

# Trophic ecology of three sympatric batoid species (*Dasyatis pastinaca, Raja clavata,* and *Raja maderensis*) from the Azores, NE Atlantic

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Abstract Describing the trophic structure and interactions of demersal elasmobranch assemblages is fundamental to understanding food web dynamics and developing ecosystem-based management approaches. Stomach content analysis (SCA) and stable isotope ratios (SIA) of carbon ( $\delta^{13}$ C) and nitrogen ( $\delta^{15}$ N) from muscle were used to examine the dietary habits and intra- and interspecific trophic ecology of three sympatric batoid species (*Dasyatis pastinaca, Raja clavata,* and *Raja maderensis*) from the Azores, Northeast Atlantic. Data were analyzed with respect to sex and maturity stages. SCA showed that *D*.

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P. Fontaine (⊠) · S. Jaquemet UMR Entropie, Faculté Des Sciences Et Technologies, Université de La Réunion, 15, Avenue René Cassin - CS 92003, 97744 Saint Denis Cédex 9, Réunion Island, France e-mail: 40007542@co.univ-reunion.fr

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cE3c- Centre for Ecology, Evolution and Environmental Changes, Azorean Biodiversity Group, CHANGE – Global Change and Sustainability Institute, Faculty of Agricultural Sciences and Environment, University of the Azores, Rua Capitão João d'Ávila, Pico da Urze, 9700-042 Angra Do Heroísmo, Portugal e-mail: joao.ps.barreiros@uac.pt pastinaca feeds mostly on crustaceans, whereas R. clavata and R. maderensis prey almost exclusively on teleosts, but not on the same species. Dasyatis pastinaca displayed higher  $\delta^{13}$ C and lower  $\delta^{15}$ N values compared to R. clavata and R. maderensis. Trophic niche breadth was variable, D. pastinaca and R. clavata had the broadest and the narrowest trophic breadth, respectively. Relative trophic position categorized D. pastinaca as a mesopredator, while R. clavata and R. maderensis occupied higher trophic positions. With size, R. clavata and R. maderensis shifted from small prey such as crustaceans to larger prey such as teleosts, and they also exhibited significant increases in  $\delta^{15}N$ with size. Dietary and isotopic overlap was overall low among species, but it was higher between R. clavata and R. maderensis, suggesting more similarity in diet and habitat use between them than with D. pastinaca. This study depicts trophic interactions and functional roles of three co-existing batoid species in the Azorean food webs. In addition to presenting new information on the trophic ecology of D. pastinaca and R. clavata, the present study provides, to our knowledge, the first description of the diet composition and trophic level of the Macaronesian endemic batoid R. maderensis.

**Keywords** Elasmobranch · Feeding ecology · Stomach contents · Trophic tracers · Macaronesia

# Introduction

Elasmobranchs often act as apex or mesopredators in marine food webs (Heupel et al. 2014). In coastal marine environments, batoids (superorder Batoidea comprising skates, stingrays, electric rays, and guitarfishes) are one of the most abundant and diverse groups of elasmobranchs, with ca. 633 species worldwide (Last et al. 2016; Flowers et al. 2021). They are important components of soft bottoms and vegetated marine systems, not only through predation (Grubbs et al. 2016; Barrios-O'Neill et al. 2017; Flowers et al. 2021) but also through bioturbation, a key process in biogeochemical cycles (Laverock et al. 2011; Takeuchi and Tamaki 2014; Flowers et al. 2021). While research on elasmobranchs has resulted in a comprehensive understanding of shark trophic ecology and their role in ecosystems (Heupel et al. 2014), such knowledge remains limited for batoids (Vaudo and Heithaus 2011). Additionally, although quantitative dietary information for batoids has progressively improved in recent years, there remains a lack of studies for many existing species in some areas, such as the Mediterranean and the Northeast Atlantic (Das and Afonso 2017; Coll-Calvo et al. 2020). The food preferences of batoids can vary regionally as a consequence of their opportunist/generalist foraging behavior in which prey availability and competition for trophic resources are important factors to determine the trophic niche of these elasmobranchs (Flowers et al. 2021). Additionally, batoids are characterized by a slow growth, late maturity and low fecundity, making their population extremely vulnerable to decline consequent to overexploitation and habitat destruction (Stevens et al. 2000; Dulvy et al. 2014).

Elasmobranch trophic ecology research has traditionally been based on stomach content analysis (SCA) (Hyslop 1980; Cortés 1999; Ebert and Bizzarro 2007). SCA provides invaluable information about the prey species consumed by the studied species (Hyslop 1980; Cortés 1999); however, the method is time consuming, typically lethal, and only captures short-term diet information (Hyslop 1980; Hussey et al. 2011). The analyses of stable isotopes (SIA) of carbon ( $\delta^{13}$ C) and nitrogen ( $\delta^{15}$ N) have been used as complemental tools to investigate long-term dietary information (Vaudo and Heithaus 2011; Shiffman et al. 2012). The stable isotope of carbon ( $\delta^{13}$ C) is used to track movements between isotopically distinct habitats and to identify different sources of carbon, whereas the stable isotope of nitrogen ( $\delta^{15}N$ ) reveals relative trophic position within a food web (Hussey et al. 2011; Kinney et al. 2011). The combination of stomach contents and stable isotope analyses ( $\delta^{13}C$  and  $\delta^{15}N$ ) provides a robust description of the trophic ecology of elasmobranchs within a food web (Shiffman et al. 2012).

