



# Variability of prey preferences and uptake of anthropogenic particles by juvenile white seabream in a coastal lagoon nursery ground

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**Abstract** Marine plastic litter, originating from land-based sources, enters the marine environment by passing through coastal ecosystems such as lagoons and estuaries. As early life history stages (ELHS) of many commercially important fish species rely on these transitional areas as nursery grounds, we hypothesized that they encounter a spatial gradient of habitat quality and pollution from inner to outer parts of their vital environment. With sizes < 5 mm, anthropogenic particles (AP), among them microplastic (MP) fibers and fragments, entail a high bioavailability for ELHS of fish, potentially facilitating AP uptake at early developmental stages which may have implications for their survival and growth. This study provides a contextualization baseline between feeding

preferences and uptake of AP by the white seabream *Diplodus sargus* (Linnaeus, 1758) in an estuarine nursery ground on the southern coast of Portugal. Juvenile fish showed a generalized, omnivorous feeding mode with differences in trophic resource utilization between individuals collected at distinct seagrass meadows in the lagoon. A total of 23.13% of the fish ( $n=147$ ) were detected with AP in the gastrointestinal tract, and the mean number of AP per AP-feeding individual was  $1.64 \pm 1.04$ , with anthropogenic fibers ( $n=47$ ) occurring more frequently than fragments ( $n=9$ ). Knowledge of the underlying factors for MP ingestion will be greatly enhanced by considering environmental conditions along with species-stage and life-stage specific feeding modes and prey preferences which shape the uptake probability of anthropogenic fibers and fragments.

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## Introduction

Coastal lagoons and estuaries are recognized as highly productive transitional environments, providing vital habitats and essential ecosystem services (Costanza et al. 1997; Beck et al. 2001; Erzini et al. 2022). Their importance as nursery grounds for many commercially valuable fish species arises from the combination of structural complexity and favorable

environmental conditions (Elliott and Hemingway 2002; Seitz et al. 2014). Irrespective of their acknowledged relevance, coastal ecosystems face severe exposure to habitat degradation (Kennish 2002; Lotze et al. 2006) and pollution as a result of the continuously increasing urbanization of onshore and offshore regions (Browne et al. 2011) along with the riverine input of anthropogenic particles (AP) from land-based sources (Rochman 2018).

Within the past decade, AP, such as microplastic (MP) fragments and fibers of < 5 mm in size, became the center of scientific and public interest. Arising from the cumulative industrial application of plastic materials and the lack of efficient waste-management (Jambeck et al. 2015; Ryan 2015), increasing quantities of MP have been documented in coastal areas around the world (Barnes et al. 2009; Cole et al. 2011; Kumar et al. 2021). Due to their size range, MP particles are available for ingestion (and potential trophic transfer) for a variety of organisms at the base of the marine food web, among them early life history stages (ELHS) of fish (Cole et al. 2013; Gove et al. 2019). Growth, condition, and survival of ELHS are strongly shaped by gradients in abiotic and biotic conditions in vital nursery grounds, with direct consequences for recruitment success to adult fish stocks (Boehlert and Mundy 1988; Beverton and Iles 1992; Ciotti et al. 2014). Therefore, in-depth research on the potential uptake and consequent physiological impact of MP is needed to holistically assess the underlying factors contributing to recruitment variability in commercially important fish species. Although plastic ingestion in fish has been reported across more than 140 families (Azevedo-Santos et al. 2019), most field studies neither assessed ELHS of commercially important fish taxa nor investigated prey selectivity by comparing diet composition with prey availability (Gamito et al. 2003; Selleslagh and Amara 2015; Müller 2021). Feeding strategy and prey preferences undergo ontogenetic changes (Galarowicz et al. 2006; Sánchez-Hernández et al. 2019) and understanding of the feeding ecology of a life stage or species is considered of major importance to establish and improve sustainable management and conservation (Braga et al. 2012).

Our study aims to fill existing knowledge gaps about the potential uptake and effects of AP for growth and survival of ELHS of a commercially important fish species on the basis of a holistic examination and

contextualization of the gastrointestinal tract of the fish. Juvenile white seabream, *Diplodus sargus* (Linnaeus, 1758), were chosen as a model organism for this field study. Due to their opportunistic prey intake and potentially disadvantageous body:AP size ratio, ELHS of fish supposedly show a higher AP ingestion probability and sensitivity towards adverse physiological effects than adults (Critchell and Hoogenboom 2018; Salerno et al. 2021). The omnivorous feeding mode of the white seabream has been hypothesized to be an influencing factor for elevated AP uptake rates (Mizraji et al. 2017; Garcia et al. 2020), yet this species is able to discriminate between natural and artificial prey items (Müller et al. 2020), challenging the abovementioned notions. The selected study area, the Ria Formosa lagoon, located at the southern Portuguese coast, is recognized as an essential nursery for the adjacent coastal fish populations, fostering significant populations of juvenile fish, including commercially relevant members of the seabream family (Sparidae) (Monteiro et al. 1990; Erzini et al. 2002; Ribeiro et al. 2006). Despite its ecological and economic importance, the ecosystem is facing persistent anthropogenic pressures (e.g., habitat degradation, impaired water quality) across a spatio-temporal gradient, with higher detrimental effects being exerted on biological communities in the interior, urbanized parts of the lagoon during summer months (Newton et al. 2003; Cravo et al. 2012; Guimarães et al. 2012).

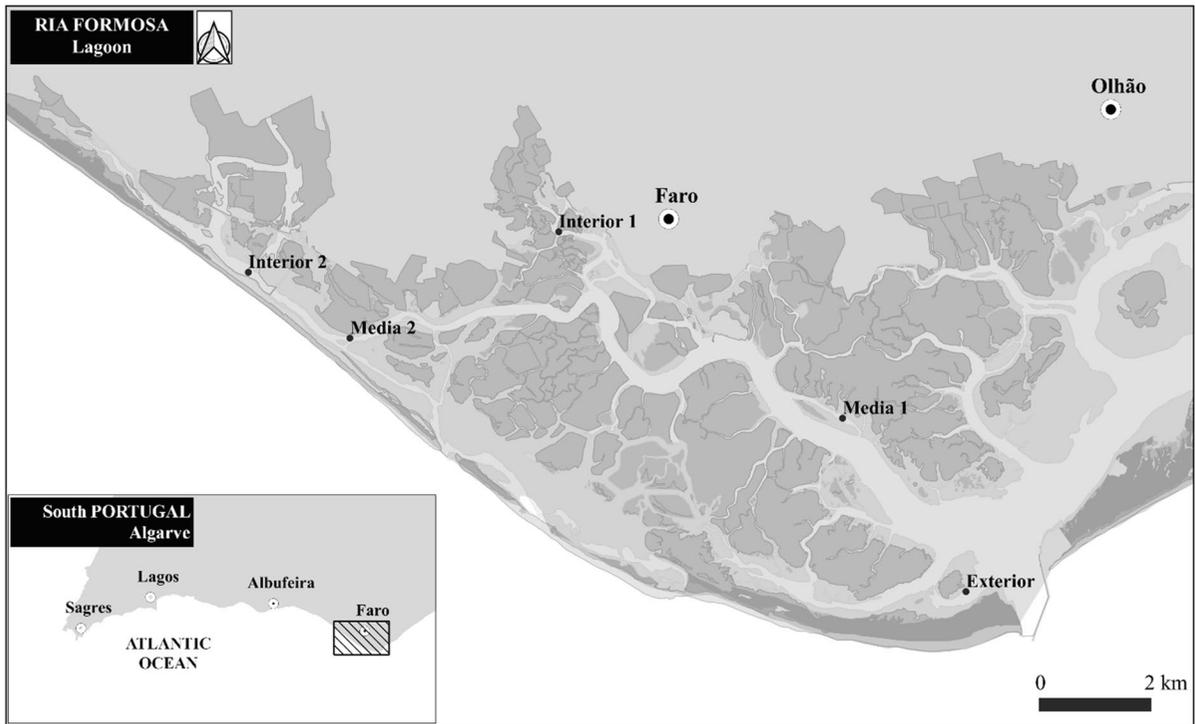
Based on the hypotheses that the gross of AP enters the marine environment through coastal ecosystems and vegetated habitats, which are vital nursery grounds for a wide range of fish ELHS, including the white seabream, act as a trap for AP (Cozzolino et al. 2020; Jones et al. 2020), our research questions were:

- i) Do juvenile *D. sargus* encounter a gradient of habitat quality and AP pollution in their vital coastal lagoon nursery area?
- ii) What are the driving factors for and the potentially detrimental effects of AP ingestion for ELHS of an omnivorous fish species?

