

# Using trace elements in otoliths to discriminate between wild and farmed European sea bass (*Dicentrarchus labrax* L.) and Gilthead sea bream (*Sparus aurata* L.)

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**Abstract** Trace elements in otoliths of sea bass (*Dicentrarchus labrax* L.) and sea bream (*Sparus aurata* L.) from fish farms and coastal wild populations in the western Mediterranean Sea were analysed by inductively coupled plasma-mass spectrometry. Results showed that concentrations of Mg, K, and Mn differed significantly between wild and farmed sea bass, while concentrations of Mg, K, Mn, Fe, Zn, Sr, and Ba varied significantly between wild and farmed sea bream. Discriminate analysis and cross-validation classification showed that the trace element profile in otoliths can be used to separate farmed fish from wild stocks with high accuracy on both sea bass (individuals correctly classified: 90.7 %) and sea bream (individuals correctly classified: 96.6 %). Moreover, trace elements in otoliths resulted to be useful to discriminate among wild fish stocks within each species.

**Keywords** Microchemistry · ICP-MS · Escapes · Aquaculture · Fish stocks · Management

## Introduction

Mediterranean aquaculture has risen during the last decades mainly producing European sea bass *Dicentrarchus labrax* L. (176,970 t in 2015) and gilthead sea bream *Sparus aurata* L. (181,442 t in 2015) in off-shore fish farms (FAO 2016). Both species are also of high commercial interest for coastal Mediterranean fisheries (FAO 2014). However, the rapid expansion of aquaculture and the problems concerning escape events, have led to environmental problems on coastal ecosystems (Dempster et al. 2007). Escaped sea bass and sea bream disperse from farms, potentially mixing with local stocks and leading to potential negative genetic and ecological consequences through interbreeding, predation, competition for food or habitat, and the transmission of pathogens to native populations (Arechavala-Lopez et al. 2011, 2012b, 2013b, 2014; Šegvić-Bubić et al. 2014). However, conflicts with local fishermen and fish markets have also arisen, regarding escape events at farms. For instance, if the number of escapees is high, they may bias estimates of wild populations if not accounted for (Fiske et al. 2005), but also fisheries landings may be unbalanced, decreasing the captures and incomes from target species (Dimitriou et al. 2007; Toledo-Guedes et al. 2014a, b; Arechavala-Lopez et al. 2015). Furthermore, frauds mislabelling the farmed fish occasionally occur at markets, because of the price premium commanded by wild fish, affecting the guarantee of fish quality for the consumer (Bell et al. 2007; Morrison et al. 2007). Therefore, there is a necessity to develop verifiable and accurate methods to

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distinguish farmed from wild fish, which will help to increase the existing knowledge about the potential environmental and economic problems of escapees.

