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Novel insights into the diet of southern stingrays and Caribbean whiptail rays

Running page head: Southern stingray and Caribbean whiptail diet

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ABSTRACT

Caribbean whiptail and southern stingrays are large bodied mesopredators, occupying shallow, nearshore ecosystems of The Bahamas, yet virtually nothing is known of their diet or potential resource competition. We used stomach content analysis via gastric lavage and stable isotope analysis to investigate the diet of 94 Caribbean whiptail rays (*Styracura schmardae*) and 112 southern stingrays (*Hypanus americanus*) across three locations in the central Bahamas. Gastric lavage was used to identify prey consumed, and compared to stable isotope analysis of $\delta^{15}\text{N}$, $\delta^{13}\text{C}$ and $\delta^{34}\text{S}$ of barb, muscle, blood and skin, representing different temporal integration periods. Both species appeared to consume a majority of crustaceans and annelids, although $\delta^{13}\text{C}$ values suggested that Caribbean whiptail rays have larger isotopic niche space across isotopes sampled (potentially broader diet) than southern stingrays, ranging from 5.82 to 3.43 ‰², and a greater variance in $\delta^{13}\text{C}$. This suggests Caribbean whiptail rays potentially feed on prey from either a wider geographic range, or from different habitats. Caribbean whiptail and southern stingrays are known to spatio-temporally overlap, and their isotopic niche overlapped by 35.6%. This study represents the first integrated diet estimate for southern stingrays using multiple techniques, and the first ever diet assessment in Caribbean whiptail rays. These data are critical for conservation of coastal ray species and management of coastal and nearshore environments throughout the region.

1. INTRODUCTION

Coastal and nearshore environments in the tropics support highly biodiverse and interconnected habitats, including coral reefs, seagrass meadows, estuaries and mangrove creeks, and are amongst the most highly productive ecosystems globally (Moberg & Folke 1999, Nagelkerken 2009). This interface between marine and terrestrial biomes is a dynamic space, particularly in the tropics, supporting a range of species, which themselves are essential to the function and health of ecosystems. Stingrays (Myliobatiformes) are ubiquitous and make up a significant portion of fish biomass in these environments (O'Shea et al. 2012). Through their foraging behavior, stingrays are significant agents of disturbance, physically modifying the habitat through the excavation of infaunal prey, which they regulate by exerting top-down control. As mesopredators, they are critical in creating links and energy transfer between lower and higher levels in food webs and may be considered keystone species in these environments (O'Shea 2012).

Despite their ecological importance in biological communities, half of the 630 recognised species of ray are considered data deficient by the IUCN (Dulvy et al. 2014), and this paucity of understanding can limit the design and implementation of effective conservation measures (Bland et al. 2015). Perhaps the most common species of stingray found in the tropical western Atlantic is the southern stingray (*Hypanus americanus*; hereafter referred to as SOST), a species widely distributed from the east coast of the USA to southeast Brazil (Grubbs et al. 2016). The southern stingray is important as a resource for eco-tourism, including supplementary feeding by snorkelers (Corcoran et al. 2013), but its trophic position and dietary preference is not well understood owing to a scarcity of broader trophic information. Previous studies have sampled stomach contents in limited numbers of southern stingrays (between two and 25), and suggested they are dietary generalists - predominantly consuming decapod and

1 portunid crabs, annelid and polychaete worms and teleost fish (Bigelow & Schroeder 1953,
2 Randall 1967, Snelson & Williams 1981, Smith & Herrnkind 1992, Gilliam & Sullivan 1993,
3 Bowman et al. 2000, Tilley et al. 2013a).

4
5 Conversely, the Caribbean whiptail stingray (*Styracura schmardae*; also known as the Atlantic
6 chupare stingray and hereafter referred to as CAWH) has been the subject of recent taxonomic
7 scrutiny (Carvalho et al. 2016) with its contemporary distribution in The Bahamas only being
8 described relatively recently (O'Shea et al. 2017). Even the most basic life-history data are
9 lacking for this species, increasing the complexity of challenges when understanding the role
10 and ecology of large-bodied, mesopredators within the coastal tropical western Atlantic. This
11 species is similar in appearance to the southern stingray, and it may thus have been
12 misclassified in the past. The extent to which southern stingrays and Caribbean whiptail rays
13 may overlap in space and time and compete for resources is currently unknown, although field
14 observations have revealed that the two species are sympatric, in that they have been observed
15 at congruent sites. Understanding niche overlap and resource partitioning however, will be a
16 critical step in developing management and conservation strategies for these two species and
17 the ecosystems that support them.

18
19 Assessing stomach content via gastric flushing in living animals provides insight into temporally
20 restricted prey preference, and dietary content (Hyslop 1980, Barnett et al. 2010, Heupel &
21 Simpfendorfer 2010, O'Shea et al. 2017), although limited in resolution to that of an individual's last
22 meal. In order to mitigate these limitations, multiple repeated sampling efforts can be conducted to
23 obtain this information over time from the same locations and cohort of animals. In addition, more
24 forensic approaches can be employed, for example measuring the chemical signatures of resources
25 integrated into consumer tissues, using stable isotope analysis can be a powerful tool when combined
26 with direct analysis of stomach content (SCA). Stable isotope analysis can provide relative and

indicative estimates of resource use (Shiffman et al. 2012, Hammerschlag 2019) by measuring the ratios of stable isotopes to standard forms of certain key elements, (N, C, H and O). While SIA cannot demonstrate absolute estimates of diet (Robinson et al. 2018), when used with putative samples from the environment, it can indicate likely dietary inputs (Phillips et al. 2014). In addition, the relative temporal period over which different tissues (e.g. blood, skin and muscle) reflect diet depends on the metabolic activity of the tissue sampled, for example, blood is more metabolically active than muscle tissue, and stingray barbs, which are metabolically inert once synthesized, so preserve a permanent isotopic signature until they erode or are shed (MacNeil et al. 2006). Because of this variability, MacNeil et al. (2006) suggest that estimation of trophic dynamics from a single tissue (e.g. muscle) can have considerable uncertainty. Thus in the present study, we use three different tissue types to compare the diets of the two study species.

The overall objective of this study was to discern and describe prey preference and dietary content in two sympatric species of stingray from multiple sites in the central Bahamas, and reconciling dietary overlap and potential resource partitioning using a multi analytical approach. More specifically, we aimed to describe the diets of both species, investigate putative temporal patterns in diet over time between these species and compare results generated from both stomach content and stable isotope analyses. Further, we aimed to assess dietary preference within and between species according to sex and size class (ontogeny).

2. MATERIALS and METHODS

The islands of Eleuthera and the Exuma Cays lie in the central Bahamas archipelago in the tropical western Atlantic (Fig. 1), where mangrove creek, coral reef and soft sediment expanses provide an abundance of suitable habitat for stingrays (Aguiar et al. 2009, Garrone Neto & Uieda 2012). Stingrays were captured from 23 sites between January 2015 and February 2017

over three geographic locations: i) the Schooner Cays (24.9008 °N, -76.3703 °W), a group of limestone islands surrounded by shallow sand flats of the Great Bahama Bank; ii) Powell Point, Southern Eleuthera (24.8317 °N; -76.3349 °W) a Cape lined with shallow mangrove creek inlets and with deep waters of the Exuma sound to the south; and iii) the Exuma Cays (24.7177°N, - 76.8223°W), a chain of islands along the Great Bahama Bank. Capture sites were selected following personal observations and capture success for a series of parallel studies. Both species of ray were observed at all 23 sites, but were not encountered, and therefore captured on the same day during the course of the study.

2.1 Ray capture and tissue sampling

Capture involved visually locating rays in shallow water (<1m) from a boat, before herding them by wading on foot into a seine net (10m x 1.5m), and encircling them into the net. Once an individual had been caught, a large dip net (1m diameter) was used to immobilise the stingray, which was then restrained by hand using puncture proof gloves, and the venomous barb secured with cloth and Velcro™ straps. Any pre-existing external identification tags were noted, and ray disc width (the distance between the two widest points of the stingray's pectoral fins, in mm) was measured using a flexible measuring tape. Rays were sexed using the presence or absence of claspers to indicate males. All new captured individuals were tagged in the dorsal surface of the left pectoral fin with a new identification tag (Hallprint.com) bearing a unique identification number, and a subcutaneous passive integrated transponder (PIT) tag was also administered. Prior to tissue collection, stingrays were placed into tonic immobility via dorso-ventral recumbence, resulting in a natural state of paralysis, which is thought to reduce stress during sampling procedures (Henningsen 1994). White muscle tissue samples (~1 cm²) were taken from the pelvic fins using sterilized scissors. Blood (3 ml) was extracted from the caudal vein using an 18-gauge hypodermic needle. A cartilage clipping (< 1 cm²) was taken from the

tip of the barb using sterilized scissors. Samples were kept on ice in the field to retard degradation of samples and once back at the lab were frozen immediately for storage. Prior to stable isotope analysis, samples were thawed and oven dried at 70°C for 24 hours.