## Materials and methods

### Field sampling design

To investigate potential spatial abiotic and biotic gradients as well as differences in AP abundance, five



**Fig. 1** Map of mainland Portugal, highlighting the location of the Ria Formosa lagoon at the Algarve coast. Enlarged map of the western part of the lagoon displays the five distinct

sampling sites selected for this study and two inlets. Sampling across the different stations took place between July and August 2018

distinct seagrass-vegetated sampling sites were selected in the western part of the Ria Formosa lagoon at the southern Portuguese coast (Fig. 1; Table 1): Two sampling sites were chosen for their proximity (linear distance < 1 km) to urbanized areas, i.e., the city center of Faro and the Praia de Faro (thereinafter called “interior 1” and “interior 2”), and two additional sampling sites were selected in intermediate linear distance (< 5 km) to the urbanized areas (“media 1” and “media 2”). One more sampling site was designated close to the inlet of the main channel to the Atlantic Ocean (“exterior”); as

the only other western Ria Formosa inlet yielded no seagrass-vegetated sites to be sampled for comparison, only one station at the greatest distance to the city (linear distance ~7 km) was considered appropriate. The field work design included three sampling campaigns in summer 2018 (04./05. July; 27. July; 10./13. August). All samplings occurred during slack tide (max. 2 h prior and after low tide); at each site, temperature, salinity, and dissolved oxygen were recorded using a YSI Professional Plus Multiparameter Instrument (YSI Pro2030), and sampling of juvenile seabream as well as

**Table 1** Overview of the sampling sites, longitude, and latitude, as well as linear distance (in km) to either the urbanized area of Faro City Center (FCC) or Praia de Faro (PdF) (x = indicates the respective reference point). Sampling occurred between July and August 2018

Station	Longitude	Latitude	Reference point FCC (-7.936640, 37.013665)	Reference point PdF (-7.994420, 37.008505)	Linear distance (km)
Interior 1	-7.947183	37.014883	x		0.9469
Interior 2	-7.999077	37.011292		x	0.5173
Media 1	-7.899833	36.991633	x		4.0893
Media 2	-7.976471	37.003297		x	1.6977
Exterior	-7.877817	36.97205	x		6.9867

of environmental AP abundance in planktic and benthic compartments was conducted.

#### Field sample collection

##### *Seabream*

Juvenile seabream were collected during all three sampling campaigns with a beach seine (25 m width, 3.5 m height, 9 mm stretched mesh size) with a cod end-type sac in the middle to facilitate handling of the catch (Erzini et al. 2002), deployed in a standardized manner (Adão et al. 2022). The net was deployed from a 6.5-m boat and towed 20 m along the shore by the boat from one side and by two to three people on the shore from the other side. After half of the towing distance, the boat headed to the shore at an angle, with the people on shore continuing towing along the shore to meet the boat when it reached the shore. Based on GPS measurements of many hauls, the average area sampled in this manner is 1087 m<sup>2</sup> (Adão et al. 2022). The catch, which was retained in the cod end, was then emptied into a box, and sorted for seabream. Individuals that were gilled in the mesh of the seine net were also collected. Juvenile seabream were stored in labeled plastic bags inside a cooler, equipped with freezer packs, and transported to the laboratory where the fish were preserved in a freezer at CCMAR, Faro, Portugal, until further processing and analyses.

#### Environmental AP abundance

Zooplankton and macrozoobenthos samples were collected during the first and third sampling campaigns to assess the AP abundance in planktic and benthic compartments. AP abundance was assessed in the water column using a conical zooplankton net (200 µm mesh size, 0.13 m<sup>2</sup> mouth opening, equipped with a HydroBios flowmeter to precisely assess the sampled water volume) which was towed 300 m just below the sea surface behind the boat around the sampling site. Additionally, macrozoobenthos samples were taken with a push net (1000 µm mesh size, rectangular opening of 20 cm height and 50 cm width), operated manually, and pushed over 10 m through the seagrass meadow. The zooplankton samples were preserved in glass jars containing a seawater-ethanol solution (70% ethanol) until further analysis. The

macrozoobenthos samples were stored deep-frozen in individual zip-lock bags until further analysis.

#### Laboratory processing

##### *Contamination control*

Potential contamination of the samples was reduced by applying the following procedural measures (Lusher et al. 2017) in the laboratories of CCMAR in Faro, Portugal, and ZMT in Bremen, Germany: Work benches and equipment (i.e., glassware, tweezers, needles, and Bogorov chambers) were cleaned with ethanol before use, the latter being additionally checked under the microscope for contamination. Contamination with fibers was minimized by washing hands and forearms thoroughly as well as by wearing cotton laboratory coats and nitrile gloves during all analytical steps performed. Care was taken also by wearing cotton clothing underneath the laboratory coat and the color of the clothing was recorded to trace back potential contamination. If in doubt about the origin, fibers found in the samples were excluded from the analysis.

To avoid misidentification of AP, the criteria proposed by Hidalgo-Ruz et al. (2012) and Lusher et al. (2017 & 2020) were adopted to the visual sorting of fragments and fibers in the fish gastrointestinal tract (GIT) and zooplankton as well as macrozoobenthos samples—if in doubt about the origin of a fragmented or fibrous object, it was excluded from the analysis, thereby applying a conservative visual classification and quantification protocol. As no subsequent polymer characterization was performed to verify the synthetic origin of the items, the detected fragments and fibers will be referred to as anthropogenic particles.

#### Environmental AP abundance

##### *Zooplankton*

Zooplankton samples were transported to the laboratory facilities of ZMT, Bremen, for further analyses. In the laboratory, samples were split into 1/4–1/128 fractions (depending on the amount of zooplanktic organisms collected) using a modified Motoda plankton sub-sampler. Zooplankton and AP images with a resolution of 2400 dpi were taken from the respective subsamples with a ZooScan (ZooScan Model

V4, Hydroptic Inc., France), following the procedures described by Gorsky et al. (2010). The scans obtained were analyzed using the software ZooProcess on the ImageJ macro language (Gorsky et al. 2010), allowing automated processing and measuring of the scanned images. ZooProcess associates the images with the available metadata and divides the scanned images into multiple single images ideally depicting a single organism or particle only. Images containing multiple or overlapping organisms and AP were manually edited using the software and subsequently processed again. For identification, all images were uploaded to the website EcoTaxa (<http://ecotaxa.obs-vlfr.fr/prj/>) where Random Forest Algorithm in combination with convolutional neural network feature extraction automatically classifies them; afterwards, the preliminary classification of potential AP, planktic organisms, and detritus was manually validated. If in doubt about the artificial origin of a fiber or fragment, the item was visually inspected both in the ZooScan scanning chamber as well as under a stereomicroscope (Gilfillan et al. 2009; Pedrotti et al. 2014). If doubt remained, the item was excluded from the analysis.

### Macrozoobenthos

The entire zip-lock bag containing the benthic sampling material was placed into a sorting tray and defrosted using tap water. Upon defrosting, one or several trays were filled with either the entire sample or several subsamples (depending on the amount of material collected) to facilitate handling. In an initial step, seagrass shoots, algae, and larger prey items (e.g., broken mollusk shells, larger crustaceans) were separated from the rest of the sample and excluded from the analysis. All remaining items, AP as well as prey organisms, were manually sorted, identified, and counted.

### Seabream

#### Fish growth

Fish were thawed at room temperature before examination. Individuals of the species *Diplodus sargus* were identified by visually inspecting distinctive external

characteristics (i.e., pigmentation and dentition), measured to the nearest mm (standard length=SL, total length=TL, height=H) and weighed before and after dissection (WW).