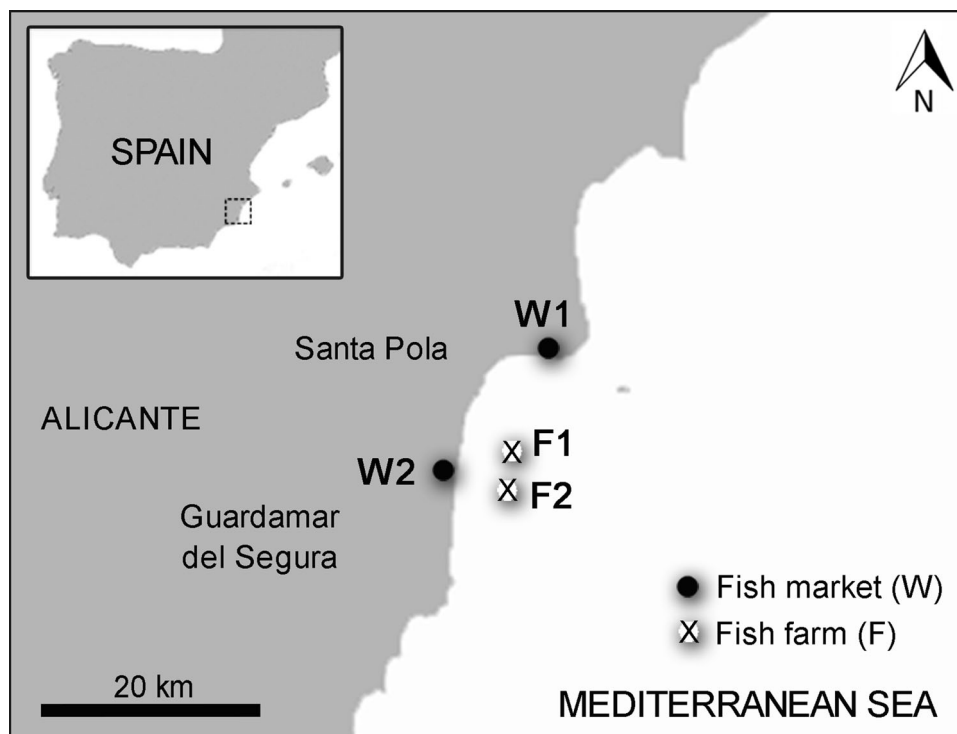
In the Mediterranean, diverse studies on sea bass and sea bream have quantified differences among wild and farmed fish through genetic analysis, chemical and molecular compositions, morphology, and organoleptic characteristics (Arechavala-Lopez et al. 2013a, and references therein). However, a wide range of applications, accuracies, time-consuming, and economic costs were showed among these techniques. Based on the assumption that farmed fish is grown in a different environment, stocking densities, and feeding regimes compared with wild fish, method based on the features of otoliths is regarded as very successful for stock discrimination (Campana and Neilson 1985; Campana et al. 2000). Otoliths are considered metabolically inert and grow throughout the life of the fish. Due to the trace element, concentrations in otoliths are a function of physiology and environment, they have been used as an indicator of fish populations inhabiting different environments or stock identity (e.g. Geffen et al. 1998; Thorrold et al. 1998; Campana et al. 2000; Gillanders and Kingsford 2000; Sanchez-Jerez et al. 2002; Rooker et al. 2003). Specifically, some studies have successfully attempted to discriminate through trace elements in otoliths, the wild and farm origin of Atlantic salmon *Salmo salar* (Veinott and Porter 2005; Perrier et al. 2011), trout *Salmo trutta* and *Oncorhynchus mykiss* (Zitek et al. 2010), and yellowtail kingfish *Seriola lalandi* (Gillanders and Joyce 2005). In addition, other studies used the differences on morphology and shape contours of the otoliths to successfully distinguish the wild or farm origin on Atlantic salmon (Hindar and L'Abéelund 1992), cod *Gadus morhua* (Higgins et al. 2010), but also on sea bass and sea bream (Arechavala-Lopez et al. 2012a). However, there is a lack of studies on the latter Mediterranean fish species, regarding the trace elements profiles in otoliths. Therefore, the aim of this study was to develop otolith trace elemental signatures of sea bass and sea bream to discriminate between wild and farmed stocks in the Mediterranean.

## Materials and methods

Fish sampling was carried out in the province of Alicante, southeast coast of Spain, and western Mediterranean Sea. A total of 30 sea bass (standard length, SL:  $40.73 \pm 0.81$  cm) and 30 sea bream (SL:  $35.43 \pm 0.21$  cm) were sampled from two neighbouring open-sea fish farms (F1 and F2), while 24 sea bass (SL:  $51.81 \pm 1.87$  cm) and 29 sea bream (SL:  $44.67 \pm 1.65$  cm) were sampled from local fish markets (i.e. Santa Pola and Guardamar del Segura; W1 and W2 respectively) (Fig. 1). The study area supports important fish farming and fisheries activities (trawl and artisanal fleets) of highly social and economic importance (García-Rodríguez et al. 2006). After fish sampling, both sagittal otoliths were collected through dissection using a stainless knife and disposable wooden chopsticks which were previously rinsed with ethanol 96°. The rest of the laboratory equipment was previously cleaned by soaking for 4 h with a chelating agent (EDTA 0.5 %) to prevent possible metal contaminations, and then moved to an acid wash in super-pure HNO<sub>3</sub> (10 %) for 24 h. Later, the equipment was carefully rinsed with 18 ohm reverse osmosis water and left to soak in sterilized conditions between sampled individuals. The collected otoliths were rinsed with 18 ohm reverse osmosis water to rehydrate the otolith covering matter for 24 h and weighted. Afterwards, each otolith was dipped in H<sub>2</sub>O<sub>2</sub> (3 %) for 5 min, then in HNO<sub>3</sub> (1 %) for 20 s, and, finally, thoroughly rinsed in 18 ohm reverse osmosis water. The otoliths were then left to dry completely and stored in 10 mL test tubes. Otoliths digestion was achieved by submerging them in 2 mL ultra-pure HNO<sub>3</sub> (10 %) and left overnight. Then, 8 mL of 18 ohm reverse osmosis water was added to each tube to raise the volume to 10 mL in each tube and drop the acid concentration to 2 % in every sample. Collecting and/or handling effects are likely to be minimal for all elements in our study based on similar treatment of fish after capture, chemical similarity, and the sterilized conditions.

