

A new insight into the influence of habitat on the biochemical properties of three commercial sea cucumber species

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Received: 12 May 2018 / Accepted: 9 November 2018
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Abstract This work makes a comparative evaluation of the biochemical profile of three sea commercial cucumber species (*Holothuria mammata*, *H. polii* and *H. tubulosa*) caught from different locations of the Mediterranean Sea (SE Spain). All species had high levels of moisture (from 73.6% in *H. mammata* to 81.2% in *H. tubulosa*), crude ash (from 9.61% in *H. mammata* to 14.7% in *H. tubulosa*) and protein (3.01% in *H. tubulosa* to 11.1% in *H. mammata*). They also had a low fat content, from 0.21% in *H. tubulosa* to 0.55% in *H. mammata*. *Holothuria polii* had intermediate values between the other two species, for all considered variables. All species had adequate protein/lipid ratios (*H. mammata*, 20:1; *H. polii*, 23:1; *H. tubulosa*, 14:1) and low lipid levels, enriched in omega-3 polyunsaturated fatty acids, especially arachidonic acid. The fatty acid profile suggests that *H. polii* is feeding on sediments more influenced by terrestrial inputs than the remaining species. *Holothuria mammata* and *H. tubulosa* are feeding on marine food sources mainly, but also with some terrestrial influence. The most abundant amino acids detected were alanine, arginine, glutamic acid, and glycine. All species had similar contents of essential amino acids (EAA) and ratios of EAA/non-essential amino acids. *Holothuria tubulosa* had a high content of toxic metals including Cr, Pb and Ni. This work highlights differences in compositional characteristics between different species of the same genus (*Holothuria*) from different locations.

Keywords Bioaccumulation · Fatty acids · Habitat · PUFAs · Toxic minerals

Introduction

Sea cucumbers (Echinodermata: Holothuroidea) are an important fishery resource, mainly in the Indo-Pacific region. They are exported to different locations, such as Hong Kong and Mainland China, where they are consumed as a dried (*bêche-de-mer*) or fresh products (Kasai 2003; Purcell et al. 2012; Purcell 2014). In those

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Asian markets sea cucumbers are a high value product and can reach up to USD300–500/kg (Purcell 2010; Wen et al. 2010). The global sea cucumber trade aiming the food market is controlled largely by China, Hong Kong SAR, Singapore and Taiwan (Jaquemet and Conand 1999; FAO 2014; INFOPECA and FAO 2013). However, sea cucumbers are being increasingly exported from Mauritania to the United States (Ann-Marie Holmes personal communication, from US Fish and Wildlife Service, US Department of Interior). Additionally, in Europe the areas and volume of sea cucumber capture are growing (Bordbar et al. 2011; FAO 2011; González-Wangüemert et al. 2016, 2018). The use of sea cucumbers has been mainly to fuel the increasing demand of Maiman based therapies also termed as zootherapies (Mahomoodally and Muthoorah 2014; Mootoosamy and Mahomoodally 2014).

It is estimated that 66 sea cucumber species are caught from more than 40 countries and exported to Asian markets (Conand and Byrne 1993; Choo 2008; Conand 2008; Kinch et al. 2008; Purcell 2010; Purcell et al. 2012). There has been a significant reduction in the sea cucumber stocks in many countries, or even a complete stock depletion in many tropical fisheries caused by the increasing market demand, unrestrained exploitation and/or inadequate fisheries (Purcell et al. 2012). Those situations boosted the catch and commercialization of species from alternative locations such as the Mediterranean Sea, being *Holothuria mammata*, *H. tubulosa* and *H. polii* some of the new target species (Aydin 2008; Aydin et al. 2011; González-Wangüemert et al. 2014; González-Wangüemert et al. 2016, 2018; González-Wangüemert and Domínguez-Godino 2016). Turkey is the main Mediterranean country actively harvesting and exporting sea cucumbers to Asian countries, including *H. mammata*, *H. polii*, *H. tubulosa* and *H. sanctori* (Çakli et al. 2004; Aydin 2008; González-Wangüemert et al. 2014; González-Wangüemert et al. 2015, 2016, 2018).

Sea cucumbers inhabit several sea floor habitats from intertidal zones to deep trenches, and from polar to tropical areas (Purcell et al. 2016). Those animals are mainly benthic and most of the exploited species are bottom detritus-feeders, consuming debris, bacteria and diatoms mixed with sediments on the seabed (Bruckner 2006; Purcell 2010; Purcell et al. 2016). Sea cucumbers have a low metabolism and its feeding source could influence their chemical composition, with strong implications for their nutritional properties (Aydin et al. 2011, Purcell et al. 2012, Yu et al. 2015). In this sense, this work makes a comparative evaluation of the chemical properties, including proximate composition, amino acids, fatty acids and essential and toxic metal contents of three commercial sea cucumber species (*H. mammata*, *H. polii* and *H. tubulosa*) caught from different locations of the Mediterranean Sea (SE Spain). Whenever possible results were compared with those obtained from the literature for the same species, caught from other Mediterranean areas such as Turkey and Italy. The amino acid levels of *H. mammata* is herein reported for the first time. Data was discussed through ecological and nutraceutical perspectives.

Materials and methods

Sample collection, processing and genetic identification

Holothuria mammata and *H. tubulosa* were collected in Los Cocedores (Murcia, SE Spain) while *H. polii* was caught in Los Urrutias (Mar Menor coastal lagoon, Murcia, SE Spain) (Fig. 1). All species were sampled during April of 2012 by scuba diving at 5 meters depth. Ten individuals of each target species were caught for analysis.