### Analysis of gastrointestinal tract (GIT)

The wet weight of the entire GIT was measured to the nearest mg; afterwards, the GIT was preserved in 70% ethanol and stored in Eppendorf tubes until further analysis. Upon content analysis, the GIT was put in a Bogorov counting chamber (filled with 70% ethanol) and opened with fine scissors and tweezers under a stereomicroscope (Zeiss Stemi 2000-C). The content was visually inspected for both artificial (i.e., AP fibers and fragments) and natural prey items, and the latter was identified to higher taxonomic levels following the approach of taxonomic sufficiency as proposed by Ellis (1985) and Ferraro and Cole (1990). Fibers detected in the GIT were carefully inspected for vegetal morphological features such as organic structures or segmentation to avoid misidentification with artificial fibers. Furthermore, only fibers were counted that were found attached to GIT content remains; free-floating fibers in the Bogorov counting chamber as well as fibers matching the clothing underneath the cotton lab-coat were also excluded. Though this conservative procedure could result in partial underestimation of AP ingestion, the potential bias caused by airborne contamination is reduced to a minimum to ensure reliable results. The GIT analysis time was not standardized for this study due to the variability of GIT content and volume; to enhance comparability of results, a fixed 20× magnification was chosen and maintained throughout all analyses.

Percentage frequency of occurrence (%FO) was chosen to be an appropriate, robust measure to analyze the GIT content of fish (Baker et al. 2014); it was calculated according to the following formula:

$$\%FO = \frac{\text{number of GIT containing prey item } i}{\text{total number of GIT containing prey}} \times 100$$

Moreover, prey-specific abundance (%P) was calculated for all countable faunal prey items:

$$\%P = \frac{\text{abundance of prey item } i}{\text{total number of prey items in all GIT containing prey item } i} \times 100$$

The analyses of the feeding strategy, prey importance, and inter-individual as well as intra-individual components of niche width were based on the two-dimensional plot (Fig. S1, supplementary information) of prey-specific abundance %P (y-axis) and frequency of occurrence %FO (x-axis), following the approach suggested by Amundsen et al. (1996) as a modification of the Costello Method (Costello 1990).

### Statistical analyses

Fish morphometrics were tested for normality of distribution (Shapiro–Wilk test) and homogeneity of variance (Fligner–Killeen test for not-normally distributed data). In case of violation of normality, the Kruskal–Wallis test was used to investigate statistical differences between the groups—if significant differences were detected, Dunn’s test (with Holm correction) was computed to perform a pairwise comparison between the groups to identify which groups differ.

Prey preferences across the different stations were examined with non-metric multidimensional scaling (nMDS) ordinations, using Bray–Curtis similarity coefficient on presence-absence data (R-package: “vegan” by Oksanen et al. 2020). Analysis of similarity (ANOSIM) was computed to examine the significance of prey item groups in the ordination pattern (Clarke 1993). A subsequent indicator species analysis (R-package: “indicspecies” by De Cáceres and Legendre 2009) was performed to identify prey items that were found more frequently in the GIT of fish from one station compared to another. Potential differences in the uptake of AP across sampling sites and campaigns were investigated using pairwise *t*-tests and Benjamini and Hochberg adjustment.

To contextualize AP uptake with other prey items, the data obtained for the MP-feeding individuals were analyzed for potential correlation by computing the Spearman rank correlation coefficient ( $r_s$ ) where values can vary between  $-1$  (indicating a strong negative correlation of the variables compared),  $0$  (indicating no association between the variables compared), and  $+1$  (indicating a strong positive correlation of the variables compared).

Redundancy analysis (RDA) was used to model the response variables (AP fragments and fibers) as a function of nominal (sampling campaign, location) and quantitative explanatory variables using PAST (Hammer et al. 2001). Three dummy variables (0, 1)

were created for each of the two nominal variables (three location types and three sampling periods). The selection of quantitative variables was based on variance inflation factors (VIF), with  $VIF > 50$  used as the criteria to remove variables. The variables retained for the RDA were low tide (m), temperature ( $^{\circ}\text{C}$ ), salinity (PSU), oxygen ( $\text{mg l}^{-1}$ ), wind speed (km/h), standard length (mm), wet weight (g), wet weight of the gastric intestinal tract (g), and the total number of prey items. Level of significance was set to  $P \leq 0.05$ .

Statistical analyses and data visualization (except for RDA) were realized with Microsoft 365 and R (version 4.0.5) (R Core Team 2020).

## Results

### Habitat quality and environmental AP availability

All five sampling sites were vegetated by seagrass and marine algae. However, slight differences were detected in the physico-chemical characteristics of the stations (Table S1, supplementary information): Temperature ranged between  $19.8$  and  $24.2$   $^{\circ}\text{C}$  across the different sampling sites and campaigns, showing a spatio-temporal gradient from inside the lagoon to the Atlantic Ocean inlet over the different campaigns, with a minor peak during the second campaign (end of July 2018). Salinity ranged between  $35.66$  and  $38.11$  PSU, showing comparable spatio-temporal fluctuations to those for temperature. Oxygen was generally higher at all stations during the first campaign as compared to the following two samplings and ranged between  $4.08$  and  $8.56$   $\text{mg l}^{-1}$ , with the highest value recorded at the station closest to the Atlantic Ocean, and the lowest oxygen concentration measured at station “interior 1”.

Following a conservative visual identification using the images produced by the ZooScan and additional visual inspection, the planktic AP concentrations varied between  $0.0$  and  $18.54$  particles  $\text{m}^{-3}$ ; the lowest concentration was measured at station “exterior” during the first campaign (beginning July 2018), whereas the highest concentrations were associated with stations “interior 2” ( $18.54$  particles  $\text{m}^{-3}$ ) and “media 2” ( $12.97$  particles  $\text{m}^{-3}$ ) during the third sampling (mid-August 2018). The most identified AP types in plankton samples were fibers and threads (Fig. S2 A–D, supplementary information). In the macrozoobenthos samples, only a few hard AP

fragments were found, which were in a size dimension larger than any prey item ingestible ( $> 5$  cm) by the target life stage and species of this study; thus, these infrequent items were omitted from the analysis.

Due to the mismatch between planktic and benthic prey availability and the GIT contents of the fish, computation of selectivity indices such as Chesson's  $\alpha$  (Chesson 1978) could not be performed; consequently, the results of the zooplankton and macrozoobenthos community analyses are not detailed here. To put the results of the fish feeding preferences into perspective, however, it has to be noted that the environmental prey composition and availability (individuals  $m^{-3}$ ) showed no significant differences (ANOSIM  $R = -0.14$ ,  $P = 0.6995$ ), neither between the sampling sites nor the sampling campaigns in the beginning of July and middle of August 2018.

### Fish morphometrics

A total of 306 juvenile *D. sargus* were collected for this study. A detailed overview of the abundances and key morphometric parameters of the juvenile fish collected at the different sites at three sampling campaigns is given in Fig. 2 (see also Table S2, supplementary information). One juvenile *D. sargus* was collected at the station closest to the inlet to the Atlantic Ocean (“exterior”); during the first sampling campaign, this individual was smaller (SL = 15 mm) and of lighter weight (WW = 0.05 g) than all other juveniles collected at the remaining four stations. Across the three campaigns, a total of 81 white seabream were sampled at the “interior” stations (“interior 1”  $n = 46$ ; “interior 2”  $n = 35$ ) while the highest numbers of individuals were recorded at the intermediate stations: “Media 1” featured 65 individuals, whereas at “media 2,” 159 individuals were collected across all three sampling campaigns.

During the first sampling campaign, fish SL varied between 15 and 47 mm, while the minimum recorded WW was 0.05 g, and the highest WW was 3.24 g. There was no statistically significant difference in standard length or total wet weight during the first sampling campaign among the five different stations (Dunn's test,  $P > 0.05$ ). During the second campaign, juvenile seabream were larger and heavier than during the second campaign, with SL ranging between 22 and 61 mm, and WW varying between 0.25 and 7.17 g. During the second campaign, fish collected

at both “interior” stations were significantly larger and heavier in comparison to both “media” stations (Dunn's test,  $P = 0.001–0.03$ ). During the third campaign, no significant differences in standard length or total wet weight were detected among the different stations. The variation in SL was 29–62 mm, and WW ranged between 0.67 and 8.38 g.