Inductively coupled plasma-mass spectrometry (ICP-MS) was applied for trace elements determination on digested otoliths, which is the most commonly utilized technique for the otolith microchemical analysis. ICP-MS is a routine method widely used to analysed trace elements in otoliths of many fish species, which allows simultaneous determination of most elements within the periodic table with limits of detection below one part per billion (e.g. Campana et al. 1994; Campana and Gagne 1995; Campana 1999; Thresher 1999; Campana and Thorrold 2001; Sturgeon et al. 2005). Each sample was then analysed in triplicate, obtaining a mean value and a standard deviation per element and sample, which were standardized with the specific otolith mass





**Fig. 1** Study area and sampling sites (Western Mediterranean Sea, Alicante province, Spain). *Black spots* represent sampling fish markets (Wild fish: W1 and W2). *White spots with black cross* represent the sampling fish farms (Farmed fish: F1 and F2)

(Campana et al. 2007). Mean dried otolith weights ( $\pm$ SD) of wild and farmed sea bass were  $0.067 \pm 0.019$  and  $0.056 \pm 0.028$  g, respectively. Mean dried otolith weights ( $\pm$ SD) of wild and farmed sea bream were  $0.065 \pm 0.023$  and  $0.051 \pm 0.021$  g, respectively. The following element isotopes were analysed: lithium ( $^7\text{Li}$ ), beryllium ( $^9\text{Be}$ ), boron ( $^{11}\text{B}$ ), sodium ( $^{23}\text{Na}$ ), magnesium ( $^{24}\text{Mg}$ ), aluminium ( $^{27}\text{Al}$ ), phosphorous ( $^{31}\text{P}$ ), potassium ( $^{39}\text{K}$ ), scandium ( $^{45}\text{Sc}$ ), vanadium ( $^{51}\text{V}$ ), manganese ( $^{55}\text{Mn}$ ), iron ( $^{56}\text{Fe}$ ), cobalt ( $^{59}\text{Co}$ ), nickel ( $^{60}\text{Ni}$ ), copper ( $^{65}\text{Cu}$ ), zinc ( $^{66}\text{Zn}$ ), strontium ( $^{88}\text{Sr}$ ), palladium ( $^{105}\text{Pd}$ ), silver ( $^{107}\text{Ag}$ ), tin ( $^{116}\text{Sn}$ ,  $^{118}\text{Sn}$ ), barium ( $^{137}\text{Ba}$ ), rhenium ( $^{185}\text{Re}$ ), mercury ( $^{202}\text{Hg}$ ), titanium ( $^{205}\text{Ti}$ ), and lead ( $^{208}\text{Pb}$ ). These trace elements were quantified on the basis of peak areas and comparison with a calibrated curve obtained with the corresponding standards. Only those elements that were above the limit of detection of the ICP-MS in all or most ( $>90\%$ ) of the samples were considered suitable for statistical analysis. Then, concentrations of  $^7\text{Li}$ ,  $^9\text{Be}$ ,  $^{121}\text{Sb}$ ,  $^{185}\text{Re}$ ,  $^{202}\text{Hg}$ , and  $^{205}\text{Ti}$  were below the limit of detection of the ICP-MS for both fish species. Concentrations of  $^{11}\text{B}$ ,  $^{23}\text{Na}$ ,  $^{31}\text{P}$ , and  $^{52}\text{Cr}$  were not detected in more than 10 % of the samples both in sea bass and sea bream. Moreover, the standard reference material was found inaccurate for  $^{11}\text{B}$ ,  $^{23}\text{Na}$ ,  $^{45}\text{Sc}$ ,  $^{51}\text{V}$ ,  $^{105}\text{Pd}$ , and  $^{116,118}\text{Sn}$ , and therefore, those elements were discarded too. Then, a total of 13 trace elements were finally selected for further analysis:  $^{24}\text{Mg}$ ,  $^{27}\text{Al}$ ,  $^{39}\text{K}$ ,  $^{55}\text{Mn}$ ,  $^{56}\text{Fe}$ ,  $^{59}\text{Co}$ ,  $^{60}\text{Ni}$ ,  $^{65}\text{Cu}$ ,  $^{66}\text{Zn}$ ,  $^{88}\text{Sr}$ ,  $^{107}\text{Ag}$ ,  $^{137}\text{Ba}$ , and  $^{208}\text{Pb}$ . Non-parametric Mann–Whitney  $U$  test was used to test the variations in each trace element from fish otoliths (dependent variables), regarding the wild or farm origin of studied fish species (independent factors). Canonical discriminant function analysis and cross-validation classification were used to determine the efficacy of the trace elements composition in otoliths to discriminate between wild and farmed fish groups. Analyses of variances were carried out using the statistical software IBM SPSS Statistics 20 and discriminant analyses were carried out using the statistical software Canoco 4.5.