2.2 Stomach content analysis (SCA)

Gastric lavage was performed following methods outlined by O'Shea et al. (2017). Briefly, a clean silicone tube (~ 10 mm diameter) was inserted through the buccal cavity into the stomach using an external marker on the ventral surface of the animal to visualise the location of the oesophageal sphincter. Sixty millilitres of ambient seawater was then introduced to the stomach via a syringe, and expelled stomach contents were captured in a tray lined with 1 mm mesh. Individuals were considered to be 'empty' after three lavage attempts (max of 180 ml of stomach flushing). Total wet mass of stomach contents was recorded to the nearest 0.1 g. Any highly digested and/or unidentifiable prey items were weighed and inspected. The remaining prey items were identified to the lowest taxonomic resolution and grouped into five phyla (Arthropoda, Annelida, Sipuncula, Mollusca and teleost fish). For each identified prey group, the number of prey items were counted and weighed to the nearest 0.1 g. Four metrics were used to describe prey; (i) numerical contribution (N_c %), calculated as total count of a given prey taxa / total count of all prey taxa x 100 (Hyslop 1980); (ii) gravimetric contribution (W_c %), calculated as total mass (g) of a given prey taxa / total mass of all prey taxa x 100 (Hyslop, 1980); and (iii) the frequency of occurrence (F_o %), calculated as total count of stomachs containing a given prey taxa / total count of stomachs with identifiable content x 100 (Vaudo & Heithaus 2011). Finally, (iv) N_c %, W_c % and F_o % were used to calculate an Index of Relative Importance (IRI (Pinkas et al. 1971)):

$$IRI = (N_c\% + W_c\%) \times F_o\% \quad (1)$$

The resulting IRI for each prey taxa (IRI_i) was then calculated as a percentage (IRI%) of the sum total of all prey taxa IRI (IRI_t), using the following equation (Cortés 1997):

$$IRI\% = (IRI_i \div \sum IRI_t) \times 100 \quad (2)$$

In order to estimate whether the number of stingrays sampled was sufficient to have described diet composition, the mean cumulative number of prey taxa found within lavage samples was plotted against the number of stingrays sampled. The number of stingrays sampled was considered to be sufficient when the curve reached an asymptote (Ferry et al. 1997).

2.3 Stable isotope analysis (SIA)

Based on the prey types recorded from stomach content analysis, prey samples (n=100) were opportunistically collected using small aquarium nets from all sites where stingrays were captured (Arthropoda n = 64, Annelida n = 14, Sipuncula n = 10 and Chordata n = 12) for context in stable isotope analyses. Tissue samples of both ray species and all prey items were freeze dried, ground to a fine powder using a pestle and mortar, and weighed into tin cups to $0.70 \text{ mg} \pm 0.05 \text{ mg}$ (for sequential $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ analysis). In a subset of 100 muscle samples, the powdered tissue was weighed into tin cups to $2.00 \text{ mg} \pm 0.05 \text{ mg}$ for $\delta^{34}\text{S}$ analysis. Earlier work suggested that chemical extraction of lipids and urea had no significant effect on $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ values in southern stingrays and Caribbean whiptail rays (Shipley et al. 2017), likely because they occur at trace levels in rays (Tilley et al. 2013a), and thus we did not remove lipids or urea before analysis. Samples were combusted in an Elementar Pyrocube purge-and-trap elemental analyser run in nitrogen, carbon and sulphur (NCS) mode, interfaced with an Elementar VisION isotope ratio mass spectrometer at the NERC East Kilbride Mass

Spectrometry Facility. Two international standards (United States Geological Survey L-glutamic acid (USGS40, analytical precision (*SD*) $\delta^{13}\text{C} = 0.07$; $\delta^{15}\text{N} = 0.16$) and IAEA-S-1, S-2 and S-3 Silver Sulphides for $\delta^{34}\text{S}$ (analytical precision (*SD*) = 0.17, 0.59, and 1.46, respectively) and three internal standards (methane sulphonamide/gelatine (MSAG2), methionine/gelatine (M2) and sulphanilamide/gelatine (SAAG2)) were run alongside the stingray samples for calibration and cross checking against other stable isotope analysis centres. Stable isotope ratios of ^{13}C to ^{12}C , ^{15}N to ^{14}N and ^{34}S to ^{32}S are expressed with delta notation (δ) as parts per million (‰) (McKinney et al., 1950), with resulting values representing the relative difference between isotopic ratios of the sample compared to a standard.

The R package Stable Isotope Mixing Models in R (SIMMR) (after Parnell et al. 2010) was used to model the proportional contribution of prey phyla to the diets of both southern stingrays and Caribbean whiptail rays ($n = 1000$ iterations per prey source), using muscle tissue samples collected from both stingray species. Arthropod prey isotopic values were highly correlated (Fig. 2), which makes discriminating between the contribution of the different prey groups challenging, so decapod shrimp, other decapods and other crustacea were grouped as ‘arthropods’ for further analyses. Following Dale et al. (2011), we used a single diet-tissue discrimination factor (DTDF's) of + 2.7 ‰ for $\delta^{15}\text{N}$, and + 0.9 ‰ for $\delta^{13}\text{C}$ for stable isotope modelling, because a single consumer tissue type was used. This value has not been validated for either Caribbean whiptail rays or southern stingrays (see Hussey et al. 2011), indeed few studies to date appear to have validated DTDFs for specific elasmobranch species (Hussey et al. 2009, Kim et al. 2012) and highlight that DTDFs can vary considerably, even within a genus (Hussey et al. 2012). We therefore cautiously apply this DTDF, accepting that future validation work may improve upon the estimates in the present study. In addition, background isotopic ratios can vary within as well as between years (e.g Scholze et al. 2003), which introduces

additional variability we were unable to account for. Ratios from muscle tissues were plotted in two-dimensional isotopic space, with DTDF adjusted isotopic ranges of the prey phyla. Sensitivity of SIMMR dietary proportion estimates to DTDFs was tested with model re-runs adjusting $\delta^{15}\text{N}$ by 0.5 and 1.0 ‰, and adjusting $\delta^{13}\text{C}$ by 0.5 and 1.0 ‰.

To test for dietary and habitat overlap between the two species, ellipses of isotopic space were generated using the R package ‘Stable Isotope Bayesian Ellipses in R’ (SIBER; [Jackson et al. 2011]). Ellipses encompass a user-defined proportion of isotopic values, and the overlapping area between two sets of isotope data can be used as a proxy for niche overlap between species (Guzzo et al. 2013). Trophic levels for prey were estimated from previous studies (Cortés 1997, Ebert & Cowley 2003, Tilley et al. 2013a). Where no trophic level was available, the closest ecological substitute for the missing prey was used. Prey trophic levels were used to calculate trophic levels for CAWH and SOST using isotope data, following (Cortés 1999):

$$TL_K = 1 + \left(\sum_{j=1}^n P_j \times TL_j \right) \quad (3)$$

Where TL_K is the trophic level of a consumer, P_j is the proportional contribution of a select prey source from the consumers diet, n is the total count of prey sources observed within the consumers’ diet, and TL_j is the trophic level of the selected prey source. This was repeated for both stingray species using prey proportion estimates from SIA.

2.4 Methodological similarity

The Czekanowski index of similarity (Feinsinger et al. 1981) was used to measure the agreement between SCA and SIA, and is calculated as:

$$CI = 1 - (0.5 \times \left(\sum_{i=1}^n P_{x_i} - P_{y_i} \right))$$

where P_{x_i} (estimate of SCA method) and P_{y_i} (estimate of SIA method) are the diet contribution estimates of the i -th prey phylum and n is the total number of prey phyla identified. The CI ranges from 0 to 1, with 0 indicating disagreement between methods, and 1 indicating perfect agreement.

2.5 Statistical analysis

Differences between the two species, two sexes and between the capture sites, were tested using Wilcoxon rank sum tests, and homogeneity of variance was tested using Fligner Killeen tests with ‘relative importance’ as the response variable. The isotopic values of the three tissue types were compared using ANOVA with repeated measures, so as to include repeated observations over time. Linear mixed effects models were used to test whether ray disc width, sex, season of capture or capture location influenced isotopic ratios and prey types consumed. Locations were grouped into Eleuthera and the Exuma Cays as there was no obvious rationale to treat them at a finer scale. To investigate whether there was a shift in habitat use for either species of stingray based on size classes, break point analysis was used to identify the disc width at which any change points in $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ occurred, using the R package ‘strucchange’. Briefly, this tests for deviations of coefficients from one stable linear regression relationship to another (i.e. where there is a structural change in the relationship between two variables), and ‘strucchange’ allows to simultaneously compute multiple breakpoints in linear regression models (Bai & Perron 2003).

All work (including stingray capture and tissue sampling) was undertaken under permits from the Bahamian Department of Marine Resources, and complied with the University of Exeter

- 1 Research Ethics framework and ethical policy, and was approved by the College of Life and
- 2 Environmental Sciences (2016/1543 and 2016/1546).