### Dietary preferences of juvenile white seabream

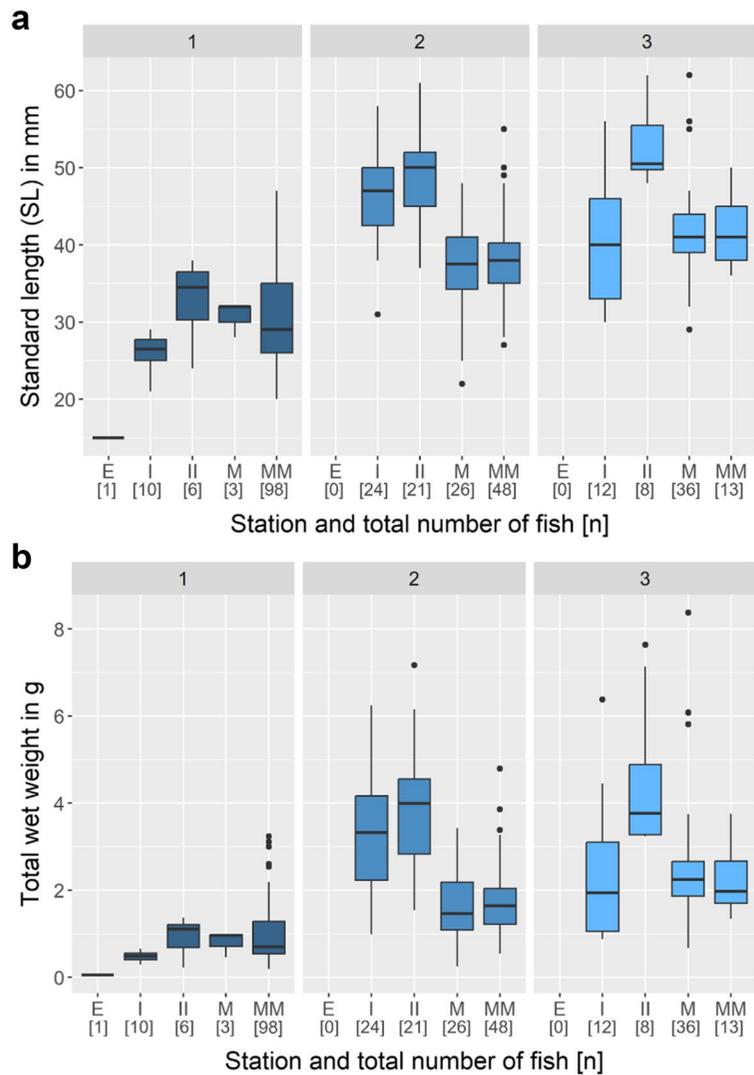
#### *Prey items*

The GIT analysis of a subset of white seabream ( $n = 147$ ) revealed a variety of different natural prey items, dominated by crustaceans, detritus, and marine flora (algae and seagrass), along with artificial items, namely AP fibers and fragments (Table 2). Crustaceans were the most abundant prey item in frequency of occurrence (94.56%FO) and prey-specific abundance (90.45%P). Within this prey taxon, noticeable differences between %FO and %P were recorded for copepods (70.75%FO; 50.87%N), Gnathiidae (49.66%FO; 12.76%P), and decapod zoea stages (14.97%FO; 85.7%P). Comparable differences in the overall presence of a prey item in the diet and its relative abundance were also detected in other prey taxa, such as polychaetes, insects, and ascidian tadpoles (Table 2). While marine flora was present in more than 2/3 of the fish GIT, its %N was not computed as the amount of algae and plant material was numerically not quantifiable.

#### Feeding strategy and prey preferences

The feeding strategy and the importance of individual prey items in the diet of *D. sargus* are visualized in a modified Costello plot (Fig. 3) after Amundsen et al. (1996). Most prey items ingested by *D. sargus* are represented in the lower part of the graph. The location of the zoea stages in the upper left of the plot implies a high specific abundance and low occurrence, whereas copepods were ingested with intermediate specific abundance and high occurrence. Most prey items (except for Copepoda) were ingested with minor frequencies and abundances (prey item points located in lower left corner of the plot, including AP fibers and fragments) thus being of relatively low prey importance (dashed line). Though detritus and algae are not included in this representation due to their uncountable amount, they appeared with a high

**Fig. 2** **A** and **B** Comparative overview of standard length (SL in mm) and total wet weight (in g) recorded for 306 juvenile *D. sargus* at five distinct sampling sites in the Ria Formosa lagoon over three sampling campaigns (1–3) in July and August 2018. Station codes: E=exterior; I=interior 1; II=interior 2; M=media 1; MM=media 2. The total number of individuals assessed at each site during each sampling campaign is indicated in brackets below the station code



frequency of occurrence (52.38–85.03%FO) as well as in high quantities, constituting an important share of the GIT fullness.

Although prey item consumption (Fig. 4) shows major overlaps across the different stations in the nMDS plot (stress = 0.179), weak yet significant differences in prey uptake were detected in relation to sampling site and campaign (ANOSIM  $R = 0.3806$ ,  $P = 0.0001$ ). According to the indicator species analysis, 8 out of 46 prey items were significantly associated with one or several groupings ( $P < 0.05$ , see supplementary information for detailed results). During the first and second campaign, fish showed a preference for Gnathiidae ( $P = 0.0478$ ) across both “interior” stations and “media 1”. For juvenile white

seabream collected during the third campaign at the aforementioned stations, crustacean prey items were of significant importance ( $P = 0.0037$ ). Except for the first sampling in “media 1” and the one individual collected at “exterior,” algae were a key item across all sampling stations and campaigns in the diet of juvenile seabream ( $P < 0.0016$ ). At “media 2,” *D. sargus* collected during the third campaign had a significant uptake tendency for Decapoda zoea stages ( $P = 0.0001$ ), and insect prey, particularly of the hemipteran order ( $P = 0.0001–0.0279$ ) which were of no importance at any other site or campaign, explaining the more distinct aggregation of this group in the nMDS plot (“media 2.3,” blue rhombus, Fig. 4).

**Table 2** Diet composition of juvenile *Diplodus sargus* ( $n = 147$ ) collected in summer 2018 at five different sampling sites in the Ria Formosa lagoon, Portugal. \* Not identified; Nc = not countable

Prey item		Number of fish with prey <sub>i</sub>	Frequency of occurrence % FO	Total abundance of prey <sub>i</sub>	Prey-specific abundance (%P)	
Crustacea total		139	94.56	6111	90.45	
	Copepoda total	107	72.79	1968	48.72	
		Copepoda*	104	70.75	1783	50.87
		Harpacticoida	25	17.01	119	6.9
		<i>Caligus</i> spp.	9	6.12	66	29.46
	Amphipoda total	20	13.61	55	9.05	
		Amphipoda*	17	11.56	39	19.8
		Gammaridae	5	3.40	5	1.23
		Caprellidae	4	2.72	11	26.83
	Isopoda total	73	49.66	366	12.94	
		Isopoda*	1	0.68	1	3.57
		Gnathiidae	73	49.66	361	12.76
		Idoteidae	1	0.68	2	22.22
		Sphaeromatidae	1	0.68	2	4.55
	Tanaidacea	Tanaididae	5	3.40	5	1.11
	Decapoda	Zoea stages	22	14.97	3452	85.7
	Ostracoda		34	23.13	63	2.35
	Branchiopoda	Cladocera	1	0.68	1	0.24
	Maxillopoda	Cirripedia—cyprid larva	50	34.01	186	10.49
	Pantopoda		1	0.68	1	0.83
	Crustacea*		11	7.48	14	1.2
	Crustacea remains		36	24.49	Nc	Nc
Ascidiacea	Tadpole	15	10.20	90	8.91	
Actinopterygii		5	3.40	5	6.49	
Polychaeta total		23	15.65	19	11.31	
	Polychaeta*	11	7.48	14	11.48	
	Polychaeta remains	13	8.84	Nc	Nc	
	Polychaeta larva	1	0.68	2	22.22	
	Serpulidae	1	0.68	1	2.70	
	Glyceridae	1	0.68	2	7.14	
Insecta total		38	25.85	282	9.85	
	Insecta*	8	5.44	13	1.12	
	Insecta remains	13	8.84	Nc	Nc	
	Diptera total	26	17.69	68	4.29	
		Chironomidae—larva	19	12.93	56	5.97
		Dolichopodidae—larva	7	4.76	8	3.65
		Meta-adult*	2	1.36	2	0.3
		Larva*	1	0.68	2	40.0
	Hemiptera total	11	7.48	201	8.16	
		Hemiptera*	4	2.72	13	1.75
		Auchenorrhyncha	11	7.48	188	7.63
Mollusca		13	8.84	22	5.29	