H. individuals had an average length of 19.32 ± 4.62 cm being located into rock crevice close to *Posidonia oceanica* meadows. *Holothuria tubulosa* specimens showing an average length of 24.71 ± 3.10 cm were caught on sandy and muddy bottom. *Holothuria polii* individuals with an average length of 14.02 ± 2.24 cm were sampled from Mar Menor coastal lagoon close to *Caulerpa prolifera* and *Cymodocea nodosa* meadows. This species is showing individuals with lower size than populations outside coastal lagoon (González-Wangüemert personal observation). Sea cucumbers were identified in situ considering its external morphology. Later this identification was confirmed using traditional taxonomic characters, such as ossicles, on the basis of the original descriptions of these species (Gmelin 1791; Grube 1840; Delle Chiaje 1824) and other relevant taxonomic contributions (Borrero-Pérez et al. 2009, 2010). This identification was further confirmed using genetic barcoding, by amplifying the cytochrome c oxidase I (COI) gene according to protocol from González-Wangüemert and Borrero-Pérez (2012). PCR fragments were sequenced and Basic Local Alignment



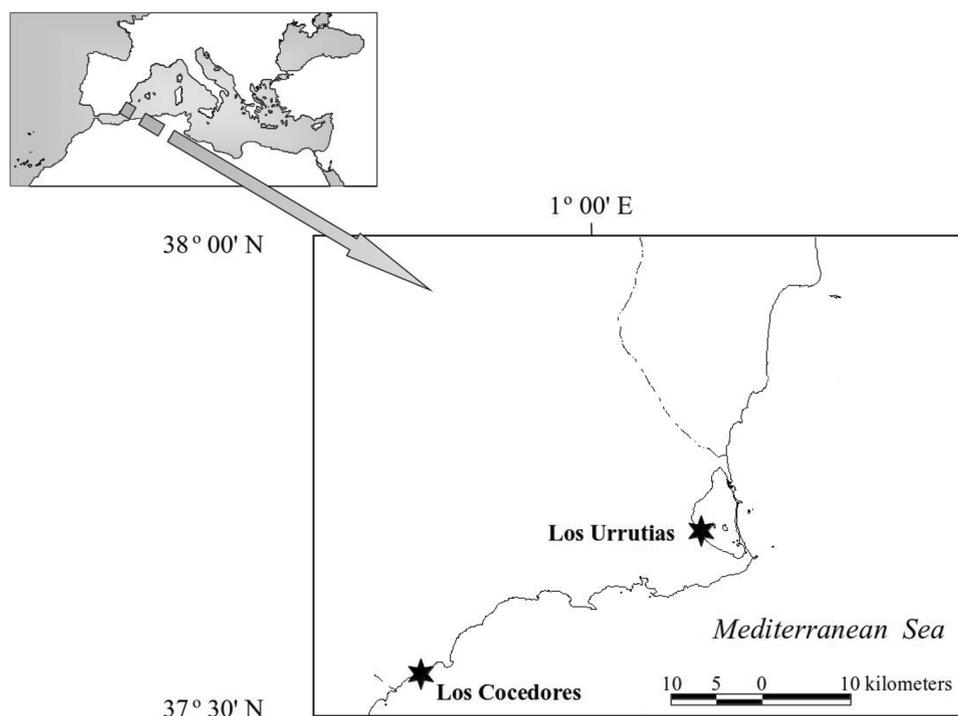


Fig. 1 Sampling sites of *H. mammata* and *H. tubulosa* (Los Cocedores/Almería, Spain) and *H. polii* (Los Urrutias—Mar Menor coastal lagoon/Murcia, Spain)

Search Toll (BLAST) was performed against GenBank nucleotide database (www.ncbi.nlm.nih.gov/genbank). BLAST finds regions of local similarity between sequences and calculates statistical significance of matches. BLAST can be used to infer functional and evolutionary relationships between sequences as well as help identify members of gene families.

For the biochemical analysis fresh adult individuals were gutted, all inner organs were removed and the body walls were rinsed with fresh water. Samples were frozen at $-20\text{ }^{\circ}\text{C}$, freeze-dried, ground into a fine powder, pooled and stored at $-20\text{ }^{\circ}\text{C}$ in tightly closed plastic bags until analyses. Each sample corresponded to ten adult individuals.

Biochemical analysis

Biochemical analysis included the determination of proximate composition, amino acids, fatty acid methyl esters (FAME) and minerals contents. These parameters were determined according to methods widely used to ascertain the biochemical profile of edible marine organisms in general (Njinkoue et al. 2016; Lah et al. 2017), and sea cucumber species in particular (Çakli et al. 2004; Zhong et al. 2007; Wen et al. 2010; Aydin et al. 2011; Roggatz et al. 2016; Gao et al. 2016).

Proximate composition

Moisture was determined by drying the samples at $52\text{ }^{\circ}\text{C}$ ($\pm 1\text{ }^{\circ}\text{C}$) until a constant weight was obtained (5 days). Ash was determined by incineration in a muffle furnace at $525\text{ }^{\circ}\text{C}$ until the samples burned completely (5 h, AOAC 1990; Gressler et al. 2010) and the crude protein content ($N \times 6.25$) was estimated by the macro-Kjeldahl method. For this method approximately 1 g of dried samples were digested using a catalyst mixture (1 pellet of kjeldahl catalyst 0.3% in $\text{CuSO}_4 \cdot 5\text{H}_2\text{O}$) and 25 mL of concentrated sulfuric acid in a digestion apparatus (BICASA Minerox MOD BE 97) under increasing temperature ($200\text{--}400\text{ }^{\circ}\text{C}$) to convert all nitrogen forms present in the samples into ammonium sulfate. Sodium hydroxide (35% solution) was then added to liberate the ammonia. The samples were distilled (P Selecta PRO-NITRO II) into 250 mL-



Erlenmeyer flasks containing 10 mL of boric acid (4% solution) and a bromocresol green indicator. The amount of nitrogen (% N) in the sample (corresponding to the amount of fixed ammonia), was determined by titration with 0.5 N HCL according to the following formula:

$$\%N = \frac{V_{\text{HCL}} \times f_{\text{HCL}} \times 0.014 \times 100}{P_{\text{sample}}}$$

where V_{HCL} is the volume of hydrogen chloride needed for the neutralization (mL), f_{HCL} the molarity of HCl used (mol L^{-1}) and P_{sample} is the exact weight of biomass (g) used for the analysis. The protein content was calculated by multiplying by 6.25, corresponding to approximately 16% of nitrogen content in proteins.