The Azores islands, the north-westernmost archipelago of the Macaronesia region, is spread across 600 km in the Northeast Atlantic (Santos et al. 1997). The absence of an insular shelf limits fishing effort to the island's slopes and the surrounding seamounts within the area (Da Silva and Pinho 2007; Diogo et al. 2015). Despite the fact that the Azores archipelago hosts a rich marine fauna and is a hotspot of diversity of elasmobranchs (41 shark species and 16 batoid species) (Barreiros and Gadig 2011; Afonso et al. 2020; Santos et al. 2020), studies investigating the trophic habits of batoids remain scarce in this area (Morato et al. 2003; Ponte et al. 2016; Torres et al. 2016a; Santos et al. 2020). In the Azores, most of the literature on the trophic ecology of batoids focuses on the description of prey items in guts, while only one study is based on carbon and nitrogen stable isotope analysis (Torres et al. 2016a). In addition, previous studies usually describe the diet of one species, while studies on inter- and intraspecific interactions remain limited (Morato et al. 2003; Ponte et al. 2016). On the other hand, many batoids have declined in abundance across the Azores waters, and most of them are considered threatened or near-threatened according to the IUCN (Das and Afonso 2017). Most batoids captured in the Azores are not directly targeted but included in multispecies fisheries or are bycatch in the few traditional fisheries targeting commercially valuable teleost species (Torres et al. 2016b).

In the coastal zone of the Azores archipelago, skates and rays comprise a large proportion of the elasmobranch community (16 species) (Barreiros and Gadig 2011; Afonso et al. 2020; Santos et al. 2020). The common stingray *Dasyatis pastinaca* (Linnaeus 1758), the thornback ray *Raja clavata* (Linnaeus 1758), and the madeiran ray *Raja maderensis* (Lowe 1838) are three batoid species occurring in the region (Barreiros and Gadig 2011; Afonso et al. 2020). *Dasyatis pastinaca* occupies benthopelagic habitat down to 200 m deep but seems to be more common in shallow shelfs (<50 m)

(Ponte et al. 2016; Santos et al. 2020). This species is common in the Azores, and is also found from southern Norway to South Africa and the entire Mediterranean (Barreiros and Gadig 2011; Ponte et al. 2016). Raja clavata is a shallow-water, bottom-living elasmobranch whose distribution extends from Iceland to South Africa and throughout the Mediterranean (Morato et al. 2003; Barreiros and Gadig 2011). In the Azores, R. clavata is frequent in the shallow shelf slope/shelf break assemblage (<200 m) over sandy grounds and represents more than 90% of the biomass of the landed skates in this region (Barreiros and Gadig 2011; Santos et al. 2020, 2021). Raja maderensis is a poorly known endemic species, restricted to waters close to Madeira and the Azores (Barreiros and Gadig 2011). The species is rare in the Azores and conversely to D. pastinaca and R. *clavata*, it belongs to the upper slope fish assemblage (200-600 m) (Santos et al. 2020). R. clavata is classified as Least Concern (NT) worldwide (Ellis et al. 2016), while D. pastinaca and R. maderensis are classified as Vulnerable (VU) (Dulvy et al. 2015; Jabado et al. 2021).

Although elasmobranchs can be fundamental to the structure and functioning of marine food webs (Navia et al. 2017), especially batoids, few studies have described their diets in the Azores (Morato et al. 2003; Ponte et al. 2016; Torres et al. 2016a). The diet of R. maderensis particularly remains poorly documented throughout the distribution range of the species. Prior studies described D. pastinaca as a broad spectrum mesopredator exhibiting epibenthic feeding habits with a diet consisting of large proportions of decapod crustaceans (Ponte et al. 2016). In contrast, *R. clavata* in the Azores preys heavily upon fish at a higher trophic level (Morato et al. 2003; Torres et al. 2016a). Based on similarity with R. clavata (Serra-Pereira et al. 2011; Ball et al. 2016; Torres et al. 2016b), R. maderensis is presumed to mostly feed on teleosts. Despite their ecological importance and the worsening of their conservation status, there is a lack of studies on the trophic ecology of these species and its variability in relation with sex and age.

In this study, the trophic ecology of three sympatric batoids was assessed by combining stomach contents and stable isotope ( $\delta^{13}$ C and  $\delta^{15}$ N) approaches. Our specific objectives were to (1) investigate the diets of *D. pastinaca*, *R. clavata* and *R. maderensis*; (2) identify sex and age-related changes in diet; and (3) assess any inter- and intraspecific dietary and trophic niche overlap between the three species in the Azores. The study provides new information about the ecological role of *D. pastinaca, R. clavata* and *R. maderensis* within the Azores region, offering new data on foraging habitats of these species.

# Materials and methods

Study area and sample collection

This study was conducted between September 2021 and May 2022 in Terceira Island, Central Azores (Fig. 1). The convoluted coastline offers a variety of coastal habitats, including rocky reefs, boulder fields, cliffs, enclosed sandy bays, which are subject to various hydrodynamic conditions (Gomes and Pinto 2004). Specimens of D. pastinaca were captured at three sites along the coast of Terceira Island that are from west to east: Cinco Ribeiras (CR), Vila Maria Bay (VM), and Fanal Bay (FB) (Fig. 1). All individuals were caught using spearfishing techniques by a shot in the skull to limit animal suffering (Diggles et al. 2011). This method prevents regurgitation of stomach contents (Bowen 1996) and makes the selection of individuals easier (Frisch et al. 2012; Coll-Calvo et al. 2020). As commercial fisheries do not catch this species, this was the only possible way to obtain specimens. Two professional spearfishers conducted one to two dives a month during the study period by daylight (no night dives were undertaken) in water depths ranging from 0.5 to 20 m. Individuals of R. clavata and R. maderensis were obtained from bottom longlines targeting demersal species. This multi-species fishery uses mainly small vessels (<14 m) operating on the rocky slopes of the island and on seamounts down to 600 m (Da Silva and Pinho 2007; Carvalho et al. 2011). Line setting started one hour before sunrise and line retrieval started about 1.5 h after sunrise (Da Silva and Pinho 2007; Carvalho et al. 2011). After recording total length (T<sub>1</sub>, to the nearest cm), sex and maturity were determined by macroscopic examination following Stehmann (2002). Life stages were classified as immature (immature, developing) or mature (spawning capable). All specimens were collected and processed in accordance with the University of Azores Ethics Committee approval (reference n°9/2021).