**Table 2** (continued)

Prey item		Number of fish with prey <sub>i</sub>	Frequency of occurrence % FO	Total abundance of prey <sub>i</sub>	Prey-specific abundance (%P)
Gastropoda	Mollusca remains	9	6.12	Nc	Nc
	Planktic*	3	2.04	5	1.27
	Benthic*	2	1.36	2	9.52
Bivalvia	Planktic*	1	0.68	1	5.58
Miscellaneous					
	Various*	28	19.05	185	15.14
	Sand grains	73	49.66	Nc	Nc
	Eggs*	5	3.40	Nc	Nc
	Detritus	125	85.03	Nc	Nc
	Alga 1*	109	70.07	Nc	Nc
	Alga 2*	77	52.38	Nc	Nc
	Seagrass*	20	13.61	Nc	Nc
	Fish scales	111	75.51	Nc	Nc
Anthropogenic particles		34	23.13	56	4.19
	Fiber	30	20.41	47	7.31
	Fragment	7	4.76	9	1.16

### Uptake of anthropogenic particles

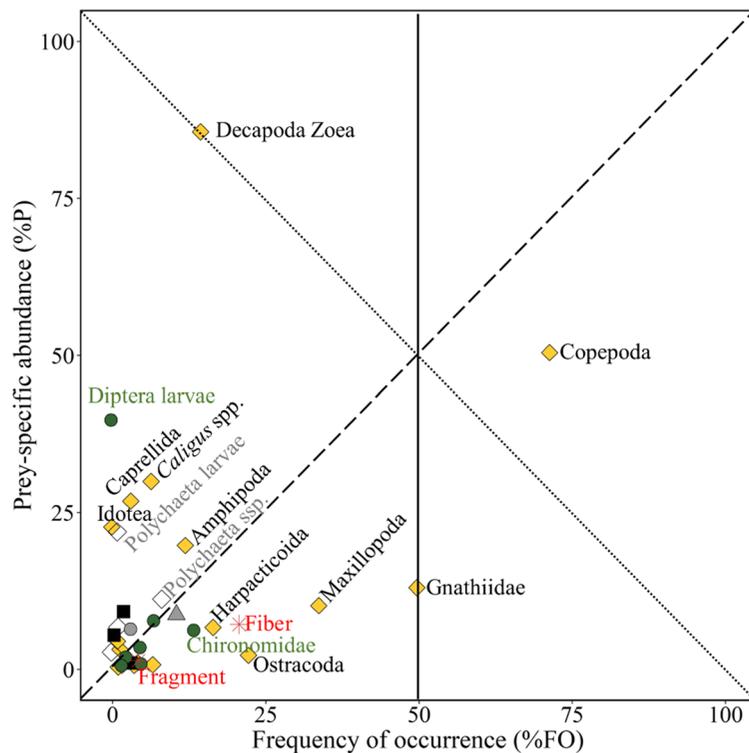
A total of 34 fish were found with AP in their GIT (23.13%FO). The proportion of AP-feeding individuals ranged between 13.64 and 31.03% across the four different stations located in the interior and middle sections of the Ria Formosa lagoon (Table 3), and the only white seabream collected at the station “exterior” did not have AP in its GIT. In total, 56 APs (4.19%P) were found in the GIT of the fish across the four different stations, most of them being fibers (47/56), and the remaining APs were fragments. The mean number of AP per AP-feeding fish was  $1.64 \pm 1.04$ . Most AP-feeding individuals had only one AP in their GIT (22 of 34 fish), two and three AP were found in five fish respectively, and four and five AP were detected in only one fish each. A pairwise comparison across sampling sites and campaigns revealed no statistically significant differences in the uptake of AP (pairwise *t*-test, adjustment: Benjamini and Hochberg,  $P = 0.46–0.93$ ).

Of the AP detected in the GIT, most were of blue color (19/56), followed by black (11/56) and red (11/56). A total of eight green/blue green APs were

found, additionally, four transparent fibers, two yellow fragments, and one purple fiber. Fibers were frequently found entangled within algae, detritus, and digested material (Fig. 5). However, the few, small fragments detected were also incorporated into the GIT content.

### Contextualization of anthropogenic particle uptake

The RDA biplot is given in Fig. 6. Locations are close to the intersection of the two axes, indicating that there is no difference in fragment or fiber ingestion between locations. The number of fibers is correlated with the first axis and fish morphometrics, namely wet weight (WW), GIT wet weight (WW\_GIT), and standard length (SL). The number of fragments is strongly associated with the second axis but is not strongly correlated with any of the explanatory variables. Contextualizing the presence of fibers and fragments in the GIT with other prey items (Fig. 7) revealed a strong, negative correlation between the two AP categories present in the GIT ( $r_s = -0.72$ ;  $P \leq 0.001$ ). The presence of fibers in the GIT was weakly to moderately positively correlated with algae (alga



**Fig. 3** Modified Costello plot, after Amundsen et al. (1996). Feeding strategy (solid line), niche width contribution (dotted line), and prey importance (dashed lined) of juvenile white seabream, *D. sargus*, are visualized based on the two-dimensional plot of prey-specific abundance (%P) and frequency of occurrence (%FO). The distribution of points along the axes and diagonals of the diagram provides the following information: specialized or generalized feeding strategy (vertical axis), rare or dominant prey items (diagonal from lower left

to upper right corner), and high between-phenotype or within-phenotype contribution to the niche width (diagonal from upper left to lower right corner). Sampling of juvenile fish took place in the Ria Formosa lagoon between July and August 2018. Red asterisk = items of anthropogenic origin; yellow rhombus = Crustacea; white rhombus = polychaeta; green circle = Insecta; gray circle = Actinopterygii; gray triangle = Ascidiacea; black square = Mollusca

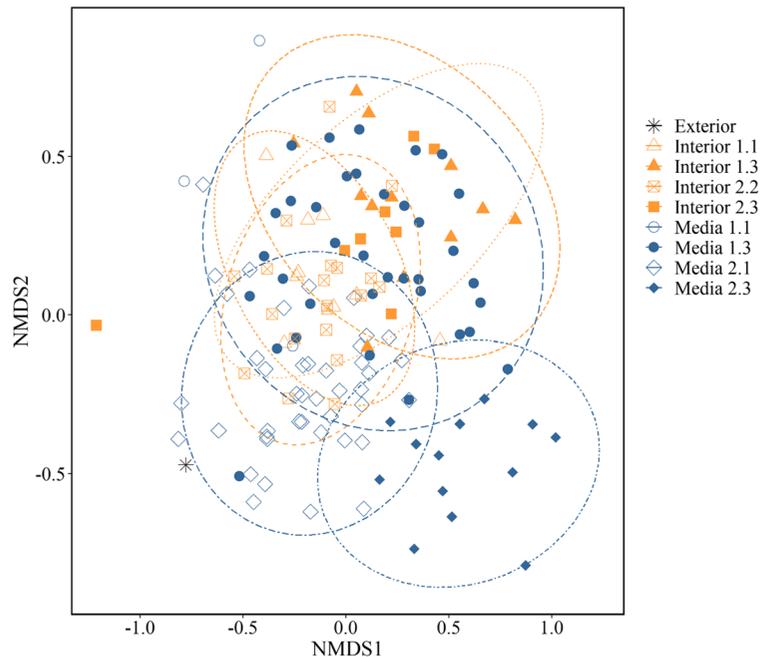
1:  $r_s = 0.44$ ;  $P \leq 0.01$ ; alga 2:  $r_s = 0.34$ ;  $P \leq 0.05$ ). Furthermore, the uptake of fibers was weakly positively correlated with the uptake of detritus ( $r_s = 0.36$ ;  $P \leq 0.05$ ). Due to their volume, both algae and detritus majorly contributed to the overall fullness of the GIT. Contrasting this, the presence of fragments was moderately negatively correlated to the uptake of detritus ( $r_s = -0.61$ ;  $P \leq 0.001$ ) and alga 1 ( $r_s = -0.4$ ;  $P \leq 0.05$ ), as well as weakly negatively correlated to the uptake of crustaceans ( $r_s = -0.35$ ;  $P \leq 0.05$ ). Additionally, a weak positive correlation was detected between the presence of fragments and ascidian tadpoles in the GIT ( $r_s = 0.35$ ;  $P \leq 0.05$ ). No other association of AP and natural prey items was of statistical significance.