Crude fat was determined by a modified method of Bligh and Dyer (Pereira et al. 2013). In brief, a solution containing methanol, chloroform and water (2:2:1) was mixed with 100 mg of dried biomass and homogenised with an IKA Ultra-Turrax disperser. Then, samples were centrifuged and a known volume of the organic phase was moved into pre-weighed tubes. The solvent was then evaporated in a gentle nitrogen flow and the tubes were reweighed to calculate the lipid content. Special care was taken to obtain a full homogenization and performing repeated washings and extractions, having in mind that sea cucumbers generally have a low lipid content. Results are expressed as g per 100 g of wet weight biomass (WW) and dry weight biomass (DW).

Amino acids

Total amino acids were determined by gas chromatography/mass spectrometry (GC/MS) after sample derivatization with *N*-methyl-*N*-tert-butyltrimethylsilyltrifluoroacetamide (BSTFA) and trimethylchlorosilane (TMCS) according to Deng et al. (2005). In brief, dried samples (100 mg) and the internal standard (*L*-norvaline, 20 mg) were mixed with 500 μL of acetonitrile and 500 μL of MBSTFA + TMCS (99:1, v/v) in screw vials under microwave irradiation (750 W, 60 s). After cooling to room temperature, samples (1 μL) were injected at 260 °C on an Agilent GC-MS, with a temperature profile of 50 °C (1 min) and 10 °C min^{-1} to 300 °C (6 min). To identify and quantify the amino acids, the total ion mode was used. A set of standards of four different concentrations containing all 22 proteinogenic amino acids was prepared and measured, and calibration curves were generated for each amino acid. Results are expressed as percentage of total amino acid content.

Fatty acid methyl esters (FAME) profile

The FAME content was determined by a modified protocol from Lepage and Roy (1984), as described previously (Pereira et al. 2013). In short, aliquots of dried samples (100 mg) were mixed with acetyl chloride and methanol (20:1, v/v) in reaction vessels and homogenized with an IKA Ultra-Turrax disperser for 2 min. Then, 1 mL of hexane was added to the mixture and heated (100 °C, 1 h). One mL of distilled water was added to the mixture, followed by centrifugation and removal of the organic phase, which was dried with anhydrous sodium sulphate. Samples were finally injected in a Varian 450-GC/240-MS (Varian 450-Gas Chromatograph/240-MS IT Mass Spectrometer, Varian Inc., Palo Alto, CA, USA), equipped with a BR-5MS capillary column (30 m \times 0.25 mm internal diameter, 0.25 μm film thickness, Bruker). The injection temperature was 300 °C, and the trap, manifold and transfer line were established to 22, 50 and 250 °C, respectively. Helium was used as the carrier and the temperature program for the GC oven was 60 °C (1 min), 30 °C min^{-1} to 120 °C, 4 °C min^{-1} to 250 °C, 20 °C min^{-1} to 270 °C, and 2.5 °C min^{-1} to 300 °C. For identification and quantification of the FAME total ion mode was used. Because of differences in the response factors, for each FAME separate calibration curves were determined in triplicate, using the Supelco® 37 Component FAME Mix (Sigma-Aldrich, Sintra, Portugal) commercial standard. Values were expressed in amount ($\mu\text{g/g}$) of dry biomass.



Minerals

Minerals were analysed by atomic absorption spectrometry-AAS (GBC Avanta Sigma, Australia) provided with a deuterium background correction. Mg, Na, K, Ca, Fe, Mn and Zn were analysed by flame (F) AAS with an air-acetylene flame. Cd, Cr, Ni and Pb were analysed with electrothermal atomisation (ET) (GBC graphite furnace 3000) using an auto-sampler (PAL 3000) (Rogatz et al. 2016). The accuracy of the analytical procedure was assessed by the analysis of certified reference material, using Community Bureau of Reference BCR60 (*Lagarosiphon major*). Procedural blanks always accounted for less than 1% of the metal concentrations in samples. Values were expressed as g/kg dry biomass (Ca, Mg, Na and K) or mg/kg dry biomass (Fe, Mn, Zn, Cr, Pb, Ni and Cd).

Statistical analysis

Results were expressed as mean \pm standard error of the mean (SEM). Significant differences were assessed by analysis of variance (ANOVA) or by Student–Newman–Keuls post hoc test ($p < 0.05$) (software “R”, R Development Core Team 2013) with the packages GAD (Sandrini-Neto and Camargo 2011) and CAR (Fox and Weisberg 2011). For each species, three samples were analysed and the assays were carried out at least in triplicate.

Results

Morphological and genetic identification

Holothuria polii showed relative small buttons (ossicles) comparing with the other two species. *Holothuria polii* has buttons more rounded, with smooth surface and edged never wrinkled. *Holothuria tubulosa* shows buttons more slender with the surface an edges clearly wrinkled. *Holothuria mammata* was distinguished from *H. tubulosa* by its buttons, which are more elongated and slightly bigger.