3. RESULTS

A total of 206 stingrays (94 Caribbean whiptail rays and 112 southern stingrays) were captured between January 2015 and February 2017, and muscle samples were collected from 193 individuals for stable isotope analysis (92 CAWH, 101 SOST), along with 56 blood samples (30 CAWH, 26 SOST) and 60 barb samples (30 CAWH, 30 SOST, sampling regime Fig. S1). For CAWH, 49 were female and 46 were male, and 92 of the SOST were female and 20 were male. CAWH ranged from 228 mm to 1,472 mm disc width (mean 629 mm \pm 283 s.d.), whilst SOST ranged from 342 mm to 1,102 mm disc width (674 mm \pm 175). Female southern stingrays were significantly larger than males (median 719 mm \pm 173 mm s.d. versus 517 mm \pm 67 mm s.d.; Wilcoxon rank sum; $W_{18,83} = 1270$, $p < 0.01$), but there was no difference between sexes in Caribbean whiptail rays.

3.1 Stomach content analysis

Gastric lavage was performed on 74 stingrays, of which 47 provided stomach content samples (SOST $n = 28$, CAWH $n = 19$, i.e. the remaining 27 rays had nothing in their stomachs). Prey items were identified from five phyla (Table 1; Arthropoda, Annelida, Sipuncula, Chordata and Mollusca) and could be resolved to sub-phylum level in 129 prey items. Cumulative prey curves for each species reached asymptotes after approximately 30 southern stingrays and 20 Caribbean whiptail rays had been sampled, suggesting that a sufficient number had been surveyed to represent the diet of both species (Fig. S2). Southern stingrays consumed a majority of arthropod prey (present in all 47 stingrays that had prey in their stomachs, constituting 62% of prey mass and 89% IRI, Table 1), in particular brachyuran crabs and penaeid shrimp (both present in 18% of rays). Teleost fish were observed in a third of rays sampled (constituting 8% by prey mass and 6% IRI). Southern stingrays had also consumed annelid and sipunculid worms, and two had consumed molluscs. Caribbean whiptail rays likewise consumed a

majority of arthropod prey (present in all CAWH sampled, constituting 59% of prey mass and 70% IRI, Table 1), in particular a genus of snapping shrimp (*Alpheus* spp.), which contributed almost a fifth of total prey mass and were present in nearly a third of the 19 CAWH sampled. The next most important prey were annelid worms (present in 68% of stomachs sampled, constituting 28% of prey mass and 27% IRI). Caribbean whiptail rays had also consumed decapod crustacea (present in one quarter of CAWH) and sipunculid worms (present in one fifth of CAWH) and one ray had consumed an un-identified teleost fish. It is not clear if this was captured or incidentally consumed during bioturbation, which tends to attract small fishes to the resultant bioturbated sediment. There were no molluscs in any CAWH stomach samples. None of stingray sex, disc width and capture location predicted the likelihood that any particular prey type was consumed in either species, with the exception of molluscs, which were predicted by capture location in southern stingrays (Table S2). However, this was driven by two of the 28 stingrays only, as none of the remaining 26 had consumed molluscs.

3.2 Stable isotope analysis

Stable isotope mixing models suggested that southern stingrays likely consumed mainly arthropods (mean $43\% \pm 5$ s.d.), annelid worms ($41\% \pm 4$), teleost fish ($12\% \pm 2$), and sipunculid worms ($2\% \pm 1$), while Caribbean whiptail rays consumed approximately the same proportion of arthropods (mean proportion $48\% \pm 6\% \pm$ s.d.), annelid worms ($45\% \pm 4$), and sipunculid worms (5 ± 3) as SOST, but far fewer teleost fish ($1\% \pm 1$, Fig. 2). Importantly, the putative prey samples collected did not cover the entire isotopic niche area (Fig. 2) suggesting that there may be additional prey species that both stingray species consume. Caribbean whiptail rays were predicted to feed at a marginally lower trophic level (TL 3.48) compared with southern stingrays (TL 3.51), based on isotope mixing models. The only variables affecting $\delta^{15}\text{N}$ were disc width and capture location in Caribbean whiptail rays, with larger rays

foraging at higher trophic levels (lme disc width: $F = 21.86$, $P < 0.01$, location: $F = 7.801$, $P < 0.01$, Table S1), but with a small effect size (e.g. doubling disc width brought about only a 20% increase in $\delta^{15}\text{N}$ (Fig. S3a). Capture location brought about an even smaller effect size, with a 5% difference between Eleuthera and Exuma (Fig. S3b). Likewise, the only variable that predicted $\delta^{13}\text{C}$ was capture location, with a difference of about 10% between locations ($F = 7.267$, $p < 0.01$, Table S1, Fig. S3c), and $\delta^{34}\text{S}$ was not predicted by sex, season, capture location or disc width. In southern stingrays, $\delta^{15}\text{N}$ varied by sex, with males foraging approximately 1‰ higher than females ($F = 9.67$, $p < 0.01$, Fig. S3d), and $\delta^{13}\text{C}$ varied with disc width, with larger rays foraging 2 to 3 ‰ lower ($F = 12.14$, $p < 0.01$, Table S1, Fig. S3e). $\delta^{34}\text{S}$ was not predicted by sex, season capture or disc width.

Compared to SCA, SIA indicated a larger contribution of arthropods, yet provided higher estimates for the contribution of annelid worms to the diets of both species (Fig. 3). It also substantially increases the contribution of molluscs to southern stingray diets. Agreement between methods was generally poorer for southern stingrays than Caribbean whiptail rays – SIA estimated that arthropods contributed four times less to diet compared to SCA, and molluscs four times more. Sipunculid worms were challenging to identify in gastric lavage samples, due to being soft bodied and easily digestible, but SIA suggested they were likely not important contributors to the diets of either species. Overall, there was a moderate to good agreement between dietary estimates from SIA and SCA for Caribbean whiptail rays. Of the three tissues sampled, dietary estimates from SIA of blood had the least agreement with SCA (Czekanowski index (CI) = 0.69), followed by barb (CI = 0.77), then white muscle (CI = 0.82). CI in southern stingrays varied from 0.94 for blood and SCA, followed by white muscle (CI = 0.48), and barb (CI = 0.33).

3.3 Inter-species differences

Southern stingrays had greater $\delta^{15}\text{N}$ values in white muscle tissues ($6.77 \text{ ‰} \pm 1.07 \text{ s.d.}$) than Caribbean whiptail rays ($4.82 \text{ ‰} \pm 1.06$; Wilcoxon rank sum $W_{96,102} = 8493$, $p < 0.001$, Fig. 2) and variances in $\delta^{15}\text{N}$ were similar for both species (Fligner-Killeen $\chi^2 = 0.0928$, $df = 1$, $p = 0.760$). Southern stingrays also had greater $\delta^{13}\text{C}$ values in white muscle tissues ($-8.76 \text{ ‰} \pm 1.05 \text{ s.d.}$) than Caribbean whiptail rays ($-9.31 \text{ ‰} \pm 1.59$; Wilcoxon rank sum $W_{96,102} = 6051.5$, $p < 0.01$, Fig. 2), although the effect size was small (difference between means 0.55 ‰). Caribbean whiptail rays had greater variance in $\delta^{13}\text{C}$ than southern stingrays (1.59 versus 1.05 s.d. , Fligner-Killeen $\chi^2 = 13.6$, $df = 1$, $p < 0.001$). Southern stingrays also had greater $\delta^{34}\text{S}$ in white muscle tissues ($9.20 \text{ ‰} \pm 3.82 \text{ s.d.}$) than Caribbean whiptail rays ($3.50 \text{ ‰} \pm 4.69$; Wilcoxon rank sum $W = 2068$, $p < 0.001$), but with similar variance (Fligner-Killeen $\chi^2 = 1.89$, $df = 1$, $p = 0.169$). Caribbean whiptail rays had a larger isotopic niche space than southern stingrays (as derived from SIBER, Fig. 4) for all combinations of $\delta^{13}\text{C}$, $\delta^{15}\text{N}$ and $\delta^{34}\text{S}$, ranging from 5.82 to 3.43 ‰^2), although isotopic niche space overlapped in all cases by 35.6% ($\delta^{13}\text{C}$ and $\delta^{15}\text{N}$), 41.1% ($\delta^{34}\text{S}$ and $\delta^{15}\text{N}$) and 61.5% ($\delta^{34}\text{S}$ and $\delta^{13}\text{C}$).