Fig. 1 (a) Location of the Azorean Archipelago in the Northeast Atlantic and (b) Terceira Island in the Azores. (c) Map of Terceira Island with the distribution of the sampling sites

Stomach content analysis (SCA)

The stomach was dissected from each individual and the contents were extracted and stored in individual sealable plastic bags. All stomach content samples were kept in a cool box following collection and thereafter frozen at -20 °C in the laboratory until further analysis.

Stomach contents were emptied, rinsed with water, and prey were sorted and identified to the lowest taxonomic level possible under a dissecting microscope with the help of taxonomic guides (Cailliet et al. 1996; Hayward and Ryland 1990; Debelius 1997; Saldanha 2003). Prey items were individually counted and weighed (wet weight). As most food remains were highly digested, the identification of prey was limited, and as such all prey items were pooled into four prey categories: (1) Polychaeta, (2) Crustacea, (3) Mollusca, (4) Teleost. A vacuity index (%V) was calculated as the percentage of empty stomachs (Hyslop 1980). Cumulative prey curves were used to determine whether the number of stomachs with identifiable preys were adequate to accurately describe the diet of the species. The final four curve points

( $\blacksquare$ =samples collected during commercial fisheries;  $\blacksquare$ =samples collected during spearfishing). *CR*, Cinco Ribeiras; *VM*, Vila Maria Bay; *FB*, Fanal Bay

for each species, and for each subgroup (female, male, immature, mature) were statistically compared with the zero slope line using a Student's t-test (Bizzarro et al. 2007).

The importance of various prey species to the diet of *D. pastinaca*, *R. clavata*, and *R. maderensis* was assessed by calculating the percent frequency of occurrence (%FO), average percent number (%N), average percent weight (%W), prey-specific number (%PN), and prey-specific weight (%PW) (Hyslop 1980; Brown et al. 2012). The prey-specific index of relative importance (%PSIRI) was used to measure the relative importance of each prey group in the diet of *D. pastinaca*, *R. clavata*, and *R. maderensis* (Brown et al. 2012). These indices were calculated as follows:

Frequency of occurrence (FO):

$$\% \text{FO}_i = \frac{n_i}{n} \times 100,$$

Average percent abundance  $(\%N_i,\%W_i)$ :

$$\%A_i = \frac{\sum_{j=1}^n \%A_{ij}}{n}$$

Prey-specific abundance ( $%PN_i, %PW_i$ ):

$$\%PA_i = \frac{\sum_{j=1}^n \%A_{ij}}{n_i}$$

Prey-specific index of relative importance (%PSIRI<sub>*i*</sub>):

$$\% PSIRI_i = \frac{\% FO \times_i (\% PN_i + \% PW_i)}{2}$$

where  $\% A_{ij}$  is the per cent abundance (by number or weight) of prey category *i* in stomach sample *j*,  $n_i$  is the number of stomachs containing prey *i*, and *n* is the total number of stomachs containing prey (Brown et al. 2012).

To measure the trophic niche breadth, the standardized Levin's index (Ba) (Krebs 1999) was calculated:

$$Ba = \frac{\frac{1}{\sum P_{ij}^2} - 1}{n - 1}$$

where Pij is the proportion of the prey j in the diet of the species i, and n is the number of prey categories. This index ranges from 0 (minimum niche breadth and maximum specialization) to 1 (maximum niche breadth and minimum specialization).

The trophic position  $(TP_j)$  was defined for species and subgroups according to Cortés (1999):

$$TP_j = 1 + \left(\sum_{j=1}^n DC_{ji} \times TP_i\right)$$

where  $TP_j$  is the trophic level of the species (j),  $DC_{ji}$  is the proportion of the prey category (i) in predator diet (j); n is the total number of prey categories, and  $TP_i$  is the trophic level of the prey group (i). Trophic levels of prey categories were assigned based on the literature (Supplemental Table 1).

Mann–Whitney–Wilcoxon rank tests for non-normally distributed data were applied using the R function "pairwise.wilcox.test" (R Core Team 2022) to test for pairwise differences in TP between species. A second series of Mann–Whitney–Wilcoxon rank tests were used to test for sex and size-effect among species. Non-metric multidimensional scaling (NMDS) ordination was used to explore variation in the diet composition among species and between subgroups (males, females, immatures, matures). Bray–Curtis dissimilarity matrices were derived based on absolute volume (weight) data of the four prey categories across all species and subgroups using the R function "vegdist" (Gardener 2014; R Core Team 2022). Given the low number of stomachs collected (see Table 1), gravimetric data pooling was designed to reduce the number of prey categories in the samples with zero values, thus increasing the efficiency of multivariate analysis (White et al. 2004). Prior to conduct permutational multivariate analysis of variance (PER-MANOVA), a non-parametric permutational analysis of multivariate dispersions (PERMDISP) was run using the R function "permdisp" of the package vegan, to test the homogeneity of multivariate dispersions of data (Anderson 2006). PERMANOVA was used as implemented in the "adonis" function of the vegan R package (Oksanen et al. 2013) to test for possible statistical differences among species and subgroups' diets. For pairwise differences between species diet compositions, PERMANOVA tests were run for all species pairs using the "adonis.pair" function of the EcolUtils R package (Salazar 2018; R Core Team 2022), and used Bonferroni correction to set significance levels for p values. In the case of significant differences in diet composition were observed, similarity percentage (SIMPER) analysis was conducted using the R function "simper" of the package vegan (Oksanen et al. 2013; R Core Team 2022) to detect which prey categories were the main contributors to the dissimilarity observed.