COI gene amplification of the specimen caught in Los Urrutias (Mar Menor, SE Spain) allowed us to obtain a sequence of 484 bp in length. The BLAST in GENBANK, identified the individual from Los Urrutias as *H. polii* with a 100% maximum identity (100% query cover) with a specimen caught in Mar Menor (SE, Spain; EU750797) and 99% maximum identity (100% query cover) with a specimen sampled in Cabo de Palos (SE, Spain; GQ214758), but showing one mutational change in the 396 position. The haplotype for this specimen was recorded as EU750793. In the case of the individual found in Los Cocedores (Murcia, SE Spain) and identified as *H. tubulosa* using external morphology, the COI amplification (532 bp) and BLAST corroborated that identification, with a 99% maximum identity (100% query cover) with a specimen from Murcia (SE Spain; GQ214750) showing three mutational changes in 260, 452 and 521 positions; the haplotype from this specimen is GQ214748. Finally, the specimen identified as *H. mammata* (external morphology) also sampled in Los Cocedores, was assigned rightly to the *H. mammata* individual GQ214747 sampled in Murcia (SE Spain), showing 99% maximum identity (100% query cover) with only 3 mutational changes in the 12, 152 and 194 positions; this haplotype was recorded as GQ214746.

Proximate composition

Proximate composition of the three sea cucumbers species is shown in Table 1, and is expressed as % of total wet biomass (WW) and dry biomass (DW). Moisture ranged from 73.6% in *H. mammata* to 81.2% in *H. tubulosa*, and the crude ash content varied from 9.61% in *H. mammata* to 14.7% in *H. tubulosa*. In general all species had a low fat content, from 0.21% in *H. tubulosa* to 0.55% in *H. mammata*, and high protein levels, ranging from 3.01% in *H. tubulosa* to 11.1% in *H. mammata*. *Holothuria polii* showed intermediate values between the other two species, for all considered variables. All the species had adequate protein/lipid ratios (*H. mammata*, 20:1; *H. polii*, 23:1; *H. tubulosa*, 14:1).



Table 1 Proximate composition of the body wall of *H. mammata*, *H. polii* and *H. tubulosa*

	<i>H. mammata</i>		<i>H. polii</i>		<i>H. tubulosa</i>	
	WW	DW	WW	DW	WW	DW
Moisture ¹	73.6 ± 1.73 ^b	–	76.3 ± 1.52 ^b	–	81.2 ± 1.55 ^a	–
Ash ²	9.61 ± 0.32 ^b	34.4 ± 1.21 ^c	13.4 ± 0.43 ^a	56.54 ± 1.81 ^b	14.7 ± 0.34 ^a	78.1 ± 1.80 ^a
Protein ²	11.1 ± 0.27 ^a	42.0 ± 1.02 ^a	7.37 ± 0.24 ^b	31.09 ± 1.01 ^b	3.01 ± 0.23 ^c	16.0 ± 1.11 ^c
Fat ²	0.55 ± 0.05 ^a	2.08 ± 0.18 ^a	0.32 ± 0.03 ^b	1.35 ± 0.12 ^b	0.21 ± 0.04 ^b	1.11 ± 0.21 ^b

Data represent the mean ± SD ($n = 3$)

In the same row values followed by different letters are significantly different at $p < 0.05$ (one-way ANOVA with Student–Newman–Keuls post hoc test)

¹Percentage (%)

²g/100 g of wet weight (WW) and dry weight (DW)

Amino acids profile

The composition of total amino acids of dried sea cucumber is shown in Table 2, and results are given in % of total amino acid content. Nineteen amino acids were detected, including 8 essential (EAA) and 11 non-

Table 2 Amino acid profile of *H. mammata*, *H. polii* and *H. tubulosa* in % of total amino acid content

	<i>H. mammata</i>	<i>H. polii</i>	<i>H. tubulosa</i>
ALA	12.2 ± 0.3 ^a	15.1 ± 2.6 ^a	14.5 ± 2.7 ^a
ARG	6.8 ± 0.2 ^d	13.4 ± 2.4 ^a	12.9 ± 1.7 ^b
ASN	1.0 ± 0.2 ^a	nd	nd
ASP	5.3 ± 0.8 ^a	4.5 ± 0.4 ^b	4.8 ± 0.7 ^b
CYS	8.2 ± 2.9 ^a	2.4 ± 0.8 ^b	2.4 ± 0.9 ^b
GLU	12.1 ± 1.9 ^{ab}	8.4 ± 1.4 ^b	8.6 ± 2.1 ^b
GLN	2.3 ± 0.7 ^b	1.0 ± 0.2 ^b	1.3 ± 0.8 ^b
GLY	11.8 ± 0.6 ^a	10.6 ± 1.2 ^a	10.6 ± 1.1 ^a
PRO	8.2 ± 0.4 ^a	10.1 ± 2.4 ^a	9.5 ± 2.0 ^a
SER	3.2 ± 0.3 ^b	3.0 ± 0.5 ^b	3.5 ± 0.7 ^b
TYR	7.1 ± 1.3 ^a	3.8 ± 1.4 ^b	3.3 ± 1.4 ^b
Non-essential (NEAA)	78.5 ± 9.7 ^a	72.6 ± 13.0 ^a	71.9 ± 14.1 ^a
HIS	2.0 ± 0.6 ^b	nd	nd
ILE	2.0 ± 0.1 ^b	2.9 ± 0.1 ^a	3.3 ± 0.3 ^a
LEU	3.7 ± 0.1 ^c	5.4 ± 0.4 ^b	5.8 ± 0.6 ^b
LYS	2.2 ± 0.7 ^b	1.1 ± 0.2 ^b	1.6 ± 0.7 ^b
MET	1.0 ± 0.0 ^b	0.7 ± 0.3 ^b	0.1 ± 0.1 ^d
PHE	3.3 ± 2.0 ^b	8.1 ± 1.5 ^a	7.9 ± 2.3 ^a
THR	3.3 ± 0.1 ^a	3.4 ± 0.6 ^a	3.7 ± 0.7 ^a
VAL	3.7 ± 0.1 ^b	5.4 ± 0.6 ^a	5.5 ± 0.7 ^a
Essential (EAA)	21.4 ± 3.6 ^a	27.3 ± 3.6 ^a	28.0 ± 5.5 ^a
EAA/NEAA	0.27	0.38	0.39
LYS/ARG	0.33	0.09	0.12