3.4 Inter-tissue differences

Although the relative rate of isotopic turnover between blood, muscle and barb has yet to be determined for either Caribbean whiptail or southern stingrays, values should be likely to reflect temporal variation since each of the three tissues has a very different metabolic turnover (e.g. see MacNeil et al. 2006 for freshwater stingrays). In Caribbean whiptail rays, blood had the highest $\delta^{15}\text{N}$ (mean $5.16 \text{ ‰} \pm 0.851 \text{ s.d.}$), followed by white muscle ($4.62 \text{ ‰} \pm 1.20$), then barb ($4.22 \text{ ‰} \pm 0.615$) (Fig. 5). In southern stingrays, there was no significant difference between blood ($6.59 \text{ ‰} \pm 1.02$) and white muscle $\delta^{15}\text{N}$ ($6.70 \text{ ‰} \pm 1.12$, ANOVA with repeated measures $p = 0.888$), but

barb had significantly lower $\delta^{15}\text{N}$ ($5.56 \text{ ‰} \pm 0.917$) than either blood or muscle (ANOVA with repeated measures $Z=11.3$, $p<0.001$, and $Z=11.6$, $p<0.001$, Fig. 5). In both species, blood had the highest $\delta^{13}\text{C}$ (Caribbean whiptail ray mean $11.8 \text{ ‰} \pm 1.28$ s.d., southern stingray $11.4 \text{ ‰} \pm 1.46$) followed by muscle (Caribbean whiptail ray $9.38 \text{ ‰} \pm 1.62$, southern stingray $8.95 \text{ ‰} \pm 1.20$) and then barb (Caribbean whiptail ray $8.37 \text{ ‰} \pm 1.51$, southern stingray $8.22 \text{ ‰} \pm 1.37$). However, we highlight that while inter-tissue comparisons suggest temporal differences in diet (Shipley et al. 2017), knowledge of tissue-specific fractionation and enrichment factors are almost absent for marine rays, so these values are presented for future utility, rather than as a robust reflection of temporal variability in diet for these two species.

3.5 Ontogenetic changes

There were no breakpoints in $\delta^{15}\text{N}$ for southern stingrays in any of the three tissues, but $\delta^{13}\text{C}$ decreased by approximately 2‰ in blood and muscle at 580 mm disc width (95% confidence interval: 468 to 609 mm) and 705 mm disc width (595 to 745 mm) respectively (Fig. 6). Caribbean whiptail rays, by contrast, showed several breakpoints in both isotopes (Fig. 6). Muscle $\delta^{15}\text{N}$ increased at 911 mm disc width (95% confidence interval: 770 to 1201 mm) by almost 2‰, suggesting an increase in the trophic level at which they were foraging, but there were no changes in blood or barb. $\delta^{13}\text{C}$ decreased in blood, muscle and barb at approximately 487, 568 and 503 mm disc width respectively, but by a smaller amount (1.2, 1.1 and 1.7 ‰ respectively), then, at 815 mm disc width (95% confidence interval 745 to 1,042 mm) $\delta^{13}\text{C}$ increased again by 2.14 ‰,

4. DISCUSSION

The present study is the first integrated diet reconstruction for southern stingrays using multiple techniques, and the first ever diet assessment for the Caribbean whiptail ray. Arthropods and annelid worms appear to make up the majority of prey for both southern stingrays and Caribbean whiptail rays, and they feed more infrequently on other benthic, and largely sessile prey. In comparison with previous studies of ray diet (Gilliam & Sullivan 1993, Ebert & Cowley 2003, Flores-Ortega et al. 2011, Jacobsen & Bennett 2012, 2013, Pardo et al. 2015), these results are perhaps not surprising, but nevertheless are an important confirmation of the prey preferences of these two data deficient species.

Southern stingrays appear to occupy a slightly higher trophic position than Caribbean whiptail rays but may have a similar dietary breadth. While both species occupy similar roles in the environment as mesopredators (Cortés, 1999) their diets likely overlap by one to two thirds; yet it remains unclear whether they compete for prey, or whether they can avoid it via temporal, spatial or microhabitat segregation (Schoener 1974). Caribbean whiptail rays likely feed on prey from a wider spatial distribution, potentially incorporating varying habitat, such as tidal mangrove creeks. Considering this species were captured over a much greater spatial extent than southern stingrays in the present study, this may explain some of the breadth of values.

There may also be some differences in the relative onshore-offshore foraging locations of the two species, but the difference may be ecologically negligible. Higher sulphur stable isotope values are often interpreted to indicate use of offshore, rather than estuarine, habitats (Barros et al. 2010). $\delta^{34}\text{S}$ in the present study thus suggests that southern stingrays either visit or use waters further offshore than Caribbean whiptail rays. Many elasmobranchs are also understood

to exhibit ontogenetic habitat partitioning, typically within defined ‘nursery’ areas that offer both shelter from predators and ample prey resources (Heupel et al. 2007).

Juvenile stingrays have been shown to inhabit mangrove and tidal creek areas (O’Shea et al. 2013) and adults will typically occupy offshore areas, with changes in carbon and sulphur isotope values reflecting this. Whether southern stingrays use any of these strategies to avoid competitive exclusion remains to be investigated; however, O’Shea et al. (2017) documented different size classes of Caribbean whiptail rays in mangrove creeks compared with sand flats, suggesting that mangroves may act as important refugia in the early developmental stages of this species. Data presented here related to $\delta^{34}\text{S}$ further corroborates the theory that Caribbean whiptail stingrays may occupy more coastal, estuarine environments. Furthermore, Caribbean whiptail rays will stay in the same creek for over three years (O’Shea et al. submitted) based on mark-recapture.

Sympatric species of ray have previously been demonstrated to segregate spatially, including southern stingrays and cownose rays in North Carolina, USA, (Bangle & Rulifson 2017), and blue-spotted mask rays and blue-spotted fantail rays on Australian coral reefs (O’Shea et al. 2013). Similarly, species of skates have been shown to segregate by depth (Brickle et al. 2003); however temporal resource partitioning in rays appears to be less common (Gilliam & Sullivan 1993, Corcoran et al. 2013).

4.1 Consistency over time

Using metabolically diverse tissues, the present study was able to gain some insight into longer term patterns in resource use consistency in southern stingrays and Caribbean whiptail rays (see also MacNeil et al. 2005, 2006), although it is important to note that tissue-specific diet

tissue discrimination factors should be derived for future work, and DTDF will almost certainly vary between the two species, and between tissues (Hussey et al. 2012). In addition, the isotopic signatures of prey, as well as predators, can change over time (Inger & Bearhop 2008), and we were unable to account for this in the present study. Caribbean whiptail rays may switch to forage on higher trophic level prey at around 900 mm disc width, which is thought to be around half their maximum size (Cervigón et al. 1994), but there is no evidence of this in southern stingrays. Estimated changes in foraging locations, however, may be more complex to discern in the present study. First, the effect size of the change in $\delta^{13}\text{C}$ between tissues is small in Caribbean whiptails, which may indicate variability in foraging location or season, rather than a net movement per se. Secondly, $\delta^{13}\text{C}$ can be challenging to interpret, with $\delta^{13}\text{C}$ increasing from terrestrial areas to coastal waters because $\delta^{13}\text{C}$ values of marine algae are much higher than terrestrial plants (Marshall et al. 2007), but then decreasing again moving offshore into oligotrophic waters that tend to be less $\delta^{13}\text{C}$ rich (Hobson et al. 1994). While size at maturity for this species is still unknown, it is likely that these small shifts in $\delta^{15}\text{N}$ representing a transfer to higher trophic prey at 900 mm disc width may reflect an ontogenetic shift; but further investigation would be required to validate this, including size at maturity. Breakpoints in $\delta^{13}\text{C}$ values for CAWH individuals is relatively well defined further lending validation that an ontogenetic shift in prey consumption may be occurring, but whether this is a result of growing larger with subsequent access to larger prey, or if it represents a shift in habitat use and association cannot be discerned without further investigation.

Given that stingrays in the present study likely move from mangroves (which extend into terrestrial zones) to offshore cays, the direction of change in $\delta^{13}\text{C}$ does not yet yield a clear understanding of their movement. Further work assessing the differences in $\delta^{13}\text{C}$ between these habitats would help, as would tracking stingrays of both species, for example with acoustic

telemetry, light geolocation tags or time depth recorders (Hussey et al. 2015) to assess the extent to which they move within a normal season (Cartamil et al. 2003). It also remains possible that some of the isotopic differences between the species are an artefact of the locations at which they were captured, with Caribbean whiptail stingrays captured over a much larger range than southern stingrays. The capture location may vary isotopically, but with few putative prey samples from each, we were unable to robustly resolve this in the present study. Further, it is unclear how far either southern stingrays or Caribbean whiptail rays move over the scale of months to years (but see Tilley et al. 2013b) and how this may therefore affect the integration of isotopic values, should they vary by site.

4.2 Methodological aspects

Studying the diet of wild sharks and rays has been predominantly approached using stomach content analysis (Hyslop 1980, Barnett et al. 2010, Heupel & Simpfendorfer 2010). More recently stable isotope analysis has been adopted as a less invasive way to infer diet in wild animals (Hussey et al. 2012, Shiffman et al. 2012, Hammerschlag 2019). It is important to note that SIA alone does not determine diet, but rather reflects relative differences, and in combination with prey sampling, can infer likely composition. The present study highlights that there can be considerable differences in the results produced by SIA and stomach content analysis, although the most important prey taxa will likely still be resolvable. The present study parsimoniously concludes that arthropod and annelid prey are likely important for Caribbean whiptail and southern stingrays, but exact contributions to diet should not be inferred. Further experimental work is needed to parameterise the rate at which $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ stable isotopes turn over in different tissue types in rays (e.g. MacNeil et al. 2005), which would improve longer-term insights into diet. In addition, future studies should attempt to collect sufficient breadth of putative prey species to cover the entire isotopic niche area.

