommend that a fragment, or voucher sample, be taken from each coral colony and banked alongside cryopreserved sperm samples to permit future genotyping and species identification (Voolstra et al., 2021). The most highly represented species in Taronga's biobank, *A. tenuis*, easily surpasses this target with sperm samples preserved from 94 colonies. However, only six colonies have been sampled from the southern GBR where *A. tenuis* is known to have higher genetic diversity compared to northern populations (Lukoschek et al., 2016) so future effort will be focussed on bolstering genets collected from this region. For other species with fewer samples, a target of 30 genets per species and population will be prioritized. Sourcing rare and poorly represented species will remain a challenge on the GBR as the majority of its reefs are tens of kilometres from the mainland; to reach them requires significant resourcing. Engagement with Traditional Owner and other community groups to prioritize culturally significant and

biodiverse sites and species will help to ensure that limited resources have the most impact for people and communities.

At present the most effective cryopreservation methodology for coral is sperm preservation, as cryopreservation technologies for other coral cell sample types are not yet sufficiently developed for biobanking. Taronga's CryoDiversity Bank and similar groups working in Hawaii, Florida, Mexico, the Red Sea, and the Caribbean have therefore focused their acquisitions on cryopreservation of the male germline (Hagedorn et al., 2006; Grosso-Becerra et al., 2021; Zuchowicz et al., 2021). Approximately 63% of coral species surveyed are broadcast spawners (Baird et al., 2009) so the capture of sperm for most species should be possible, albeit with some notable challenges such as gonochores (e.g., *Porites* species) and species for which reproductive timing remains unknown (Baird et al., 2021). Coral sperm cryopreservation can help diversify shrinking populations through selective breeding, but currently relies on availability of fresh oocytes; we also need additional strategies to mitigate species extinctions by preserving whole organisms, or somatic cells. Cryopreservation and laser-warming of coral larvae (Daly et al., 2018) is a technology with promise but may be too complex for large-scale or field application. Newer, simpler cryopreservation technologies may provide additional opportunities for coral; for example, cryo-grid technology can easily be used to cryopreserve large numbers of *Drosophila* embryos and pancreatic cells from various species (Zhan et al., 2021; Zhan et al., 2022). Taronga's CryoDiversity Bank coral acquisitions have only recently expanded to include experimental preservation of other cell types (e.g., embryonic cells: Hagedorn et al., 2012b), life history stages (e.g., larvae: Daly et al., 2018; Cirino et al., 2019), and the microalgal endosymbionts of corals (Symbiodiniaceae: Hagedorn et al., 2010; Hagedorn and Carter, 2015; Lin et al., 2019; Kihika et al., 2022). While cryopreservation of sperm should remain a priority in the near term, the inclusion of new sample types into existing biorepositories and restoration programs will be rapidly adopted as the technologies mature and become ready for broader application.

The most efficient and effective use of cryopreserved sperm samples for reef restoration will likely be for genetic management of brood-stock within *ex situ* coral aquaculture systems, where sexually produced juveniles are generated under controlled conditions (Hagedorn et al., 2017; Randall et al., 2020) and microbiome manipulations are possible (Buerger et al., 2020; Santoro et al., 2021; Maire and van Oppen, 2022) prior to deployment onto selected reefs (e.g., Quigley et al., 2021; Randall et al., 2021). The use of cryopreserved sperm is a cost-effective way to increase the effective breeding population (e.g., Howell et al., 2021), providing greater control over brood-stock and genetic management for trait selection *via* selective crosses among individual colonies. Alternatively, pooled samples enable "batch" multi-colony fertilisation mixtures to maximise the number and diversity of larvae produced, and fit with the planned automation of *ex situ* coral spawning activities to generate restoration-scale quantities of material. Although the Taronga CryoDiversity Bank

has tended towards cryopreservation of sperm from individual colonies in recent years, future collections should aim to achieve a mix of both individual and pooled samples.

Cryopreserved sperm could also be used to overcome temporal differences in spawning between colonies. Importantly, cryopreserved sperm can easily be moved between institutions and populations to facilitate assisted gene flow (e.g., Hagedorn et al., 2021; Daly et al., 2022), reducing the cost and risk of disease transfer associated with moving coral colonies. An example of where this approach may be important is the Florida Coral Rescue program<sup>4</sup>, which has brought over 2300 colonies from 20 species of coral into the care of 27 facilities managed by 28 partners (including zoos and aquaria) in 15 states across the USA in response to extensive coral mortality in the wild from stony coral tissue loss disease (SCTLD). To manage population genetics within this *ex situ* breeding program and permit genetic crosses amongst institutions, cryopreserved sperm could be utilized to produce new corals for reef restoration and for biobanking to ensure that founder population genetics are retained (B. Firchau, pers. comm.).

Depending on the scale to which fertilization using cryopreserved sperm can be expanded, it may be possible to generate larvae directly for settlement and deployment using biobanked sperm samples. *In vitro* fertilization methods for some coral species are well established and typically involve a standard fertilization ratio of 10 eggs per ml exposed to  $1 \times 10^6$  sperm/ml, corresponding to approximately  $1 \times 10^5$  sperm per egg (Pollock et al., 2017; dela Cruz and Harrison, 2020). In the case of bulk fertilization of freshly collected gametes, fertilization volumes can reach up to 200 L for large mixed-batch cultures, generating hundreds of thousands of embryos (Negri and Heyward, 2000). Cryopreservation methods were developed to preserve 1 mL aliquots of sperm at a  $1 \times 10^9$  sperm/mL, with a predicted efficiency of 50% sperm survival and consequently a reduction in overall motility (Hagedorn et al., 2017). Existing IVF production methods using cryopreserved coral sperm (Hagedorn et al., 2021; Daly et al., 2022) are capable of producing tens of thousands of coral larvae, suitable for the genetic maintenance of brood-stock or for small-scale restoration activities (e.g. Grosso-Becerra et al., 2021), especially if combined with advanced microfragmentation and husbandry techniques (Page et al., 2022). Current protocols for sperm cryopreservation and use may therefore already be suitable for many restoration applications; however, to support the production of millions of coral larvae, different approaches to sperm cryopreservation, sample packaging, and fertilisation methods, such as those used in the agriculture and aquaculture industries, may be required.