## Results

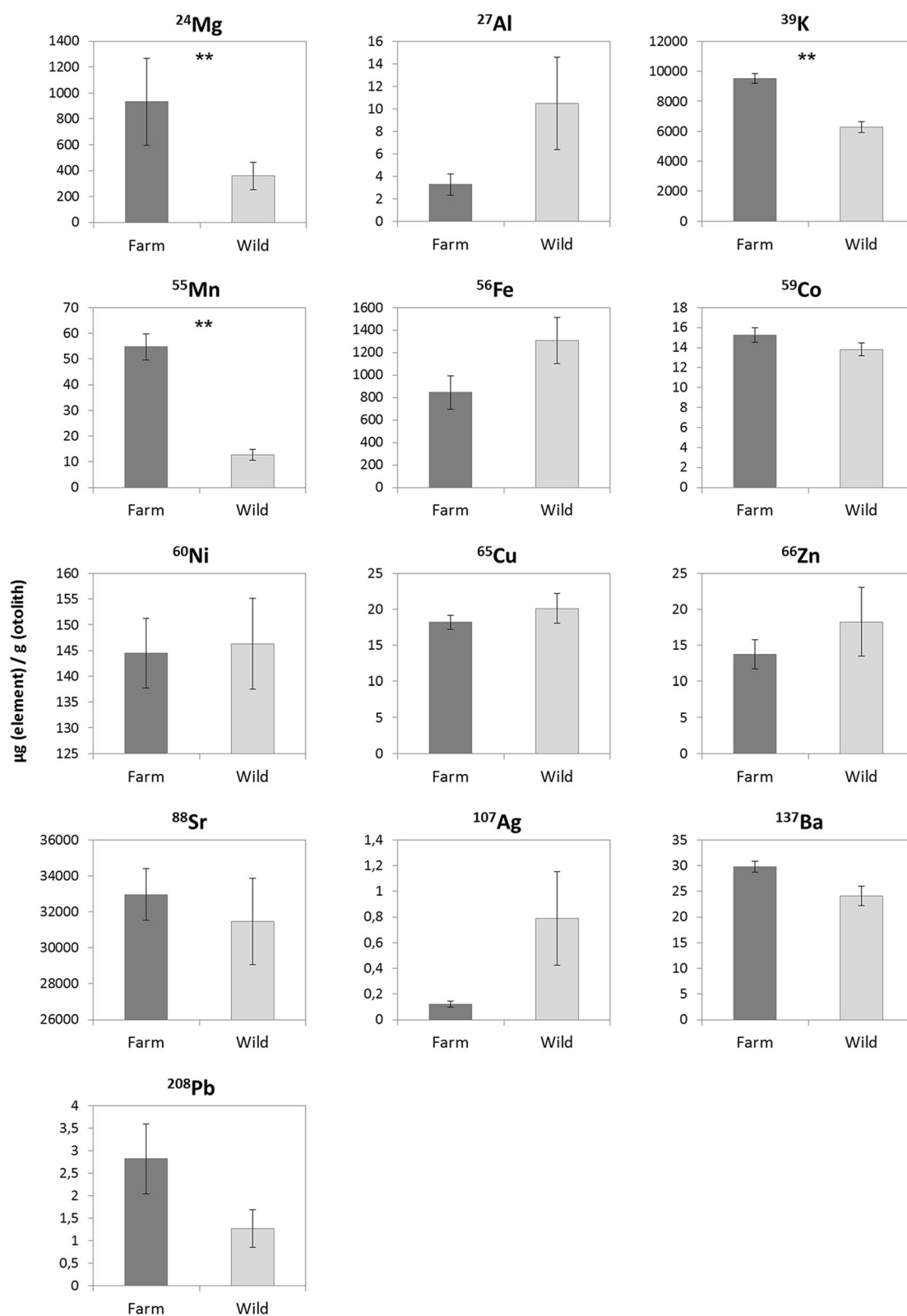
Concentrations of some trace elements differed significantly between wild and farmed individuals (Table 1). Non-parametric analysis (Mann–Whitney  $U$  test) determined that concentrations of  $^{24}\text{Mg}$  ( $Z = -4.613$ ,