#### Stable isotope analysis (SIA)

Stable isotope analysis was performed on a subset of individuals for each species (see Table 1). Approximately 5 g of white muscle tissue was collected from the dorsal side of rays. Each muscle tissue was rinsed, frozen, dried at 50 °C for 48 h, and then ground into a homogeneous powder using a mortar and pestle. Ground samples were weighed (0.5-1 mg) into tin capsules.  $\delta^{13}$ C and  $\delta^{15}$ C were determined using a continuous flow mass spectrometer coupled to an elemental analyzer, at the Stable Isotope Laboratory of the University of Caen (icore.unicaen.fr). The isotopic composition is reported in the conventional delta ( $\delta$ ) per mil notation (%), relative to Vienna Pee Dee Belemnite ( $\delta^{13}$ C) and atmospheric N<sub>2</sub> ( $\delta^{13}$ C). Lipids were not extracted from samples. Thus, correction equations were applied for  $\delta^{13}C$  ( $\delta^{13}C_{LE} = 0.967 \times \delta^1$  ${}^{3}C_{bulk} + 0.861$ ) and  $\delta^{15}C$  ( $\delta^{15}N_{LE} = 1.018 \times \delta^{15}N_{bulk} + 0.020$ ), where  $\delta^{13}C_{LE}$  and  $\delta^{15}N_{LE}$  are corrected with

<b>Table 1</b> Size range ( $T_L$ , cm), number (% of empty) of stomach
content (SCA) and stable isotope samples (SIA), standardized
Levin's index (Ba), estimated trophic position based on the
diet (TP), Bayesian isotopic ellipse area corrected (SEAc), and
estimated trophic position by $\delta^{15}N$ (TPSI) for <i>D. pastinaca</i> , <i>R</i> .

*clavata*, *R. maderensis*, and each of the subgroups. SE=standard error. Different letters indicate significant difference in TP and TPSI values between species (capital letters) and between sexes or maturity stages (lowercase) within each species

Species	Size range (cm)	SCA	SIA	$\delta^{13}C \pm SE$	$\delta^{15}N\pm SE$	Levin's index (Ba)	TP±SE	SEAc	TPSI±SE
D. pastinaca	32–59	15 (13%)	12	$-15.2 \pm 1.9^{A}$	12.1 ± 1.9 <sup>A</sup>	0.51	3.48±0.31 <sup>A</sup>	0.68	$3.29 \pm 0.53$ <sup>A</sup>
Males	36–51	6	7	$-15.3 \pm 2.2^{a}$	$10.9 \pm 1.0^{a}$	0.44	$3.43 \pm 0.26$ <sup>a</sup>	0.71	$3.31 \pm 0.47$ <sup>a</sup>
Females	32–59	7	5	$-14.9\pm1.5^{a}$	$11.0 \pm 1.4^{a}$	0.54	$3.46 \pm 0.32^{a}$	0.66	$3.25 \pm 0.67$ <sup>a</sup>
Matures	45–59	7	6	$-15.8\pm1.8^a$	$11.5 \pm 0.9^{a}$	0.58	$3.52 \pm 0.38$ <sup>a</sup>	0.59	$3.52 \pm 0.53$ <sup>a</sup>
Immatures	32–43	6	6	$-14.5 \pm 1.9^{a}$	$10.4 \pm 1.2^{a}$	0.46	$3.41 \pm 0.36^{a}$	0.65	$3.27 \pm 0.49^{a}$
R. clavata	35-91	29 (41%)	11	$-17.3\pm1.7^{\rm B}$	$12.1 \pm 0.5$ <sup>B</sup>	0.17	$4.19 \pm 0.39^{\text{ B}}$	0.30	$3.57 \pm 0.32^{\text{ A, B}}$
Males	35-85	7	5	$-18.3 \pm 1.1^{a}$	$12.1\pm0.5$ <sup>a</sup>	0.21	$4.13 \pm 0.42^{a}$	0.27	$3.53 \pm 0.51$ <sup>a</sup>
Females	39–91	10	6	$-16.4 \pm 1.7^{a}$	$12.2 \pm 0.6^{a}$	0.16	$4.21 \pm 0.44$ <sup>a</sup>	0.35	$3.58 \pm 0.44$ <sup>a</sup>
Matures	76–91	6	5	$-18.5 \pm 0.9$ <sup>a</sup>	$12.3 \pm 0.4^{a}$	0.12	$4.35 \pm 0.17$ <sup>a</sup>	0.23	$3.60 \pm 0.58$ <sup>a</sup>
Immatures	35-69	11	6	$-16.3 \pm 1.6$ <sup>b</sup>	$11.9 \pm 0.6^{a}$	0.23	$3.87 \pm 0.21$ <sup>b</sup>	0.34	$3.34 \pm 0.45$ <sup>a</sup>
R. maderensis	31-87	23 (39%)	12	$-18.3 \pm 1.4$ <sup>B</sup>	$12.7 \pm 1.1^{B}$	0.23	$4.14 \pm 0.33$ <sup>B</sup>	0.52	$3.76 \pm 0.42$ <sup>B</sup>
Males	31–79	5	7	$-17.9 \pm 1.6$ <sup>a</sup>	$13.2 \pm 1.0^{a}$	0.25	$4.12 \pm 0.38^{a}$	0.59	$3.69 \pm 0.52$ <sup>a</sup>
Females	40-87	9	5	$-18.6 \pm 1.5$ <sup>b</sup>	$12.6 \pm 1.1^{a}$	0.22	$4.23 \pm 0.41$ <sup>a</sup>	0.50	$3.73 \pm 0.57$ <sup>a</sup>
Matures	64–87	7	5	$-19.5 \pm 0.3$ <sup>a</sup>	$13.6 \pm 0.7$ <sup>a</sup>	0.16	$4.31 \pm 0.19^{a}$	0.43	$3.88 \pm 0.65$ <sup>a</sup>
Immatures	31–58	7	6	$-17.3 \pm 1.4$ <sup>b</sup>	$12.2 \pm 1.0^{b}$	0.27	$3.85 \pm 0.17$ <sup>b</sup>	0.58	$3.62 \pm 0.49$ <sup>b</sup>

lipid extraction (LE) (Logan et al. 2008). In addition, there was not prior urea extraction from the samples. To ensure a confident interpretation, the mathematical corrections developed for elasmobranchs by (Li et al. 2016) were applied to correct the  $\delta^{15}N$  ( $\delta^{15}N_{LE + UE} = 0.856 \times \delta^{15}N_{LE} + 2.813$ ), where  $\delta^{15}N_{LE+UE}$  is the value of  $\delta^{15}N$  corrected with LE and urea extraction (UE).