1 *Conclusion*

2 Results presented here offer a critical insight into the trophic ecology and position between two
3 species of large bodied stingray from The Bahamas. Given that rays represent the most
4 vulnerable group of cartilaginous fishes globally (Dulvy et al. 2014), and the conservation
5 status of these species remains data deficient, this is a critical step in elucidating elements of
6 their life history to better inform the effective management of species and the habitats that
7 support them. This is particularly relevant for the Caribbean whiptail ray which is considered
8 rare throughout its known range (Nunes & Nunes 2020), and provides the very first assessment
9 of its trophic ecology. Finally, understanding the dietary niche preferences of co-occurring
10 species allows for a better understanding when assessing the ecological gradients at which the
11 partitioning of resources occurs between and among tropical batoids.

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FIGURES

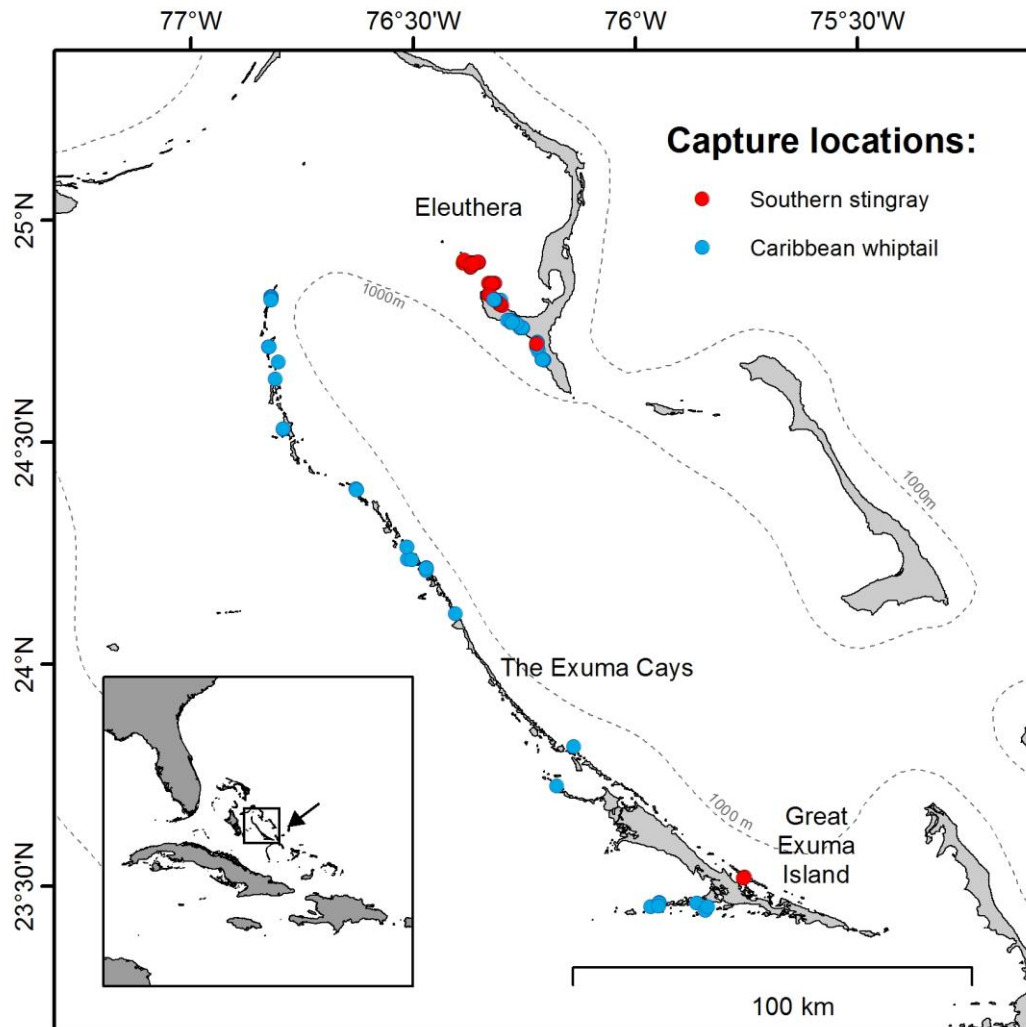


Figure 1: Map showing the capture locations of Caribbean whiptail rays (blue points) and southern stingrays (red points) at sites surrounding the Exuma Cays, and Eleuthera, The Bahamas.

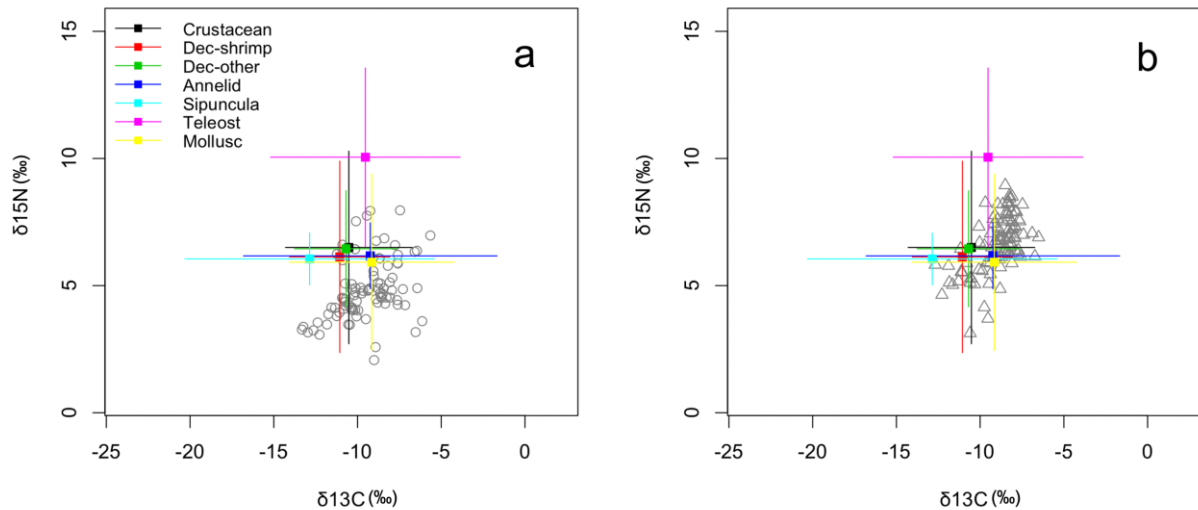


Figure 2: Bivariate scatter plot of $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ values for white muscle samples from (a) Caribbean whiptail rays and (b) southern stingrays, after applying a diet tissue discrimination factor of + 2.7 ‰ for $\delta^{15}\text{N}$, and + 0.9 ‰ for $\delta^{13}\text{C}$. The $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ values (mean \pm s.d. in both dimensions) for seven putative prey groups (Decapod shrimp, other decapods and other crustacea are detailed separately, but grouped elsewhere as arthropods) from samples collected at the same time and locations as the rays were captured are shown (Arthropoda in black, red and green, Annelida in dark blue, Sipuncula in turquoise, Teleost fish in pink and Mollusca in yellow).