## Discussion

### Quality control and MP identification limitations

Although the development and application of different digestion protocols (e.g., chemical digestion, enzymatic digestion) to extract MP from biological samples have been advanced over recent years, visual identification and optical analyses remain a rapid and relatively cheap method to classify and quantify plastic (Lusher et al. 2017 & 2020). However, without subsequent polymer analysis, these methodologies bear the risk of observer biases and identification inaccuracies, especially in relation to smaller MP size spectra (Lenz et al. 2015; Hanvey et al. 2017; Angelini

**Fig. 4** Non-metric multidimensional scaling (nMDS) analysis ordination biplot based on Bray–Curtis coefficient of similarities between presence/absence data of prey items found in the GIT of juvenile white seabream *D. sargus* at five different sampling sites in the Ria Formosa lagoon (stress=0.179, dimensions=3, non-metric fit  $R^2=0.968$ , linear fit  $R^2=0.799$ ; ellipses drawn based on 95% confidence interval where applicable). Station names, e.g., “interior 1,” are extended by the sampling campaign (i.e., 1, 2, 3)



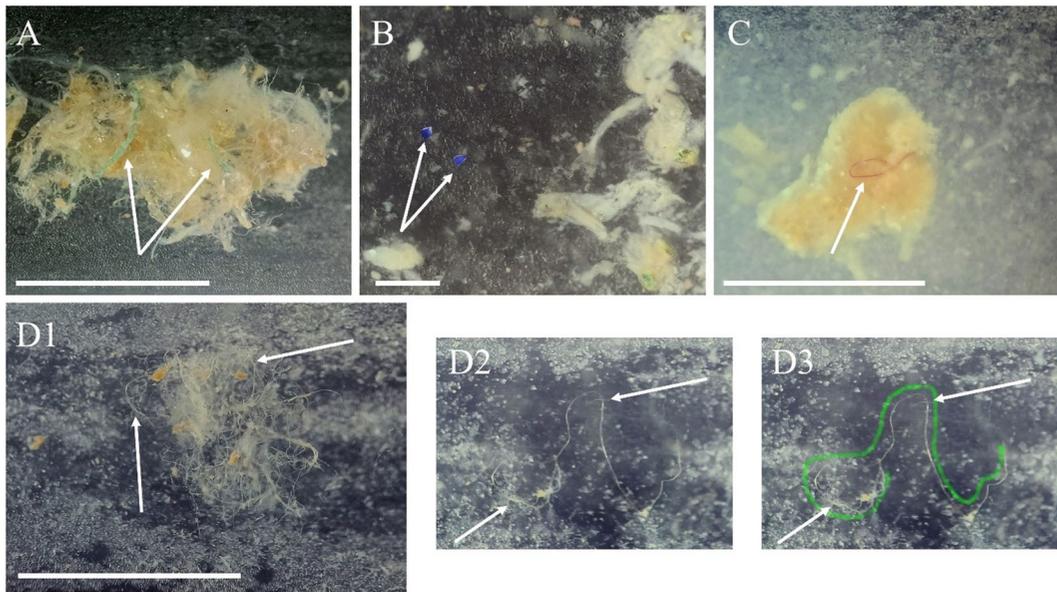
et al. 2019). In the present study, the abovementioned risks were partially overcome by applying strict quality control measures and a conservative identification and quantification approach. By omitting fibers and fragments of questionable origin, we present a trend in AP distribution in the Ria Formosa lagoon while not claiming to depict a precise quantification of MP abundances in the respective region. The overall goal of this ecological investigation was to provide a contextualization of AP uptake in a juvenile fish species from an estuarine nursery. Thus, the visual assessment of both natural and anthropogenic prey items

was necessary despite lowering the reliability of AP detection to fragments and fibers >150  $\mu\text{m}$ . Although the benefits of the ZooScan lie in the quick, semi-automated analysis of zooplankton samples, its limitations in MP detection are entailed in the depiction of images in gray-scales only (Fig. S3 A-C, supplementary information). A reliable AP identification based on color or shades (other than within the black-white-range) cannot be achieved and, therefore, requires additional visual inspection (Gilfillan et al. 2009; Pedrotti et al. 2014). Digestion of scanned subsamples, subsequent second image analysis on the ZooScan

**Table 3** Overview of anthropogenic particle (AP) uptake by juvenile *D. sargus* across four different stations sampled between July and August 2018. The single white seabream, collected at station “exterior,” had no AP in its GIT and is therefore, not represented here. %FO=frequency of occur-

rence;  $n_{\text{AP}}/n_{\text{total}}$ =number of AP-feeding individuals per total number of individuals investigated; the AP category (fiber or fragment, plus total numbers per category) detected in the GIT; the mean number of ingested AP plus standard deviation along with the range of AP ingested by AP-feeding individuals

Station	%FO ( $n_{\text{AP}}/n_{\text{total}}$ )	AP category ( $n_{\text{total}}$ )	AP mean $\pm$ SD (min–max)
Media 1	28.21% (11/39)	Fiber (12), fragment (3)	1.36 $\pm$ 0.81 (1–3)
Media 2	19.6% (11/56)	Fiber (15), fragment (4)	1.73 $\pm$ 1.01 (1–4)
Interior 1	13.64% (3/22)	Fiber (3)	1 $\pm$ 0 (1)
Interior 2	31.03% (9/29)	Fiber (17), fragment (2)	2.11 $\pm$ 1.36 (1–5)

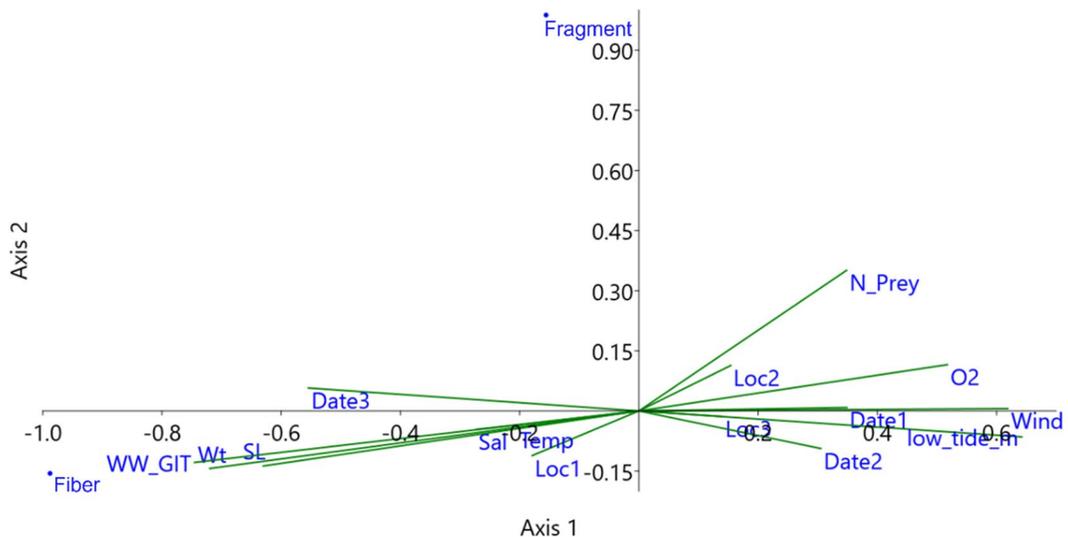


**Fig. 5** A–D Exemplary photographs showing different anthropogenic fibers and fragments detected in the GIT of juvenile white seabream, *Diplodus sargus*. **A** Green fiber entangled within algae and detritus. **B** Two blue fragments with partially

digested zoëa for size comparison. **C** Red fiber in digested material. **D1–D3** Transparent fiber entangled within algae—separated from algae—separated and outlined in green for enhanced visibility. Scale bar=500 µm

for verification along with a polymer identification, using methods such as Fourier-transform infrared spectroscopy (FTIR), would be advised