The normality and homoscedasticity of isotope data were tested using the Shapiro–Wilk and Levene's test, respectively. The nonparametric Wilcoxon signed-rank test was used to determine statistical differences in  $\delta^{13}C$  and  $\delta^{15}N$  among batoid species and subgroups (sexes and maturity stages).

To evaluate the isotopic niche for each species and for each sex-maturation stage within each species, the smallest convex hulls that contain all individual  $\delta^{13}$ C and  $\delta^{15}$ N values within a group were calculated to represent the total niche breadth of each species (Layman et al. 2007). As a measure of the trophic breadth, the Bayesian isotopic ellipse area corrected (SEAc) was calculated to reduce bias for small sample size (Jackson et al. 2011). The niche area is defined as the area occupied in bi-plot space in  $\%^2$  and is considered to be a proxy for the extent of trophic diversity (or trophic breadth) exploited by species or each sexmaturation stage within each species (high SEAc values indicate high trophic breadth). Due to the absence of isotopic signals of prey, the use of isotopic ellipses as a proxy of trophic diversity should be taken with caution because stable isotope analysis can have difficulty separating groups when the food web base has similar  $\delta^{13}$ C signatures (Shipley and Matich 2020). Inversely, when the food web base has different  $\delta^{13}C$ signatures, stable isotope analysis can indicate high trophic diversity while predators feed on few prey items (Shipley and Matich 2020). Bayesian standard ellipses corrected were also used to calculate isotopic niche overlap among groups and were measured as a percentage (%) of the shared isotopic space of each species (Jackson et al. 2011). Differences in SEAc size were considered significant if the 95% credibility intervals of posterior draws did not overlap. These analyses were done using the "SIBER" library in R statistical package v. R 4.0.0 (Jackson et al. 2011).

To estimate the trophic position for species and subgroups using nitrogen stable isotopes, a scaled  $\Delta^{15}N$  framework approach based on a dietary  $\delta^{15}N$  value-dependent model was used (Hussey et al.

2014). With knowledge of the  $\delta^{15}$ N value of a known baseline consumer ( $\delta^{15}$ N<sub>base</sub>), the  $\delta^{15}$ N value of the consumer ( $\delta^{15}$ N<sub>TP</sub>), the dietary  $\delta^{15}$ N value at which  $\delta^{15}$ N incorporation and  $\delta^{15}$ N elimination are equal ( $\delta^{15}$ N<sub>lim</sub>), and the rate at which the ratio between  $\delta^{15}$ N incorporation and  $\delta^{15}$ N elimination changes relatives to dietary  $\delta^{15}$ N averaged across the food-web (*k*), TPSI, is calculated as:

$$\text{TPSI} = \frac{\log \left(\delta^{15} N_{lim} - \delta^{15} N_{\text{base}}\right) - \log \left(\delta^{15} N_{lim} - \delta^{15} N_{\text{TP}}\right)}{k} + \text{TP}_{\text{base}}$$

Given its abundance throughout the Central and North East Atlantic and since it has been reported to be the thornback ray's main prey (Morato et al. 2003), the boarfish *Capros aper* was used as the  $\delta^{15}N_{hase}$ estimate.  $\delta^{15}N_{base}$  and  $TP_{base}$  were used according to data obtained from Torres et al. (2016a) for this species in the Azores. Values for  $\delta^{15}N_{lim}$  and k are estimated from meta-analysis (see Hussey et al. 2014). Mann-Whitney-Wilcoxon rank tests were applied to test for pairwise differences in TPSI between species. Differences in TPSI by species were assessed using Mann-Whitney-Wilcoxon rank (all sex-maturation stage from one species pooled together compared to all sex maturation stages of the other species). Differences in TPSI among species by subgroups were also analyzed using Mann-Whitney-Wilcoxon rank.

# Results

#### Stomach content analysis

A total of 67 stomachs were analyzed, belonging to 15 D. pastinaca individuals measuring between 32 and 59 cm (T<sub>I</sub>), 29 R. clavata and 23 R. maderensis individuals measuring between 35-91 cm and 31-87 cm, respectively (Table 1). Cumulative curves of prey suggested that enough stomachs were examined to accurately describe the diet of the three studied species and of each subgroup as the cumulative prey curves always reached the asymptote (Supplemental Table 2, Supplemental Fig. 1). Of the analyzed stomachs, 2 D. pastinaca (13%V), 12 R. clavata (41%V), and 9 R. maderensis (39%V) stomachs were empty (Table 1). Overall, 634 individual prey items were found, belonging to four prey categories (Polychaeta, Crustacea, Mollusca, Teleost), from which five were identified to the species level (Supplemental Table 3). These four prey categories were found in all studied species in different proportions (Fig. 2).

Stomach content analysis (SCA) revealed that D. pastinaca, R. clavata, and R. maderensis feed on a wide range of prey items, with crustaceans and teleosts as main categories of prey (Fig. 2, Supplemental Table 3). Crustaceans dominated the diet of D. pastinaca in frequency of occurrence, abundance and weight, while teleosts dominated the diet of R. clavata and R. maderensis (Fig. 2, Supplemental Table 3). Decapods were the most abundant and diversified of all prey species, and they were consumed by the three studied species. Even if teleosts were the main prey categories in R. clavata and R. maderensis, at lower taxonomic levels, differences in %FO, %PN, %PW, and %PSIRI were clear. Raja clavata preyed preferentially upon Macroramphosus scolopax (Linnaeus 1758) and R. maderensis on Pagellus bogaraveo (Brünnich 1768) but unidentified fish were also important for the two species (Fig. 2, Supplemental Table 3). Capros aper (Linnaeus 1758) was also an important fish prey of these two batoids.