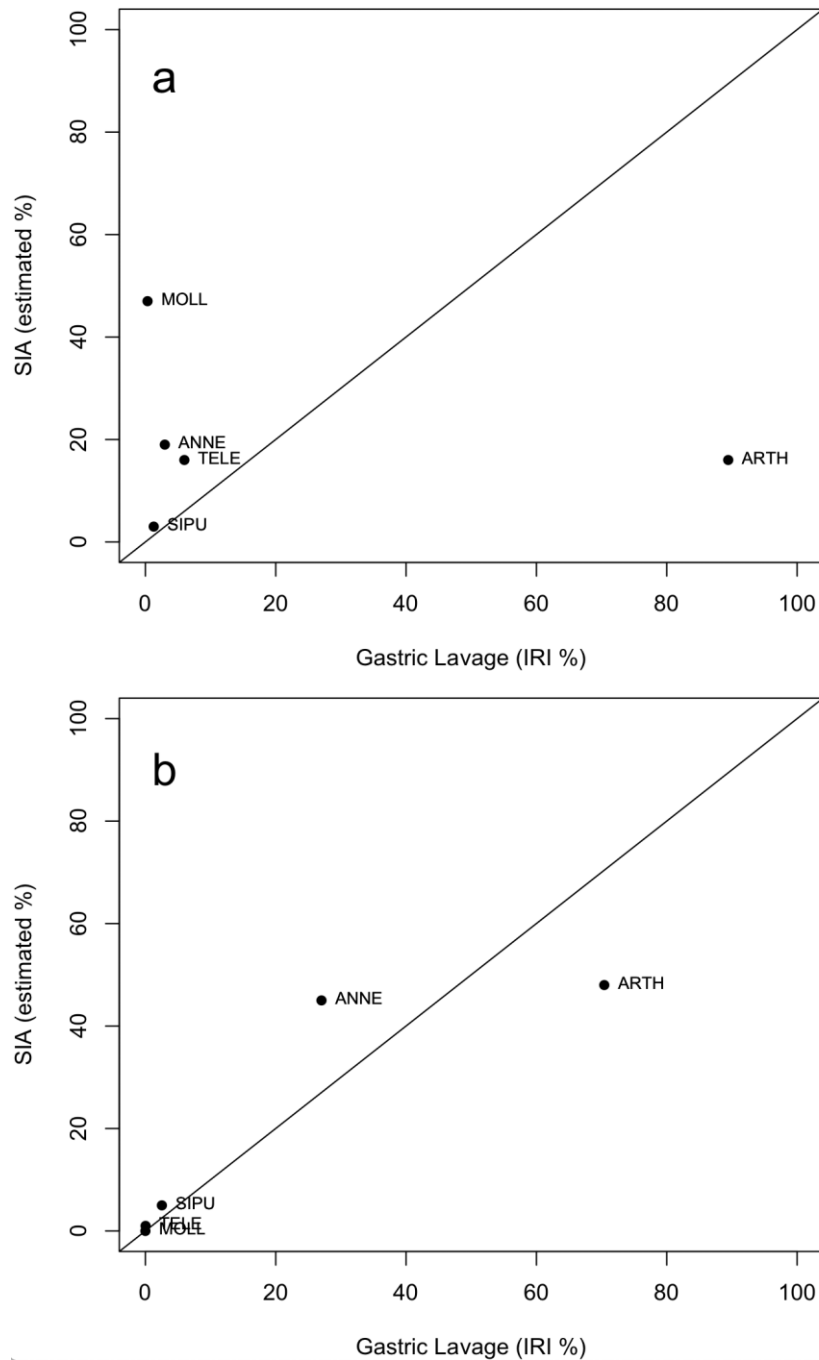


Figure 3. Plot showing estimated importance of five prey phyla (ARTH = arthropods, ANNE = annelid worms, SIPU = Sipunculid worms, TELE = teleost fish and MOLL = molluscs) estimated by SCA and SIA for a) Southern stingrays and b) Caribbean whiptail rays. Diagonal line shows line of equivalence, where SCA and SIA would produce the same result.

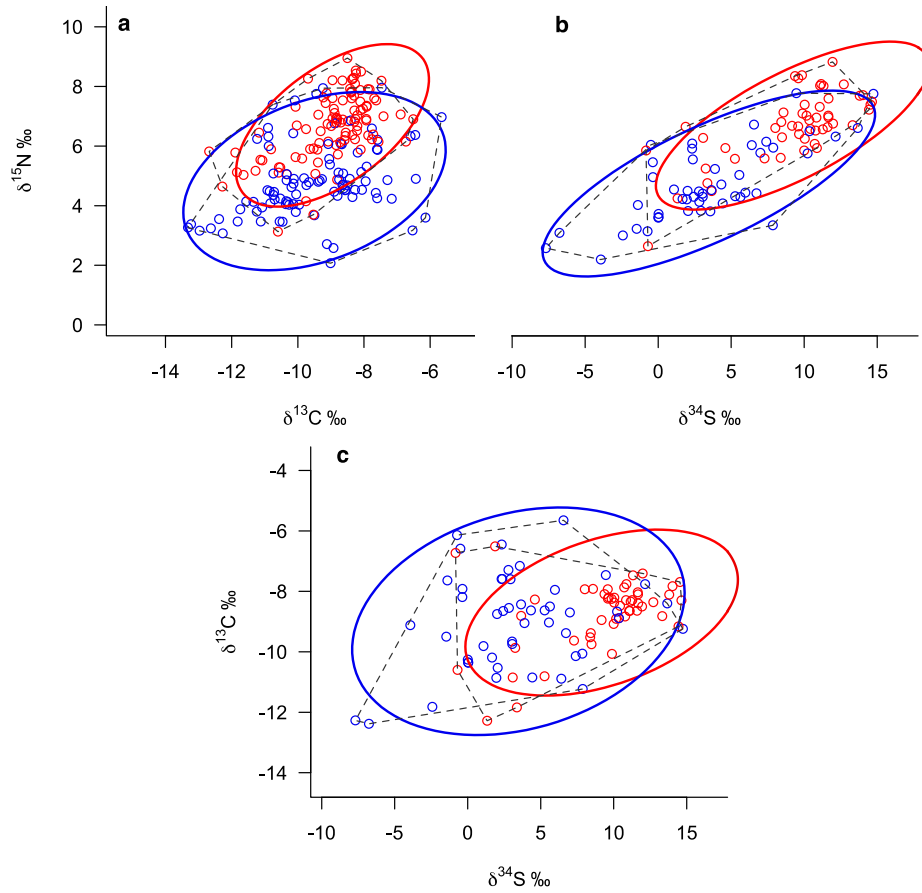


Figure 4: Bivariate isotope plots showing Standard Ellipses of isotopic spaces from white muscle samples of Caribbean whiptail rays (blue) and southern stingrays (red) for (a) $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$, (b) $\delta^{15}\text{N}$ and $\delta^{34}\text{S}$, and (c) $\delta^{13}\text{C}$ and $\delta^{34}\text{S}$. Solid lines enclose standard ellipse areas (SEA) for each species which could be used to represent the total isotopic space occupied by each species. Dashed lines represent convex hulls which encompass all data points for each species.

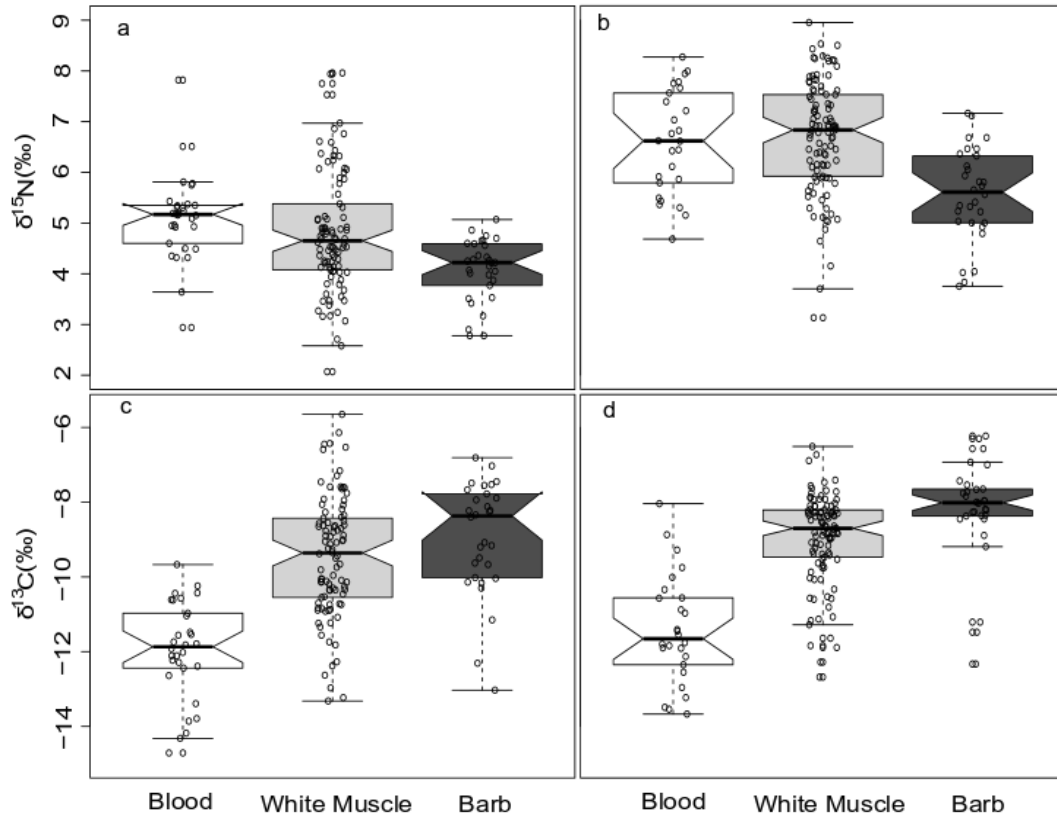


Figure 5: Boxplots showing the distribution of $\delta^{15}\text{N}$ (a and b) and $\delta^{13}\text{C}$ (c and d) isotope values for Caribbean whiptail ray (left) and southern stingray (right) blood (white), white muscle (light grey), and barb (dark grey) tissues. Boxes show interquartile ranges, horizontal line shows median value and notches indicate statistical differences.