Data represent the mean ± SD ($n = 2$)

nd not detected, ALA alanine, ARG arginine, ASN asparagine, ASP aspartic acid, CYS cysteine, GLN glutamine, GLU glutamic acid, GLY glycine, HIS histidine, ILE isoleucine, LEU leucine, LYS lysine, MET methionine, PHE phenylalanine, PRO proline, SER serine, THR threonine, TRP tryptophan, TYR tyrosine, VAL valine, TAA total amino acids, EAA essential amino acids, NEAA non-essential amino acids, nd not detected

In the same row, values followed by different letters are significantly different at $p < 0.05$ (one-way ANOVA with Student–Newman–Keuls post hoc test)

essential amino acids (NEAA). The most abundant amino acids were alanine, arginine, glutamic acid, and glycine. All species had similar contents of EAA (21.4–28%), and as a consequence, similar ratios of EAA/NEAA. The lysine/arginine ratios ranged from 0.09 in *H. polii* to 0.33 in *H. mammata*.

FA content and profile

The FA contents of the three sea cucumber species included in this work are represented in Table 3. They had high amounts of polyunsaturated FA (PUFA) ranging from 402 $\mu\text{g/g}$ in *H. polii* to 1028 $\mu\text{g/g}$ in *H. mammata*. PUFA accounted for 32–42% of total FA content. The least concentrated FA in these species was the SFA (321–526 $\mu\text{g/g}$) which accounted for 21–27% of total FA. Arachidonic acid (AA, C20:4n-6) was the most abundant FA in *H. mammata* (643 $\mu\text{g/g}$) and *H. tubulosa* (365 $\mu\text{g/g}$), while oleic acid (C18:1n-9c) was predominant in *H. polii* (246 $\mu\text{g/g}$). All species had similar amounts of EPA (115–221 $\mu\text{g/g}$). In contrast, regarding docosahexaenoic acid (DHA, C22:6n-3) *H. mammata* (89.7 $\mu\text{g/g}$) and *H. tubulosa* (63.7 $\mu\text{g/g}$) had the highest contents. Besides palmitic acid (C16:0), and the two monounsaturated FA (MUFA) eicosenoic (C20:1n-9) and nervonic acids (C24:1n-9), the other FA contributed to less than 5% of the total amount of FA.

Table 3 FAME profile ($\mu\text{g/g}$ DW) of *H. mammata*, *H. polii* and *H. tubulosa*

	Common name	<i>H. mammata</i>	<i>H. polii</i>	<i>H. tubulosa</i>
C14:0	Myristic acid	45.5 \pm 1.7 ^a	31.3 \pm 6.5 ^b	47.5 \pm 1.5 ^a
C15:0	Pentadecanoic acid	21.3 \pm 0.7 ^a	10.0 \pm 1.1 ^c	19.7 \pm 1.7 ^b
C16:0	Palmitic acid	107 \pm 4 ^a	40.9 \pm 9.2 ^c	91.9 \pm 11.3 ^b
C17:0	Margaric acid	25.3 \pm 2.0 ^a	14.6 \pm 2.9 ^b	24.8 \pm 3.4 ^a
C18:0	Stearic acid	88.7 \pm 9.6 ^a	39.9 \pm 8.5 ^c	73.1 \pm 6.5 ^b
C19:0	Nonadecanoic acid	45.9 \pm 3.8 ^a	27.7 \pm 4.2 ^b	39.2 \pm 3.2 ^b
C20:0	Arachidic acid	67.9 \pm 4.3 ^a	42.0 \pm 6.9 ^b	51.2 \pm 1.8 ^a
C21:0	Heneicosanoic acid	61.9 \pm 5.4 ^a	35.9 \pm 7.8 ^b	41.5 \pm 4.0 ^c
C22:0	Behenic acid	nd	37.4 \pm 7.0 ^b	42.8 \pm 3.3 ^a
C23:0	Tricosanoic acid	27.5 \pm 2.7 ^b	16.9 \pm 5.0 ^a	18.9 \pm 2.6 ^b
C24:0	Lignoceric acid	34.9 \pm 7.9 ^b	24.8 \pm 1.7 ^a	27.7 \pm 0.9 ^b
Σ SFA		526 \pm 35 ^a	321 \pm 17 ^c	478 \pm 29 ^b
C16:1	Palmitoleic acid	76.0 \pm 3.1 ^b	23.5 \pm 7.3 ^c	60.1 \pm 6.8 ^{ab}
C18:1n-9 c	Oleic acid	350 \pm 23 ^c	246 \pm 32 ^a	288 \pm 18 ^b
C18:1n-9 t	Elaidic acid	58.9 \pm 5.9 ^b	24.1 \pm 5.2 ^c	49.1 \pm 6.6 ^a
C20:1n-9	Eicosenoic acid	137 \pm 12 ^b	90.1 \pm 18.3 ^a	91.2 \pm 14.6 ^c
C22:1n-9	Docosenoic acid	67.0 \pm 6.3 ^b	44.1 \pm 7.0 ^a	43.0 \pm 4.8 ^c
C24:1n-9	Nervonic acid	182 \pm 19 ^a	85.6 \pm 17.4 ^b	99.1 \pm 10.9 ^c
Σ MUFA		870 \pm 113 ^b	514 \pm 79 ^a	631 \pm 90 ^b
C18:2n-6	Linoleic acid	35.8 \pm 3.4 ^a	16.1 \pm 1.8 ^b	27.3 \pm 1.5 ^a
C20:4n-6	Arachidonic acid (AA)	643 \pm 85 ^a	228 \pm 54 ^c	365 \pm 46 ^b
C20:5n-3	Eicosapentaenoic acid (EPA)	221 \pm 31 ^a	115 \pm 27 ^a	157 \pm 19 ^a
C20:2n-6	Eicosadienoic acid	38.2 \pm 3.3 ^a	16.3 \pm 1.8 ^b	30.4 \pm 3.6 ^a
C22:6n-3	Docosahexaenoic acid (DHA)	89.7 \pm 9.5 ^a	26.1 \pm 4.3 ^b	63.7 \pm 6.4 ^a
Σ PUFA		1028 \pm 202 ^a	402 \pm 75 ^b	643 \pm 116 ^{ab}
Σ n-3		311 \pm 32 ^a	141 \pm 27 ^b	220 \pm 20 ^a
Σ n-6		717 \pm 85 ^a	260 \pm 54 ^c	422 \pm 46 ^b
Σ n-6/n-3		2.3	1.8	1.9