Dietary breadths were similar in R. clavata (Ba=0.17) and R. maderensis (Ba=0.23) and smaller than in *D. pastinaca* (Ba=0.51) (Table 1). According to the niche breadth size, immature individuals in R. clavata and R. maderensis were found to be generalist feeders, whereas adults exhibited a specialist feeding strategy (Table 1). The trophic position of D. pastinaca, estimated with the diet composition, was significantly lower (Table 1) than those of R. clavata and R. maderensis, which were not significantly different from each other (Table 1). There were no significant differences in TP between sexes within species. In all species, the trophic position increased with the maturity of the individuals, but this increase was significant for R. clavata and R. maderensis only (Table 1).

The NMDS revealed significant differences between the diet composition of *D. pastinaca* and both *R. clavata* and *R. maderensis*, but not between *R. clavata* and *R. maderensis* (Fig. 3a, Supplemental Table 4). Diet composition did not vary significantly between males and females in any of the studied species (Fig. 3b, d, Supplemental Table 4) and between maturity stages of *D. pastinaca* (Fig. 3e, Supplemental Table 4). PERMANOVA revealed significant differences between maturity stages in *R. clavata* and *R. maderensis* (Fig. 3f, g, Supplemental Table 4). The



Fig. 2 Diet composition of three batoids from the coast of Terceira Island by sex and stage of maturity. Bar charts show the prey-specific index of relative importance (%PSIRI) of the four major prey categories (Crustacea, Mollusca, Polychaeta,

overall diet dissimilarity between D. pastinaca and R. clavata was 77.42%, with teleosts and crustaceans contributing the most to this dissimilarity (38.95% and 32.36% respectively, Supplemental Table 5). The second greatest dissimilarity occurred between the diets of D. pastinaca and R. maderensis (68.11%, Supplemental Table 5), with teleosts and crustaceans as the main drivers of this dissimilarity (33.71%% and 29.19%, respectively, Supplemental Table 5). In R. clavata, the overall diet dissimilarity between immature and mature individuals was 46.36%, and this was significantly made up of crustaceans (23.03%), teleosts (21.56%), and mollusks (1.5%) (Supplemental Table 5). There was 51.45% dissimilarity between immature and mature individuals in R. maderenesis diets, with teleosts and crustaceans being the prey categories that contributed the most to the dissimilarity (26.10% and 23.74%, respectively, Supplemental Table 5). Crustaceans dominated the diet of immature

and Teleost) found in (A) *D. pastinaca*, (B) *R. clavata*, (C) *R. maderensis*. Numbers in parentheses above bars indicate stomach sample size for each combination

*R. clavata* and *R. maderensis*, while teleosts were the most important prey in mature individuals.

Stable isotope analysis (SIA)

 $δ^{13}$ C and  $δ^{15}$ N values were measured in 35 white muscle tissue samples, 12 for *D. pastinaca*, 11 for *R. clavata*, and 12 *R. maderensis* (Table 1). Results for stable isotope analysis for each species and subgroups are represented in Table 1. *D. pastinaca* exhibited  $δ^{13}$ C values significantly enriched compared to those of *R. clavata* and *R. maderensis* (Table 1). *Raja clavata* and *R. maderensis* muscles were slightly  $δ^{15}$ Nenriched compared to *D. pastinaca* muscle (Table 1). In all studied species, stable isotope values did not vary significantly between males and females; however, Mann–Whitney–Wilcoxon rank tests revealed significant differences between stages of maturity in *R. clavata* and *R. maderensis*. Immature and mature *R. clavata* exhibited a significant difference in  $δ^{13}$ C values (Table 1). In *R. maderensis*, mature individuals had lower  $\delta^{13}$ C and higher  $\delta^{15}$ N than immature individuals (Table 1), as shown by their relative positions in isotopic niche space (Fig. 4g).

The niche breadths of the studied species were calculated from Bayesian isotope analysis as 0.68, 0.30, and 0.52 in D. pastinaca, R. clavata, and R. maderensis, respectively (Fig. 4a, Table 1). Among three pair-wise combinations between the studied species, isotopic overlap was only found between R. clavata and R. maderensis (Fig. 4a), with low SEAc overlap value (24%). No SEAc overlap between D. pastinaca and R. clavata or R. maderensis was found (Fig. 4a). There were no significant differences for D. pastinaca, R. clavata, and R. maderensis in isotopic niche breadths between sexes, with high SEAc overlap value (Fig. 4b, c, d, Table 1). The intraspecific analysis by maturity stages for D. pastinaca showed that SEAc values were close, with 29% of overlapping (Fig. 4e). Size-related SEAc differed in R. clavata and R. maderensis, with the immature stage having the higher SEAc values (Fig. 4f, g, Table 1), and the overlap between maturity stages was low in these two species (Fig. 4f, g).

Dasyatis pastinaca exhibited the lowest average trophic position based on  $\delta^{15}$ N values, with a value of  $3.29 \pm 0.53$ . Raja clavata and R. maderensis exhibited averaged values of  $3.57 \pm 0.32$  and  $3.76 \pm 0.42$ , respectively. Raja maderensis has significantly higher TPSI values than D. pastinaca (Table 1). Raja clavata TPSI did not significantly differ from both D. pastinaca and R. maderensis (Table 1). Neither significant interactions between sex and TPSI nor significant differences between sexes were detected in D. pastinaca and R. clavata or R. maderensis (Table 1). Dasyatis pastinaca and R. clavata had similar trophic positioning between immature and mature (Table 1). Mature R. maderensis showed higher trophic positions than immature (Table 1).