Reef recovery programs developing and applying reef restoration practices at the scale required to help large reef

ecosystems (e.g., Bay et al., 2019; National Academies of Sciences, Engineering, and Medicine, 2019) will necessitate a prioritization of species and populations; this is especially true for the GBR, which encompasses hundreds of species across thousands of individual reefs along thousands of kilometres of coastline. Taronga's coral cryo-collection will, in part, support the production needs of an established aquaculture program (Gibbs et al., 2019; Worley Parsons Services and Australian Institute of Marine Science, 2019) whose species priorities will be informed by restoration modelling and species selection tools (Madin et al., 2021). Large-scale aquaculture of this type will likely target a smaller number of species to produce large quantities of coral recruits for direct deployment onto reefs, potentially leading to a further increase in sampling from species that are already well represented in the bank (e.g., *A. tenuis* and *A. hyacinthus*). Of equal importance, however, will be the need to establish a more diverse collection that provides a reservoir of biodiversity from which to draw once initial coral cover is restored. This strategy will warrant continued collection from lesser represented species and regions, which will be supported by competitive funding streams and philanthropy in partnership with Traditional Owners and various community groups. Striking the balance between securing broad biodiversity and storing large volumes of prioritized species to support aquaculture will be an important logistical challenge for all coral biobanking programs going forward. Modelling efforts aimed at prioritizing the array of suggested interventions (e.g., Condie et al., 2021) will also be required before we can fully understand the scale at which cryopreservation will integrate with coral production in Australia and therefore the quantities of cryopreserved material that will be required.

## Conclusions and future work

Coral reefs are complex ecosystems to conserve; ongoing applied research and modelling will be needed to improve our understanding of the necessary interventions, their scale, and their design (Sivapalan and Bowen, 2020). Based on our experience, biobanking activities should ensure that: each colony is uniquely identified and either genotyped or vouchered; collection and cryopreservation of sperm is prioritized and new sample types are introduced as technologies become available; and collections target a mix of individual and pooled samples matching production strategy and restoration goals. To achieve this will require increased resourcing for infrastructure to support the required biobanking capacity, along with continued refinement of metadata and database management to ensure that these collections can be maintained in perpetuity.

Aquaculture and biobanking on the scale required for coral conservation worldwide will require strategic partnerships

<sup>4</sup> <https://myfwc.maps.arcgis.com/apps/dashboards/eba7dc2cabc64f60819e6d4b084d94cd>

between science, industry, restoration specialists, government, and community. Taronga scientists, along with scientists from other institutions utilizing coral cryopreservation strategies globally, are collaborating to refine coral cryopreservation technologies and build capacity through groups such as the Coral Restoration Consortium (CRC) Cryopreservation and Biobanking Working Group and the Coral Biobank Alliance. It is hoped that these initiatives will help to achieve the shared goal of minimizing the loss of biodiversity on coral reefs globally.

Taronga Conservation Society commits to working in a way that respects, recognizes and includes First Nations people. Taronga will consult with Traditional Owners to ensure that the proposed cryopreservation interventions are socially and culturally safe, including that: consent is obtained to collect material from sea Country; movement of cryopreserved material to Taronga's biobank facilities is performed in collaboration with Traditional Owners of sea Country and Cammeragal (North Sydney, NSW) and Wiradjuri (Dubbo, NSW) people; and access to biobanked material only occurs within the consent provided by the relevant Traditional Owner group. Moreover, Taronga recognizes the current and ongoing heritage and spiritual connection of Traditional Owners to their sea Country and will permanently track Traditional Custodianship within the CryoDiversity Bank electronic database. Recognition of the Traditional Custodians of living cells within the biobank into perpetuity will be vital to the long-term stewardship of these samples and harmonizes with the integrated and inclusive approach to species conservation being undertaken by Taronga and RRAP.

The prospects of the Taronga CryoDiversity Bank to support reef restoration are promising; however, the reality is that the causes of climate change progress unabated. Recent fine-scale modelling (Dixon et al., 2022; Kalmus et al., 2022) predicts that an elevated temperature of 1.5°C could see <1% of coral surviving by the mid-2030s. Securing biodiversity through continued sperm cryopreservation, and the development of simple, field-ready, technologies for coral embryo cryopreservation, will be crucial to stop-gap species loss in the coming years and decades. We hope the results of our work and efforts may give future generations options for healthier, more diverse reef ecosystems.

## Data availability statement

The raw data supporting the conclusions of this article will be made available by the authors, without undue reservation.

## Author contributions

RH compiled and analysed the data, wrote the manuscript and prepared the figures. JD did the main manuscript revision and editing. MH, RS and JO'B contributed significantly to the establishment and ongoing management of the Taronga CryoDiversity Bank coral collection and reviewed the manuscript.

All other authors listed have made a substantial, direct, and intellectual contribution to the work over the life of Taronga's reef recovery program, reviewed the manuscript and approved the manuscript for publication.

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## Conflict of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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## Supplementary material

The Supplementary Material for this article can be found online at: <https://www.frontiersin.org/articles/10.3389/fmars.2022.960470/full#supplementary-material>

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