Data represent the mean \pm SD ($n = 3$)

Σ SFA total saturated fatty acids (FAs), Σ MUFA total monounsaturated FAs, Σ PUFA total polyunsaturated FAs, Σ n-3 total omega-3 PUFAs; Σ n-6 total omega-6 PUFAs; Σ n-6/n-3 ratio of omega-6 to omega-3 fatty acids, nd not detected

In the same row, values followed by different letters are significantly different at $p < 0.05$ (one-way ANOVA with Student–Newman–Keuls post hoc test)



The amount of $\Sigma n-6$ FAME (260–717 $\mu\text{g/g}$) was always higher than the percentage of $\Sigma n-3$ (141–311 $\mu\text{g/g}$) in all species. The ratio between the $\Sigma n-6$ and $\Sigma n-3$ (1.8–2.3) was similar for the three species.

Minerals

As shown in Table 4, the sea cucumbers under study contain many dietary essential minerals, such as sodium (56.9–75.2 g/kg), potassium (3.37–5.58 g/kg), calcium (41.1–145 g/kg), magnesium (12.7–21.4 g/kg), iron (33.7–4400 g/kg), zinc (8.9–227.7 mg/kg), and manganese (4.53–86.6 mg/kg). *Holothuria polii* had the highest levels of calcium (145 g/kg) and magnesium (21.4 g/kg), while *H. tubulosa* had the highest content of residual metals; especially iron (4400 mg/kg), which was 100 times higher than the amounts found in the remaining species. A similar trend was observed with the other residual minerals such as manganese and zinc and also with the toxic metals chromium, lead and nickel (Table 4).

Discussion

The proximate composition of fresh sea cucumbers differs between species, catching season and feeding patterns (Bordbar et al. 2011). Holothurians are on the bottom of the food chain and help to recycle detritus, since most of them are deposit feeders that consume bacteria and diatoms mixed with sediments on the seabed (Purcell 2010; Purcell et al. 2016). Bacterial densities in sediments can differ greatly from site to site being generally associated with the type of organic compounds found in those sediments. For example, significant differences in chlorophyll *a* concentration and bacterial cell concentrations were observed between the Eastern and Western Mediterranean sediments and water, associated with a shift from carbohydrates to amino acids as the dominant biochemical components (Danovaro et al. 1999). Such diversity in feeding source could explain the variation in the chemical composition of sea cucumbers (Aydin et al. 2011).

The high-moisture levels registered in the sea cucumbers studied on this work are in accordance with the values found in the literature for the same species sampled in the Eastern Mediterranean regions (Aydin et al. 2011), and in the Southern Adriatic Sea (Sicuro et al. 2012), and also for other sea cucumber species belonging to the same genus and inhabiting the SW Mediterranean region, as for example, *H. arguinensis* (Roggatz et al. 2016). The crude ash contents was higher than the values reported by other authors in the same species from Turkey (Çakli et al. 2004; Aydin et al. 2011) and from the Southern Adriatic Sea (Sicuro et al. 2012), and also than those reported for important commercial sea cucumbers, such as *Stichopus japonicus* L. (Gao et al. 2016).

Table 4 Mineral content of the body wall of *H. mammata*, *H. polii* and *H. tubulosa* in g/kg dry biomass (Ca, Mg, Na and K) or mg/kg dry biomass (Fe, Mn, Zn, Cr, Pb, Ni and Cd)

	Symbol	<i>H. mammata</i>	<i>H. polii</i>	<i>H. tubulosa</i>
Calcium	Ca	41.1 ± 0.05 ^c	145 ± 0.13a	95.8 ± 0.86 ^b
Magnesium	Mg	12.7 ± 0.04 ^c	21.4 ± 0.10 ^a	16.6 ± 0.09 ^b
Sodium	Na	66.5 ± 0.45 ^b	56.9 ± 0.38 ^c	75.2 ± 1.01 ^a
Potassium	K	3.86 ± 0.01 ^b	3.27 ± 0.07 ^b	5.58 ± 0.15 ^a
Iron	Fe	33.7 ± 2.28 ^c	40.6 ± 0.30 ^b	4400 ± 77.94 ^a
Manganese	Mn	4.53 ± 0.14 ^c	46.2 ± 0.83 ^b	86.6 ± 3.47 ^a
Zinc	Zn	10.5 ± 0.08 ^b	8.90 ± 0.04 ^c	227.7 ± 0.48 ^a
Chromium	Cr	0.87 ± 0.02 ^b	0.77 ± 0.02 ^b	15.2 ± 0.42 ^a
Lead	Pb	nd	3.07 ± 0.02 ^b	6.5 ± 0.07 ^a
Nikel	Ni	0.46 ± 0.03 ^b	nd	2.5 ± 0.07 ^a
Cadmium	Cd	nd	0.09 ± 0.01 ^a	nd

Data represent the mean ± SD ($n = 3$)

nd not detected

In the same row, values followed by different letters are significantly different at $p < 0.05$ (one-way ANOVA with Student–Newman–Keuls post hoc test)



These results could be related to higher levels of minerals in the tissues, which could be linked with the higher availability of diatoms and microalgae in this geographical area (SW Mediterranean Sea; Rohling et al. 2015), which are consumed by cucumbers.