**Table 1** Trace elements concentrations ( $\mu\text{g g}^{-1}$ ) in sea bass and sea bream otoliths from wild and fish farms, and results from non-parametric Mann–Whitney *U* test (*Z* value)

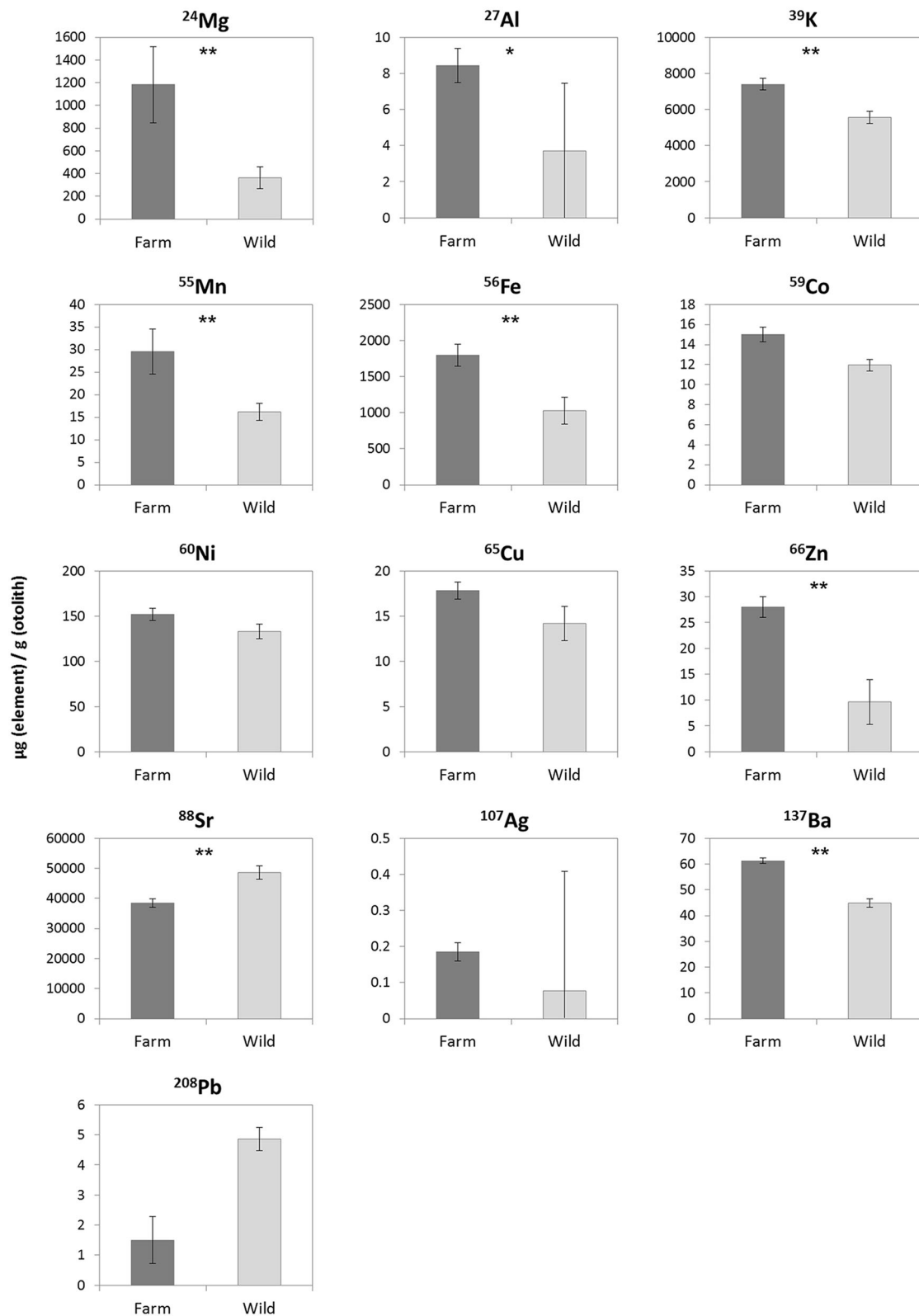
	Sea bass								<i>Z</i> (sig.)
	Wild ( <i>N</i> = 24)				Farm ( <i>N</i> = 30)				
	Min.	Max.	Mean	SD	Min.	Max.	Mean	SD	
<sup>24</sup> Mg	128.22	2769.88	358.20	522.25	0	9179.18	931.39	1839.49	−4.613**
<sup>27</sup> Al	0	92.54	10.48	20.13	0	20.32	3.26	5.24	−1.463
<sup>39</sup> K	1919.08	8514.12	6280.81	1836.88	6121.45	17141.76	9516.36	1739.28	−5.988**
<sup>55</sup> Mn	2.07	50.59	12.64	10.18	30.14	161.96	54.71	27.47	−5.971**
<sup>56</sup> Fe	0	3923.39	1307.58	1008.33	0	3142.76	845.32	822.01	−1.918
<sup>59</sup> Co	7.47	19.54	13.80	3.17	10.54	27.81	15.25	4.10	−1.132
<sup>60</sup> Ni	98.32	300.01	146.28	43.13	105.89	268.41	144.48	36.76	−0.052