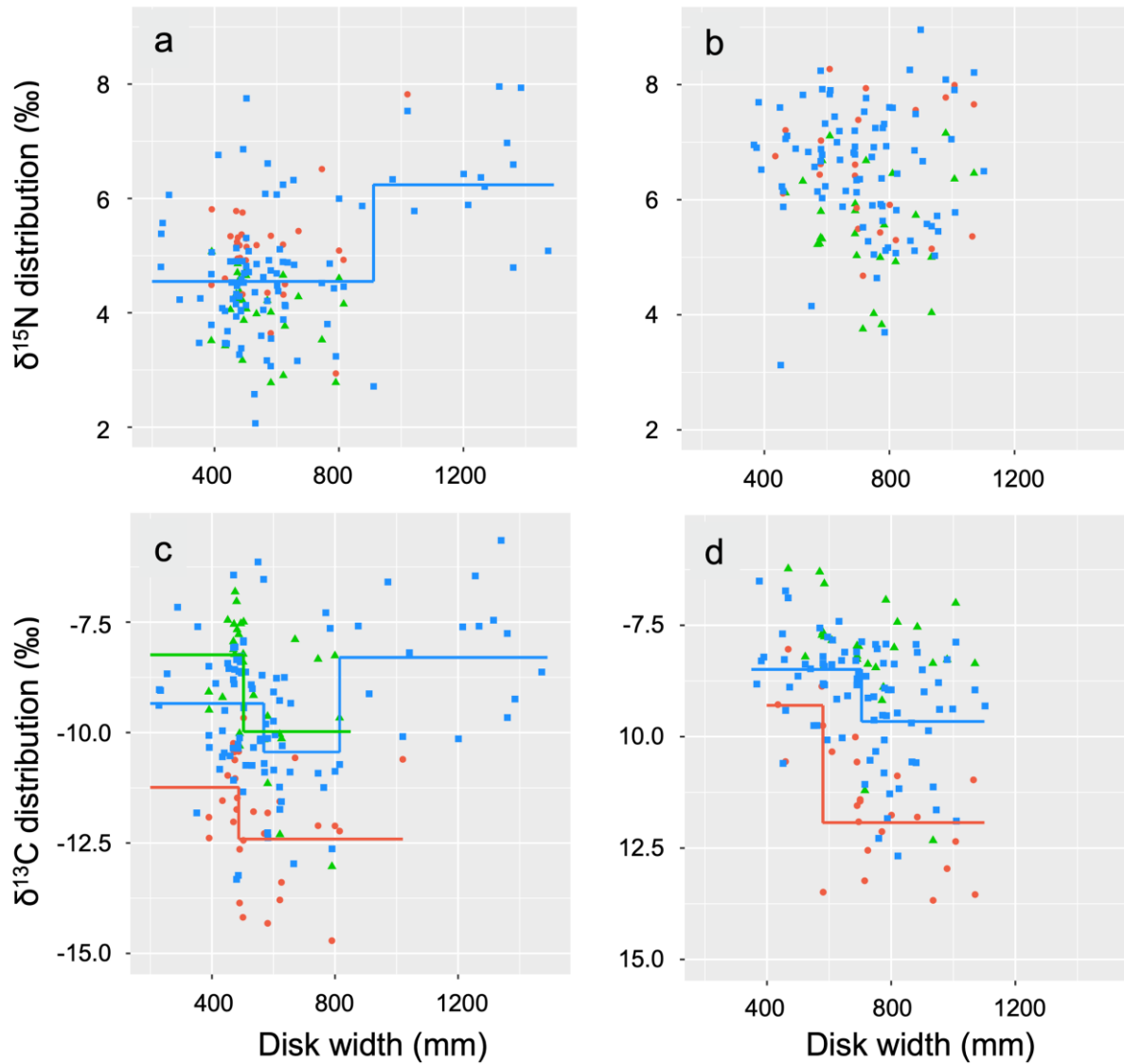


Figure 6: Scatterplots showing breakpoints in $\delta^{15}\text{N}$ (top row) and $\delta^{13}\text{C}$ (bottom row) and disc width for Caribbean whiptail rays (left panel) and southern stingrays (right panel). Points of different shape and colour depict muscle (blue squares), blood (red circles), and barb (green triangles). Breakpoints are demonstrated by vertical changes in solid lines (disc width point at which the mean isotopic value shifts).

Table 1: Dietary compositions of Caribbean whiptail rays and southern stingrays as from stomach content analysis. Identified prey items are listed below the appropriate phylum and noted with the lowest taxonomic level to which they were identified; phylum (p), sub-phylum (s-p), infraclass (ic), order (o), infraorder (io), family (f) and genus (g). Prey proportions are represented as percentage numerical composition (N_c %), percentage weight composition (W_c %), percentage frequency of occurrence (F_o %), and overall percentage index of relative importance (IRI %).

	Caribbean whiptail rays				Southern stingrays			
	<i>N_c</i> %	<i>W_c</i> %	<i>F_o</i> %	<i>IRI</i> %	<i>N_c</i> %	<i>W_c</i> %	<i>F_o</i> %	<i>IRI</i> %
Arthropoda	51.38	58.63	100	70.41	67.57	62.46	100	89.43
Crustacea ^(s-p)	1.83	2.30	10.53	0.69	17.57	16.56	35.71	32.28
Stomatopoda ^(o)	2.75	1.92	10.53	0.78	6.76	1.91	14.29	3.28
Decapoda ^(o)	18.35	13.09	26.32	13.06	1.35	3.32	3.57	0.44
Brachyura ^(io)	3.67	1.65	10.53	0.88	10.81	5.46	17.86	7.69
Portunidae ^(f)	0	0	0	0	5.41	2.60	14.29	2.02
Diogenidae ^(f)	0	0	0	0	1.35	0.33	3.57	0.16
Palaemonidae ^(f)	6.42	16.13	15.79	5.62	5.41	3.07	10.71	2.40
Penaeidae ^(f)	0.92	4.23	5.26	0.43	9.46	16.00	17.86	12.04
<i>Alpheus sp.</i> ^(g)	17.43	19.30	31.58	18.31	5.41	5.50	3.57	1.03
<i>Metapenaeopsis sp.</i> ^(g)	0	0	0	0	4.05	7.72	7.14	2.23
Annelida	40.37	27.83	68.42	27.02	10.81	4.37	21.43	2.98
Annelida ^(p)	29.36	26.68	57.89	51.22	10.81	4.37	21.43	8.62
Arenicolidae ^(f)	11.01	1.15	10.53	2.02	0	0	0	0
Mollusca	0	0	0	0	2.10	5.91	7.14	0.33
Mollusca ^(p)	0	0	0	0	2.70	5.91	7.14	1.63
Sipuncula	7.34	13.41	21.05	2.53	5.41	19.51	10.71	1.28
Sipuncula ^(p)	7.34	13.41	21.05	6.90	5.41	19.51	10.71	7.07
Chordata	0.92	0.13	5.26	0.03	13.51	7.75	32.14	5.98
Teleostei ^(ic)	0.92	0.13	5.26	0.09	13.51	7.75	32.14	18.10

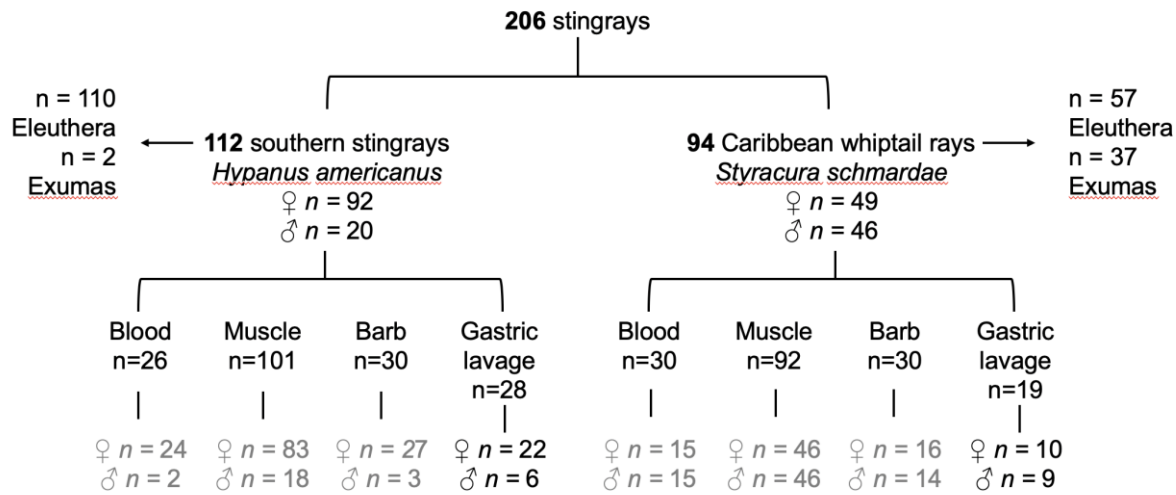


Figure S1: Sampling regime and sample numbers of southern stingrays and Caribbean whiptail rays for stable isotope analysis (all samples except ‘gastric lavage’). A subset of 100 ray muscle samples (48 Caribbean whiptail and 52 southern stingrays) were analysed for $\delta^{34}\text{S}$.