# Discussion

In the present study, we provide new information on the trophic ecology of three co-existing batoids in Terceira Island, in the Central Azores. Even if a limited number of studies were carried out in this area, they never attempt to understand inter- and intraspecific variability in the diet of *D. pastinaca*, *R clavata*, and R. maderensis. Both stomach content and stable isotope analysis showed differences in the trophic habits (diet composition, trophic niche, and trophic level) among the three studied species. In addition, R. clavata and R. maderensis showed a change in diet composition and foraging habitat with ontogeny. Although the results of the cumulative prey curves suggest a good sampling effort to describe the diet of the studied species, the low number of samples, especially in D. pastinaca, limits strong conclusions. The degree of prey identification and the absence of reconstituted biomass of prey are other weaknesses of the study. The information provided in this study remains useful to complement those dealing with trophic ecology of these species in the Azores and other locations for these three vulnerable species. Especially, this study provides new information on the trophic ecology of R. maderensis, a poorly studied endemic species of the Macaronesian region.

# Intraspecific assessment

There was some evidence for ontogenetic changes in the diets of R. clavata and R. maderensis although the results must be treated with caution because sample sizes may have not been sufficient to completely characterize the diet of juveniles. Both immature and mature individuals fed on fish, but this prey category was more important in mature individuals, whereas crustaceans were more abundant in the stomach contents of smaller individuals. The ontogenetic shift of R. clavata revealed in this study is consistent with previous studies, which reported that immatures preyed more on crustaceans than adults and individuals shift their diet toward teleosts as size increased (Santić et al. 2012; Kadri et al. 2014; Daban et al. 2022; Gül and Demirel 2022). SIA also showed significant variations of carbon and nitrogen values with age in R. clavata and R. maderensis. Immatures of R. clavata and R. maderensis showed broader isotopic niches than adults. This suggests that these immatures may display movements across an isotopically heterogeneous environment and/or have more diverse food sources, which is in agreement with the diet results. Raja maderensis exhibited significant increases of  $\delta^{15}N$  values with age, indicating that as individuals grow, the contribution of prey



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◄Fig. 3 Non-parametric multidimensional scaling (nMDS) plot for (a) *D. pastinaca*, *R. clavata*, and *R. maderensis*; (b–d) for males (●) and females (▲) *D. pastinaca*, *R. clavata*, and *R. maderensis*; (e–g) for matures (+) and immatures (■) of *D. pastinaca*, *R. clavata*, and *R. maderensis*. Gravimetric data (% prey weight) of the four major prey categories (Polychaeta, Crustacea, Mollusca, and Teleost) were used to examine differences in diet composition

of higher trophic levels increases (Hussey et al. 2011). Although the sampling size is limited, the low dietary overlap between stages of maturity of R. clavata and R. maderensis may also indicate that these species change their diet as they grow, or experience ontogenetic habitat shifts. Ontogenetic changes have been reported for different species of batoids (Marshall et al. 2008; Treloar et al. 2009; Mulas et al. 2019). Several reasons explain this, including increased energy requirements as individuals mature (Lucifora et al. 2009), larger gap size and stronger bite as animals increase in size (Kolmann et al. 2015), and greater movements across foraging habitats as individuals get larger (Wetherbee et al. 2004). Morato et al. (2003) found in the Azores that specimens of R. clavata from different depths fed on different prey, with increasing consumption of penaeids, seabreams, and myctophids with depth. Interestingly, both mature groups of R. clavata and R. maderensis showed depleted  $\delta^{13}$ C values in comparison to immature groups. Because  $\delta^{13}$ C values decrease with distance from shore or depth (Hussey et al. 2011; Shipley et al. 2017), this finding could suggest that in R. clavata and R. maderensis, juveniles and adults segregate by depth or distance to the shore (Rigby et al. 2015). It may also reveal changes in habitat use or hunting strategies with age (Afonso and Hazin 2015). Whether this reflects foraging on different prey taxa, habitat segregation, or both remains to be elucidated. In contrast to the other two species, there was no significant change in diet composition and stable isotopic values with size for D. Pastinaca. Ismen (2003) observed a shift to bony fish consumption among adults in Iskenderun Bay in the eastern Mediterranean. Therefore, the lack of a relationship between age and diet composition or stable isotopic values could either be related to the sample size or be related to the prey availability in Terceira Island compared to Iskenderun Bay.

While sample sizes were limited, no evidence for sex-related dietary differences in the studied species was detected, in agreement with the results from several studies of batoids (Jacobsen and Bennett 2013; Mulas et al. 2019). Other studies have, however, noticed that some batoid species exhibit differential diet and habitat use patterns between sexes (Ajemian and Powers 2014; White et al. 2014; Jargowsky et al. 2020; Serrano-Flores et al. 2021).

#### Interspecific assessment

Overall, the results showed that the three species prey mostly upon crustaceans and fishes, but displayed different feeding habits. Polychaetes and mollusks were of minor importance. Crustaceans, especially decapods, were major prey for D. pastinaca in accordance with previous studies carried out in the Azores (Ponte et al. 2016), the Mediterranean Sea (Ismen 2003; Yeldan et al. 2009; Mulas et al. 2019; Tiralongo et al. 2020) and the Black Sea (Saglam et al. 2010). The diet of *R. clavata* was primarily composed of teleosts, confirming previous observations from the Azores (Morato et al. 2003; Torres et al. 2016a), the Black Sea (Demirhan et al. 2005), the Aegean Sea (Eronat and Ozaydın 2015), and the Mediterranean Sea (Kadri et al. 2014), but our results indicate that some immature individuals feed on crustaceans. Several authors reported a dominance of crustaceans in the diet of R. clavata in the Mediterranean Basin (Kabasakal 2002; Saglam and Bascinar 2008; Santić et al. 2012; Mulas et al. 2019; Daban et al. 2022). Although our results indicated that R. clavata has a specialist feeding strategy, previous studies have described the species as a generalist and opportunistic predator (Kadri et al. 2014; Mulas et al. 2019). The foraging plasticity of some batoids could allow them to exploit the most abundant prey, resulting in the unequal prey contribution (Serrano-Flores et al. 2021). In the Azores, the high degree of piscivory may result from the high availability of fish prey (Morato et al. 2003; Torres et al. 2016a). The diet composition of R. maderensis could not be compared with other studies, since none exist to our knowledge. Results agree with the proportion of crustaceans and teleosts consumed by several skates (Dolgov 2005; Ebert and Bizzarro 2007; Treloar et al. 2009), including the sympatric R. clavata, which has many ecological