Sea cucumbers are generally characterized by low amounts of fat and high protein contents (Bordbar et al. 2011). The protein levels of the sea cucumbers were high, although slightly lower than those reported in the literature for different sea cucumber species, as for example, *Actinopyga mauritiana* and *Bohadschia argus* (Wen et al. 2010). *Holothuria tubulosa* had a lower protein content than the same species caught in Turkey (Aydin et al. 2011) and in the Southern Adriatic Sea (Sicuro et al. 2012). *Holothuria polii* had a similar crude protein content that the same species caught in the later area (Sicuro et al. 2012). The sampling location had little effect on the protein levels of *H. mammata*. The species under study had a low crude fat content, but higher than the values reported for the same species caught in the Southern Adriatic Sea (Sicuro et al. 2012), and also than the reference amount reported for other commercial sea cucumber species (Bordbar et al. 2011). The fat contents was, however, lower than the values reported for other seafood products, namely skipjack tuna (*Katsuwonus pelamis*; Khodabux et al. 2007). All species had an appropriate protein/lipid ratio which is nutritionally relevant in view of the important role of proteins for the human body.

The amino acid profile of the studied species agrees with the reported in literature for the same species (Sicuro et al. 2012) and also for different sea cucumbers genera, such as *Cucumaria* and *Actinopygia* (Zhong et al. 2007; Wen et al. 2010). The richness of glycine is highly relevant, since the intake of food rich in that amino acid is associated with the reduction in total cholesterol levels in serum (Aljawad et al. 1991; Ikeda et al. 1993). The lysine/arginine ratios are usually linked to hypocholesterolemic effects (Bordbar et al. 2011) and were lower than those usually described in different sea cucumbers species (Bordbar et al. 2011).

All species showed a higher proportion of PUFA relatively to other FA, while SFA were present in the lowest relative amount. PUFA are significant components in tissue repair and wound healing, inflammation prevention, reduction of the incidence of coronary heart disease and cancer (Fredalina et al. 1999; Gill and Valivety 1997; Roynette et al. 2004; Harper and Jacobson 2005). AA was the most representative FA, as observed for most species of sea cucumbers (Bordbar et al. 2011). This FA is a major component of the cell membrane phospholipids and has a potential role in growth and in blood clotting process leading to wound healing (Gill and Valivety 1997; Roynette et al. 2004; Harper and Jacobson 2005; Harris et al. 2008). The levels of EPA and DHA together ranged from 141 $\mu\text{g/g}$ in *H. polii* to 311 $\mu\text{g/g}$ in *H. mammata*. Both PUFAs are associated with several health benefits, from the prevention of cardiovascular diseases to the improvement of brain and eye function (Harris et al. 2008). The levels of such PUFA in these species are, however, very low considering the daily intake requirements advised by the WHO (around 500 mg/d of EPA + DHA). Nonetheless, the FA profile of these species, with $n-6$ to $n-3$ FA ratios (1.8–2.3) considerably lower than 10 is highly beneficial to human health (Sánchez-Machado et al. 2004; Schmitz and Ecker 2008). Hence, the consumption of these species is not likely to have an adverse effect on cardiovascular diseases.

When compared to previously reported studies, some significant differences can be deduced from the profiles obtained herein, probably due to the distinct geographical origins of the species used. The FA profile of *H. polii*, *H. tubulosa* and *H. mammata* reported by Aydin et al. (2011) for example, shows a similar FA pattern to our results in which AA and EPA are among the most abundant FA. However, contrary to our results, in that study, oleic acid was not a dominant FA and α -linolenic (C18:3 $n-3$) and eicosadienoic (C20:2) acids were also detected (Aydin et al. 2011). Our results were not comparable with those reported for *H. tubulosa* and *H. polii* caught from Italian Southern Adriatic Sea due to the different processing method used, which consisted in drying samples at 70 °C for 24 h prior to chemical analysis (Sicuro et al. 2012), since this drying method can modify the fatty acids contents of the samples (Bordbar et al. 2011).

Fatty acids have been increasingly used as biomarkers by ecologists to trace the origin and trajectory of organic matter in ecosystems (Parrish et al. 2000). According to Alfaro et al. (2006) long chain FA (with 24 or more carbons) are characteristic of terrestrial plants and thus their presence indicates organic material of terrestrial origin. On the other hand, the presence of EPA and DHA are indicative of diatoms and dinoflagellates, respectively, in the sea cucumbers food sources. As sea cucumbers are bottom feeders with low metabolism, it is probable that the FA composition of their body walls reflects the FA composition of their food source (Ramón et al. 2010; Aydin et al. 2011). In this context, species enriched in C24:0 FA would feed mainly on food sources of terrestrial origin, while those with prevalence of EPA and DHA would feed mainly on marine food sources. *Holothuria polii* was collected in a location more influenced by terrestrial inputs than



the remaining species. In fact, *H. polii* was collected from Los Urrutias (SW Mar Menor coastal lagoon, SE Spain), a location close to several temporary and stable watercourses harbouring waters and sediments from agriculture and urban areas (González-Wangüemert and Vergara-Chen 2014). *Holothuria mammata* and *H. tubulosa* sampled from Los Cocedores (Murcia, Mediterranean Sea, SE Spain) presented a mixed signature, because of the terrestrial inputs in this marine region are limited and only produced during the rain season. In general, bacteria present a distinct lipid profile, usually enriched in SFA and MUFA, contrary to microalgae, which are showing high content in PUFA (Lebreton et al. 2011).