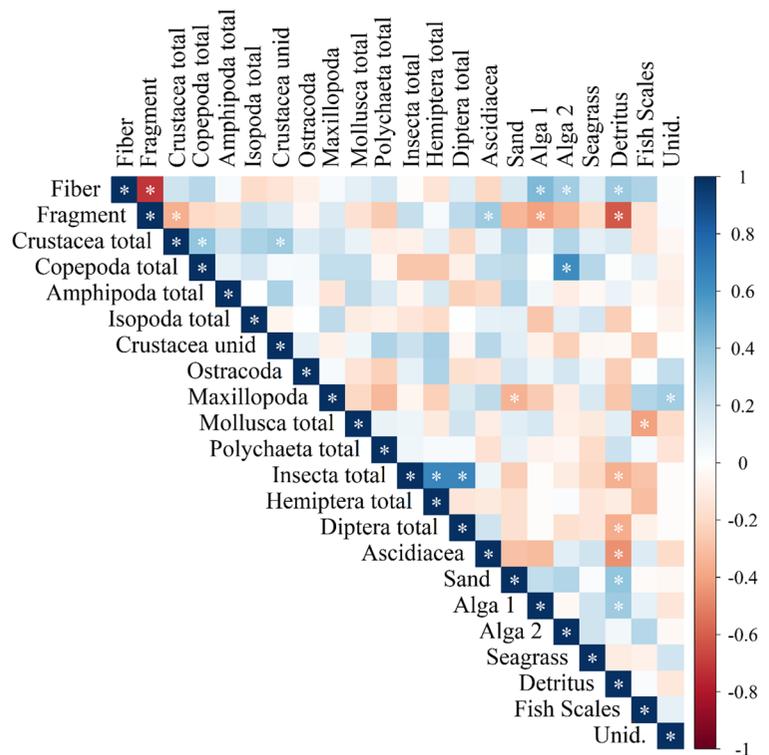
for future studies aiming at precise quantification of MP concentrations in coastal ecosystems (Lins-Silva et al. 2021).



**Fig. 6** Biplot of redundancy analysis (RDA) for abundance of plastic fibers and fragments found in the gastrointestinal tracts of *Diplodus sargus* across three sampling location categories (exterior, media, and interior) and over three sampling campaigns in the Ria Formosa lagoon. Abbreviations: SL: standard

length [mm]; Wt: total wet weight [g]; WW\_GIT: wet weight of the GIT [g]; Temp: temperature [°C]; Sal: salinity [PSU]; O2: oxygen [mg<sup>-1</sup>]; low\_tide\_m: depth at low tide [m]; Wind: wind speed [kmh<sup>-1</sup>]; N\_Prey: total number of prey items in GIT

**Fig. 7** Correlogram of AP uptake and other key prey items, computation based on Spearman rank correlation coefficient for AP-feeding individuals of *D. sargus*. Negative correlation ( $r_s - 1$  to 0) is indicated by shades of red, while positive correlation ( $r_s 0$  to +1) is given in shades of blue. Significant associations ( $P \leq 0.05$ ) are highlighted by \*



#### Distribution of seabream along spatio-temporal gradients of habitat quality

Members of the family Sparidae frequently use structurally complex, nearshore habitats as nursery grounds with juvenile sparids showing a high site fidelity (Erzini et al. 2002; Ribeiro et al. 2006; Abecasis et al. 2009; Vinagre et al. 2010); thus, the chosen field study design was considered appropriate to reflect potential differences in feeding preferences and AP uptake probabilities. The distribution of ELHS of white seabream in the Ria Formosa lagoon suggests a preference of this species for nursery sites inside the lagoon, which were generally characterized by higher temperatures and salinities along with lower concentrations of dissolved oxygen, and muddier, finer sediment.

The recorded fluctuations of abiotic parameters from inside the Ria Formosa lagoon to the inlet to the Atlantic Ocean verified the spatio-temporal trends previously described (Newton and Mudge 2003). As oxygen availability, salinity, and temperature significantly impact the swimming and feeding behavior of fish (Fry 1971; Blaber and Blaber 1980; Pandian and

Vivekanandan 1985), the AP uptake probability may also vary in relation to these environmental parameters as indicated by our results (Fig. 6).

The slightly higher values in fish standard length and wet weight recorded at both “interior” stations at the end of July 2018 may be indicative for the variability in ingress rates into the Ria Formosa, with individuals of high body condition and thus high swimming capabilities advancing to nursery sites further inside the lagoon (Baptista et al. 2019 & 2020). However, this observation needs further verification to clearly define the significance of different micro-habitats as nursery grounds inside the Ria Formosa and thus the potential installation of customized protection and conservation measures for the white seabream.

Next to the detrimental effects of climate change scenarios, anthropogenic disturbances, or pollutants entering the Ria Formosa (Bebiano 1995; Cortesão et al. 1986; Newton et al. 2003; Newton and Mudge 2005), plastic litter represents an additional stressor to this vital ecosystem and its biological communities which has been rarely accounted for in the past (Velez et al. 2020; Cozzolino et al.

2020; Oliveira et al. 2020). Regardless of the applied conservative approach in AP quantification and the potential methodological detection limits of the ZooScan, the results of this study suggest a tendency of higher AP concentrations at stations inside the lagoon, confirming a spatial trend that has been detected in the Ria Formosa lagoon by Velez et al. (2020) as well as in other estuaries worldwide (Lima et al. 2014; Hitchcock and Mitrovic 2019). Taking the niche partitioning behavior of seabream into consideration (Sánchez-Velasco and Norbis 1997; Erzini et al. 2002; Gamito et al. 2003; Ventura et al. 2015), the AP encounter probabilities across distinct seagrass meadows and thus the potentially detrimental effects for ELHS and recruitment success may vary between different seabream species. This variability may pose a currently unexplored risk to this viable ecological and economical resource for local artisanal fisheries and aquaculture (Leitão et al. 2009; Bonanno and Orlando-Bonaca 2020).

#### Feeding ecology of juvenile white seabream in the Ria Formosa

The GIT content analysis revealed a generalized, omnivorous feeding mode of juvenile white seabream in the Ria Formosa: floral, faunal, and detrital prey were ingested across the different sampling sites and campaigns, with slight differences in resource utilization detected between individuals of the different stations. Algae and seagrass were important prey items across all stations in July and August 2018, which is in line with previous studies describing the feeding preferences of white seabream (Sala and Ballesteros 1997; Figueiredo et al. 2005; Merciai et al. 2018). The importance of crustacean and insect prey organisms, which are less rapidly digested than smaller, softer items, may be slightly over-estimated (Windell and Bowen 1978; Buckland et al. 2017); however, the relevance of these prey taxa in the diet of *D. sargus* has also been reported before (Rosecchi 1987; Sala and Ballesteros 1997; Osman and Mahmoud 2009). Noteworthy is the presence of two fish ectoparasites, namely *Gnathiidae* and *Caligus* spp., in the GIT of the juvenile fish, further establishing evidence for a facultative cleaning behavior contributing to the trophic resource utilization of *D. sargus* (Rosecchi 1987; Mariani 2001; Neto et al. 2019).

#### Contextualization of anthropogenic particle uptake

Ichthyofaunal communities have been proven to be useful biological indicators to monitor ecosystem health and environmental quality (Whitfield and Elliott 2002; Ribeiro et al. 2008; Ramos et al. 2012). Thus, assessing the feeding ecology of ELHS of the omnivorous white seabream was considered an important first step towards a holistic understanding of the potential impacts of plastic pollution in an estuarine nursery ground. The assessment of the entire GIT content allows for the evaluation of inter-individual and species-specific prey preferences along with feeding habits potentially facilitating the disproportionate uptake of AP and enables a sound estimation of the ecological threat anthropogenic pollutants present (Ory et al. 2017; Cardozo et al. 2018; Lopes et al. 2020). As management and conservation measures rely on a sound scientific basis, the GIT analysis performed herein was complemented by the assessment of environmental parameters and AP availability, a necessary data integration only infrequently accounted for in previous studies on MP uptake by fish (Gamito et al. 2003; Cardozo et al. 2018; McGregor and Strydom 2020; Müller 2021; Wootton et al. 2021). Although several in situ studies on the feeding ecology of juvenile white seabream have been published over the past decades, MP uptake by this omnivorous species of commercial importance has been rarely investigated; thus, intraspecific comparisons of MP uptake can be drawn only to a limited extent.