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Fig. 4 Isotopic niche overlap plot of: (a) *D. pastinaca, R. clavata,* and *R. maderensis;* males and females of (b) *D. pastinaca,* (c) *R. clavata,* (d) *R. maderensis;* immature and mature individuals of (e) *D. pastinaca,* (f) *R. clavata,* (g) *R. maderensis.* Convex hulls are indicated by black dashed lines. Standard Bayesian ellipses corrected (SEAc) are indicated by solid-colored lines

and phylogenetic affinities (Dolgov 2005; Ebert and Bizzarro 2007; Treloar et al. 2009; Serrano-Flores et al. 2021; Ball et al. 2016; Torres et al. 2016b).

Although based on a small sample size, interspecific differences in isotopic values were highlighted. As isotopic carbon ratios typically increase from offshore to near-shore ecosystems (Hussey et al. 2011), the significantly higher  $\delta^{13}$ C values in *D. pastinaca* supported that the species is predominantly a coastal feeder. Furthermore, the species occupied a limited spatial extent, in agreement with its neritic foraging habitat (Torres et al. 2016b; Das et al. 2022). The significantly lower  $\delta^{13}$ C values in *R*. *clavata* and *R*. maderensis suggested that these species derive their carbon from both benthopelagic and oceanic prey, as the organic matter resulting from pelagic sources is more depleted in  $\delta^{13}$ C when compared to the benthos (Hussey et al. 2011). Dasyatis pastinaca preyed mainly upon crustacean and other low trophic level preys, explaining the significantly lower  $\delta^{15}$ N compared to *R*. *clavata* and *R*. maderensis that rely mostly on fish. While D. pastinaca and R. clavata were described as generalist and specialist respectively, R. maderensis showed an intermediate feeding strategy based on SEAc. Important isotopic niche overlap was observed between R. clavata and R. maderensis in accordance with the diet results. This high overlap does not necessarily mean that species are feeding on the same prey (Hussey et al. 2011) because different prey species can have similar isotopic values. It has also been suggested that the metabolic turnover of white muscle may be too low (> 250 days) to reveal fine-scale insights, resulting in high overlap between potential interactive species (MacNeil et al. 2005). Therefore, future work should incorporate isotopic signal of prey and faster isotopic turnover rate tissues (e.g., plasma and whole blood) to provide even greater insight into the diets, foraging habitats, and home ranges of the studied species.

Role of D. pastinaca, R. clavata, and R. maderensis in food webs

Our trophic positions estimated from stomach contents were generally close to the standardized values reported for skates (Ebert and Bizzarro 2007) and stingrays (Jacobsen and Bennett 2013), and consistent with values reported using stable isotopebased trophic positions (Torres et al. 2016a; Gül and Demirel 2020, 2022). Our estimation of trophic level based on stable isotopes was lower than based on stomach contents. Stable isotope and stomach content analyses can produce different results for trophic position due to the different temporal integration of the diet, and in our case, to the limited information available on baselines to estimate trophic position (Olin et al. 2013). Trophic position estimates ranged from  $3.25 \pm 0.67$  to  $4.35 \pm 0.17$  according to the method, the species, and the individuals. These estimates are coherent with a previous study conducted in the Azores (Torres et al. 2016a), confirming the mesopredator role of D. pastinaca and high trophic level of R. clavata and R. maderensis.

The combination of the diet and isotope results indicates that R. clavata and R. maderensis occupy similar foraging habitats and exhibit close related dietary habits. This suggests that these species have very similar functions in the food web off Terceira Island. This functional redundancy is typical of high trophic level predators and it is associated with the wide sharing of food resources, and less trophic specialization (Heupel et al. 2014). High trophic level predators impact the prey community through diffuse predation, as several predators forage upon the same prey species (Heupel et al. 2014). Diffuse predation is important in maintaining structure and robustness of food webs (Bascompte et al. 2005; Navia et al. 2017). Raja clavata and R. maderensis exhibited ontogenetic changes in their diet. As they mature, their trophic levels change and, with that, the roles they play as predators change. Consequently, these changes can lead to a more important role of the species in structuring the prey community (Bascompte et al. 2005).

In conclusion, this study presents new information on the feeding ecology of *D. pastinaca*, *R clavata*, and *R. maderensis* in the Azores, where limited information on the ecology of these species is available. The results revealed a clear feeding preference for crustaceans for *D. pastinaca* and teleosts for *R. clavata* and *R. maderensis* off Terceira Island. The results are in overall agreement with previous studies on these species in other areas and with the general knowledge on batoids. They remain preliminary, as the number of samples was low, therefore limiting the robustness of the conclusions. Further studies should be conducted to better understand ontogenetic, sexrelated, and seasonal changes in diet and habitat use. In addition, the variation of diet of these species in the Macaronesian region should be investigated, especially for the endemic *R. maderensis*, which is listed as vulnerable by the IUCN and is regularly by-caught in local fisheries.

Author contributions All authors contributed to the study conception and design. Material preparation, data collection, and analyses were per Paco Fontaine, João Pedro Barreiros, and Sébastien Jaquemet. The first draft of the manuscript was written by Paco Fontaine and all authors commented on previous versions of the manuscript. All authors read and approved the final manuscript.

**Data availability** The datasets generated during and/or analyzed during the current study are available from the corresponding author on reasonable request.

#### Declarations

**Conflicts of interest** All authors certify that they have no affiliations with or involvement in any organization or entity with any financial interest or non-financial interest in the subject matter or materials discussed in this manuscript.

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