Sea cucumbers usually contain a good mineral profile, with high amounts of Ca, Mg, Fe and Zn (Bordbar et al. 2011). In this work, *H. polii* had the highest content of Ca, Mg and K, but the highest levels of Fe and Zn were observed in *H. tubulosa* (Table 4). The human consumption of ingredients rich in nutrients such as K, Ca, Mg, Fe, Zn, and Mn can have positive health implications, since they are an essential part of many important enzymes and they play different roles as catalysts and antioxidants. *Holothuria tubulosa* had a very high content of these essential metals, but also the highest levels of toxic metals such as Cr, Pb and Ni. *Holothuria tubulosa* had higher amounts of Cr, Fe, Pb and Zn than the same species from Southern Adriatic Sea (Sicuro et al. 2012); while *H. polii* had lower levels of Fe and Zn, and a higher content of Pb (Sicuro et al. 2012). Working with the same species caught in the Northern Mediterranean Sea, Tunca et al. (2016) observed that *H. mammata* significantly accumulated some metals, such as Cd and Mn in its body wall, while *H. polii* accumulated Cu, Cd, Cr and Mn at the lowest amount. The levels of Pb found in our study for *H. tubulosa* are lower or similar than those reported for the same species caught in the NW Mediterranean (Warnau et al. 2006), while the Fe and Zn contents were much higher. The accumulation of minerals in living organisms is influenced by several factors, as for example, the environment, concentration and type of the mineral in the bottom, exposure time, presence of other elements in the environment, condition, metabolic rate and/or sex of the organism, physico-chemical characteristics of the medium and feeding habits (Warnau et al. 2006). The differences in the levels of particular minerals observed in our study can be ascribed, for example, to possible differences in feeding habits between species (Aydın 2008; Aydın and Erkan 2015). Previous works showed that the decomposition of plant tissues, including seagrass, may be a source of metals to the marine environment, which are released through leaching and mineralization, and sink after adsorption to litter (Weis and Weis 2004). The high metal contents in *H. tubulosa* tissues could, therefore, be partially explained by its feeding behaviour associated to sediment from seagrass meadows which are showing a potential high metal concentration (Marín-Guirao et al. 2005; Warnau et al. 2006). In fact, it has been suggested that due to the peculiar ecological characteristics of *H. tubulosa*, this species could be used to complement the small set of bioindicators available so far for surveying metal contamination in the *Posidonia oceanica* ecosystem from Mediterranean Sea (Warnau et al. 2006). *Holothuria tubulosa* was caught from Los Cocedores, a location with vast areas of *P. oceanica* and *C. nodosa* meadows; some authors have described the bio-accumulation of heavy metals on these marine phanerogam species (de Leon et al. 1982; Marín-Guirao et al. 2005). *Holothuria tubulosa* is usually associated with those meadows with 24 h feeding activity (Costa et al. 2014), while *H. mammata* is found during the day under the rocks, and only gets out at night for feeding in areas not close to *P. oceanica* (González-Wangüemert et al. 2016). Moreover, *H. tubulosa* could be a more efficient metal bio-accumulator than *H. mammata* (Warnau et al. 2006). Although 50% lower than in *H. tubulosa*, the Pb concentration in *H. polii* was significantly lower than those quantified on the body wall tissues of sea cucumbers from the Southern Adriatic Sea (Storelli et al. 2001). Moreover, *H. polii* was the only species in which Cd was found in concentrations similar to those reported by other authors (Storelli et al. 2001). *Holothuria polii* was collected in the south basin of the Mar Menor lagoon, which is one of the largest coastal lagoons in Europe. Mar Menor has fresh-water inputs and its southern part is enclosed by mountains, which were subjected to intense mining activity in the last two centuries (Marín-Guirao et al. 2005). Although mining activity stopped in 1991, during flood seasons the metals of mine tailings, including Pb and Cd, are released to Mar Menor and accumulate in the sediments and in the seagrass *C. nodosa* (de Leon et al. 1982; Marín-Guirao et al. 2005) which could explain the high levels of those elements detected on *H. polii*.



Conclusions

All species had high moisture, ash and protein levels, low lipid content, and adequate amounts of most essential amino acids, coupled with low lysine to arginine ratios. The FA profiles of all species were characterized by high levels of omega-3 PUFA and considerable amounts of DHA and EPA. *Holothuria polii* had a FA profile enriched in C24:0 FA, while *H. mammata* and *H. tubulosa* had a prevalence of EPA and DHA. *Holothuria tubulosa* had a high content of toxic metals such as Cr, Pb and Ni. This work pinpoints the influence of some physical parameters of the environment, namely mineral loads and terrestrial inputs, on the nutritional properties of sea cucumbers.

Acknowledgements This work was supported by the CCMAR/Multi/04326/2013, SEABIOMED (PTDC/MAR/103957/2008), CUMFISH (PTDC/MAR/119363/2010) and CUMARSUR (PTDC/MAR-BIO/5948/2014) projects funded by the Foundation for Science and Technology (FCT) and the Portuguese National Budget. Luísa Custódio and Mercedes González-Wangüemert were supported by FCT Investigator Programme (IF/00049/2012 and IF/00998/2014, respectively). The authors acknowledge Dr. Zélia Velez (CCMAR) and Dr. Maribela Pestana (FCT) for their assistance with the analysis of amino acids and total proteins, respectively.

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