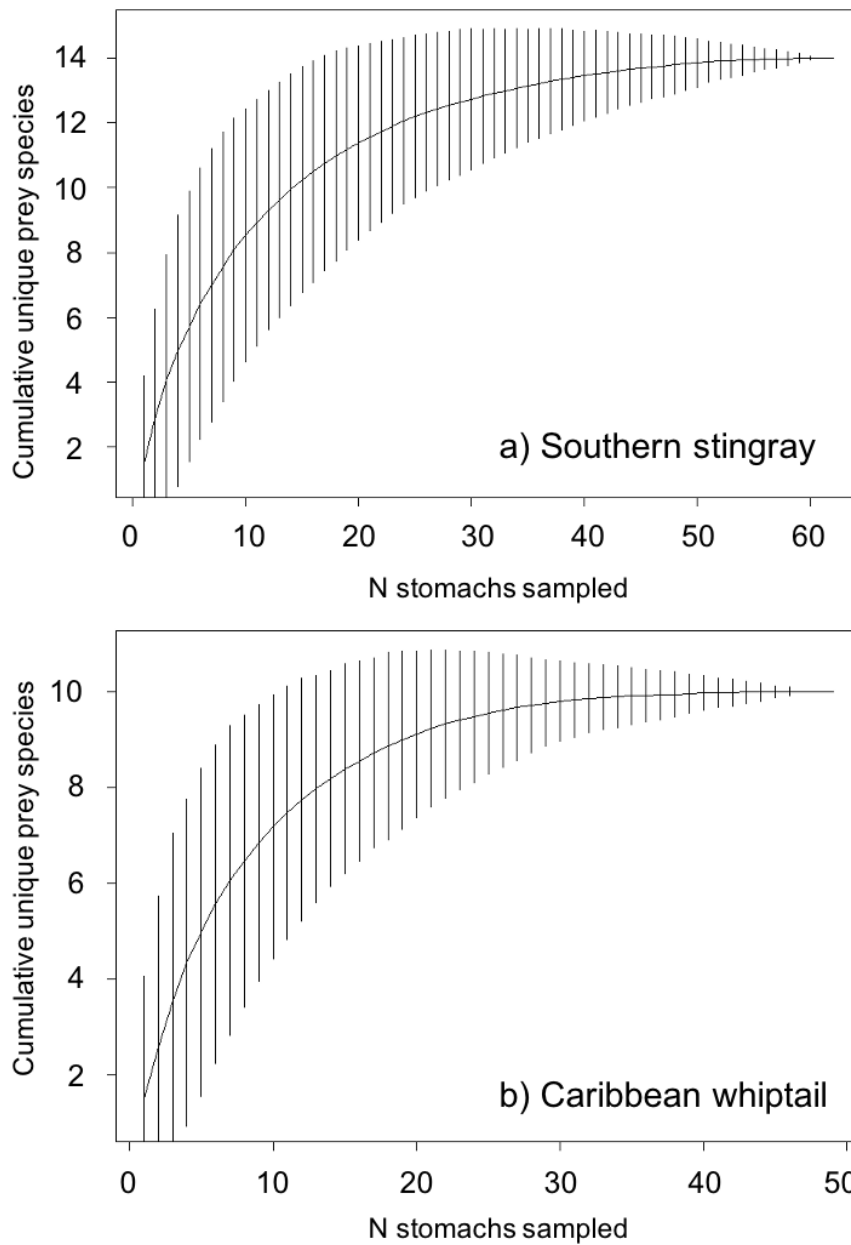


Figure S2. Plots showing the mean cumulative number of unique prey species discovered after sampling stomach contents of a) southern stingrays and b) Caribbean whiptail stingrays. The order in which individuals were sampled was randomised, and the number of cumulative unique prey discovered was resampled 1000 times, and the mean value calculated across runs. Vertical bars denote standard deviation around each mean value. A sufficient number of sampled stomachs is represented with an asymptote in the graphs curve (Ferry et al. 1997).

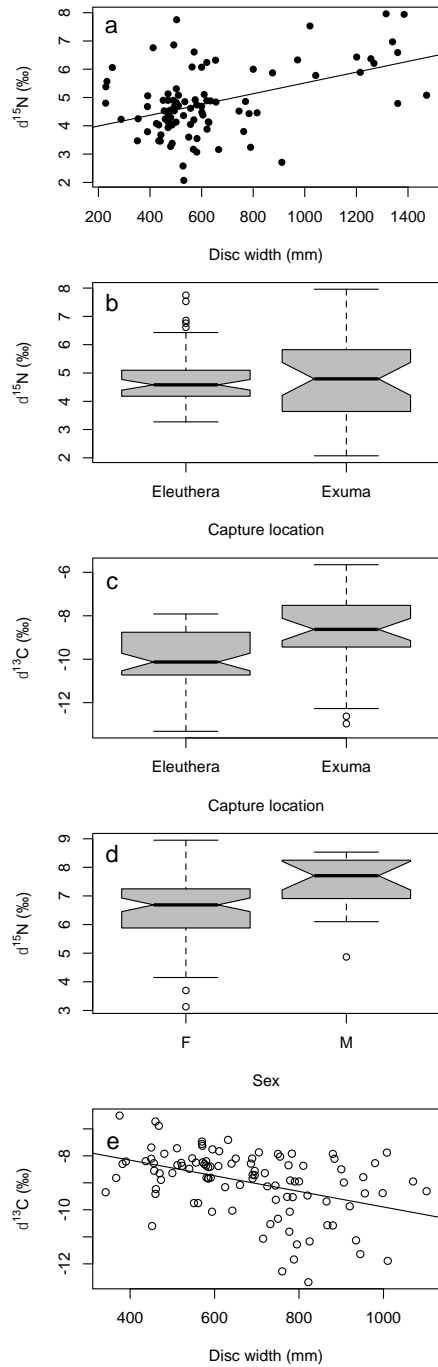


Figure S3: The relationship between stingray disc width (in mm, a, e), capture location (b, c) or sex (d) and $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ isotope values in white muscle of Caribbean whiptail rays (a, b, c) and southern stingrays (d, e), showing variables that significantly predict isotope values (see also Table S1). Parts (b, c and d) are boxplots, where notches indicate statistically significant differences between the two datasets. Lines on (a) and (e) show significant least squared linear regression relationships.

Table S1: Linear modelling of factors (sex, season of capture, disc width and Island) that influence isotopic values of $\delta^{15}\text{N}$, $\delta^{13}\text{C}$ and $\delta^{34}\text{S}$ from white muscle tissue. Bold text indicates statistical significance. Island was not used in linear modelling for southern stingrays because only two of 112 were caught outside of Eleuthera.

	$\delta^{15}\text{N}$		$\delta^{13}\text{C}$		$\delta^{34}\text{S}$	
	F	p-value	F	p-value	F	p-value
Southern stingray						
Sex	9.67	<0.01	0.981	0.324	0.269	0.606
Season of capture	3.367	0.069	1.902	0.171	0.417	0.522
Disc width	0.014	0.907	12.137	<0.01	0.239	0.627
Island	(-)	(-)	(-)	(-)	(-)	(-)
Caribbean whiptail						
Sex	0.699	0.405	0.014	0.907	1.041	0.313
Season of capture	3.274	0.074	0.332	0.566	1.351	0.251
Disc width	30.112	<0.01	1.206	0.275	0.923	0.342
Island	7.807	<0.01	7.267	<0.01	1.770	0.190

Table S2: Linear modelling of factors (sex, season of capture, disc width and Island) that influence the relative importance of Arthropods, Annelid worms, molluscs, sipunculid worms and teleost fish to the diets of southern stingrays and Caribbean whiptail rays as determined from stomach content analysis. Bold text indicates statistical significance. Mollusca was not used in linear modelling for Caribbean whiptail rays because they were no identified in any stomach samples.

	Arthropoda		Annelida		Mollusca		Sipunculid		Teleost	
	F	p-value	F	p-value	F	p-value	F	p-value	F	p-value
Southern stingray										
Sex	0.576	0.455	1.992	0.171	0.044	0.835	2.661	0.116	0.010	0.919
Disc width	1.231	0.278	0.112	0.741	1.657	0.210	0.417	0.525	0.746	0.397
Island	0.177	0.678	0.060	0.809	5.624	<0.05	0.023	0.880	0.249	0.622
Caribbean whiptail ray										
Sex	0.585	0.456	1.839	0.195	(-)	(-)	1.266	0.278	0.780	0.391
Disc width	4.359	0.054	3.429	0.084	(-)	(-)	0.491	0.494	0.423	0.530
Island	0.151	0.703	1.267	0.278	(-)	(-)	1.295	0.273	0.084	0.777

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