#### Spatio-temporal factors

Across the different stations considered for this study, the proportion of individuals with AP detected in their GIT ranged between 13.64 and 31.03%FO, with a mean load of  $1.64 \pm 1.04$  AP per AP-feeding individual. The recorded proportion of AP-feeding fish in situ is slightly higher than the MP uptake verified by a laboratory feeding experiment conducted on juveniles of the same species. In the laboratory set-up, the ingestion rate of polystyrene particles (500–1000  $\mu\text{m}$ ) across the different treatment groups varied between 9.8 and 17.65%FO. However, higher mean MP loads per MP-feeding

individual as well as pronounced inter-individual differences in MP uptake were reported (Müller et al. 2020). A multi-species assessment conducted at the Egyptian coast found an MP-feeding proportion of 100%FO ( $n=40$ ) along with high average loads of  $3593 \pm 3985$  particles in juvenile *D. sargus* (Shabaka et al. 2020). Upon closer examination of the substantial variation in the reported %FO and MP loads in juvenile *D. sargus*, the studies showed substantial variation in several factors previously identified to impede the comparability of findings. Next to the study/sampling characteristics (i.e., number and age/size of fish examined, sampling location), the applied quality control measures varied. Moreover, there were remarkable differences between the MP detection and quantification methodologies, as well as the MP (or AP) size ranges and types considered across the investigations (Collard et al. 2019; Markic et al. 2020).

No significant correlation between AP uptake and station was detected, presumably due to the lack of pronounced differences caused by the rather short time frame of the field study along with the high turnover rate of the water volume in the Ria Formosa during each tidal cycle. Previous investigations from estuarine environments were also not able to establish a significant correlation between MP uptake and environmental parameters even if a spatio-temporal pattern of MP ingestion was verified (e.g., Ferreira et al. 2016; Vendel et al. 2017; Silva et al. 2018).

The observation of higher loads of anthropogenic fibers than anthropogenic fragments in the GIT of the fish agrees with previous investigations reporting fibrous MP to be more abundant both in the marine realm (Rochman et al. 2015) and in fish GIT (Lusher et al. 2013; Bessa et al. 2018). Given the intense commercial fishing activity in the Ria Formosa lagoon and adjacent coastal waters, the most probable source of the fibers is fishing gear. Although the threads frequently detected in the plankton samples (Fig. S3 B+C, supplementary information) were beyond the ingestible size range for juvenile white seabream, their fragmentation products, such as smaller fiber bundles or individual fibers (Fig. 5 D1–D3), were found in the GIT of the fish. Hence, the spatial trend described for bigger sized plastics may be still considered ecologically relevant when assessing the potential risk of micro-sized AP.

### Prey preferences and feeding mode

The role of feeding mode in MP uptake is still debated and no common consensus has been achieved (Mizraji et al. 2017; Markic et al. 2020). The present study detected correlations between the ingestion of fibrous AP and the presence of vegetal prey as well as detritus. The correlation between uptake of AP fibers and fish morphometrics can be explained by the fact that both prey items majorly contributed to the overall GIT fullness and GIT weight. Based on the observations made while analyzing the GIT contents, anthropogenic fibers seemed to be frequently attached or incorporated in detritus and vegetal materials, which may suggest a higher potential of herbivorous, detritivorous, or omnivorous fish species to take up synthetic fibers along with their natural, soft-bodied prey (Peters et al. 2017; van der Hal et al. 2020; Wootton et al. 2021). In benthic invertebrates, the combination of MP characteristics (i.e., size, shape) and the feeding habits of a species were more decisive in relation to MP ingestion than the trophic guild, a finding which still requires further verification for ichthyofaunal taxa (Piarulli et al. 2020). The sensory perception of prey before and during intake along with their specialized food handling apparatus, notably the different types of teeth, enables members of the sparid family to utilize a broad food spectrum. All prey items are initially sucked in, those without a carapace directly reach the pharyngeal jaws while the buccal jaws hold back hard-bodied prey items for seizing, crushing, and rejecting via the mouth (Vandewalle et al. 1995). Ontogenetic dietary shifts have been described for the white seabream, with larger individuals ( $> 150$  mm SL) showing a tendency to ingest hard-bodied prey, such as gastropods and echinoderms (Figueiredo et al. 2005). Owing to the specialized feeding mode, ontogenetic shifts in dietary preferences may not necessarily affect the amount and type of plastic intake with varying age/size of the fish. However, smaller individuals, as the ones investigated here, frequently use vegetal prey such as algae or seagrass which potentially bear the risk of accidental ingestion of incorporated fibrous AP. The present study detected only a small number of hard-bodied natural and artificial prey items in the GIT of the juvenile white seabream, which has been found to discriminate polystyrene fragments

from crustacean prey (Müller et al. 2020), further supporting the abovementioned specialization in food intake.

Considering the overall small amount of ingested AP (Table 3), as well as their rather negligible size relation in comparison to natural prey items and digested materials (Fig. 5), the overall importance of AP in the diet of juvenile white seabream appears to be marginal. This is also confirmed by the fact that neither anthropogenic fibers nor fragments were significant contributors to any observed trends in the analyses of feeding strategy and preferred prey uptake (Figs. 3 and 4).

#### Effects of AP ingestion for juvenile white seabream

Most studies on AP uptake in fish examined adult individuals; thus, ambiguities remain regarding the potential effects plastic pollutants exert on vulnerable early life history stages in estuarine ecosystems (Browne et al. 2011; Steer et al. 2017; Vendel et al. 2017; Critchel and Hoogenboom 2018; Müller 2021; Wootton et al. 2021). Fish morphometrics such as standard length, weight, or condition factors can be used both as an explanatory variable for AP uptake and as an indicator for potential detrimental effects (Müller 2021). In the present study, the uptake of the few AP > 150 µm was positively associated with the number of prey items ingested as well as the GIT wet weight and GIT fullness, indicating a good nutritional status of the AP-feeding fish and an accidental co-ingestion of fibers along with voluminous vegetal or detrital prey. The present study could not verify the accumulation of AP and consequent blockage of the GIT of juvenile white seabream or any detrimental effects of AP ingestion on fish condition (e.g., inferior sizes and weights of AP-feeding fish).

Especially smaller plastic items, however, and associated chemicals may be translocated to other organs and tissues, posing an ecotoxicological risk to the fish and potentially also for human health upon consumption of contaminated tissues (Rochman et al. 2014; Avio et al. 2015; Barboza et al. 2020). The arising implications, particularly for commercially relevant fish species, need further evaluation by assessing the amount of smaller-sized microplastics and nanoplastics taken up accidentally along with the food, through trophic transfer or via drinking, making use of advanced identification and quantification

methodologies such as FTIR spectroscopy (Setälä et al. 2014; Roch et al. 2020; Veerasingam et al. 2020).

## Conclusion

Integrative studies, considering both the feeding biology of a species and the environmental availability of plastic pollution, have the potential to enhance our understanding of the extent to which fish deliberately or unintentionally ingest AP. Despite the continuously increasing number of studies on AP ingestion by fish, the extent of AP uptake by ELHS, being crucially dependent on nursery grounds in transitional ecosystems, needs further scientific exploration. Though juvenile white seabream show an omnivorous feeding habit and may encounter a gradient of habitat quality and elevated plastic encounter rates in their nurseries, their specialized mode of prey uptake may prevent them from ingesting high concentrations of hard-bodied AP fragments. Yet, they may still be prone to taking up fibrous AP as well as smaller-sized microplastics and nanoplastics along with their natural vegetal prey or detritus. Using ichthyofaunal communities as biological indicators for AP pollution in coastal ecosystems should be realized only under consideration of species-stage and life-stage-specific feeding modes and prey preferences as these factors may affect the uptake probability of different AP shapes, sizes, and colors.

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**Author contribution** Carolin Müller: conceptualization, methodology, validation, formal analysis, investigation, writing—original draft, visualization, funding acquisition. Karim Erzini: supervision, conceptualization, methodology, validation, investigation, resources, writing—review and editing, project administration, funding acquisition. Tim Dudeck: formal analysis, investigation, writing—review and editing, visualization. Joana Cruz: validation, investigation, writing—review and editing. Luana Santos Corona: investigation, writing—review and editing. Felipe Eloy Abrunhosa: investigation, writing—review and editing. Carlos Manuel Lourenço Afonso: validation, investigation, writing—review

and editing. Miguel Ângelo Franco Mateus: validation, investigation, writing—review and editing. Cristina Orro: investigation, writing—review and editing. Pedro Monteiro: investigation, writing—review and editing. Werner Ekau: supervision, conceptualization, methodology, validation, resources, writing—review and editing, project administration, funding acquisition.

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**Data availability** The data that support the findings of this study are available from the corresponding author upon reasonable request.

The authors declare no competing interests.

## Declarations

**Ethical approval** The collection of samples was approved by the Instituto da Conservação da Natureza e das Florestas for Carolin Müller, Karim Erzini, and Isidoro Costa. Treatment of samples was under the Animal Care regulations at the respective research institutions.

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