<sup>65</sup> Cu	11.98	64.07	20.11	10.15	11.78	34.09	18.19	5.26	−0.905
<sup>66</sup> Zn	2.74	90.65	18.26	23.27	3.33	46.04	13.74	10.99	−0.001
<sup>88</sup> Sr	7121.73	52783.38	31464.48	11763.91	24973.46	67811.66	32958.18	7878.05	−0.522
<sup>107</sup> Ag	0	8.35	0.78	1.78	0	0.54	0.12	0.13	−1.119
<sup>137</sup> Ba	9.34	41.48	24.13	9.27	23.74	55.31	29.80	5.75	−1.915
<sup>208</sup> Pb	0.16	10.43	1.27	2.06	0.45	16.62	2.81	4.26	−1.132
	Sea bream								<i>Z</i> (sig.)
	Wild ( <i>N</i> = 29)				Farm ( <i>N</i> = 30)				
	Min.	Max.	Mean	SD	Min.	Max.	Mean	SD	
<sup>24</sup> Mg	129.31	888.62	365.64	261.40	268.77	7965.40	1184.49	2054.46	−3.154**
<sup>27</sup> Al	0	49.52	3.71	9.20	0	59.58	8.44	12.26	−2.175*
<sup>39</sup> K	1537.33	7987.50	5579.26	1277.16	0	28326.36	7399.01	4873.28	−3.093**
<sup>55</sup> Mn	4.18	39.26	16.13	7.96	11.37	103.95	29.57	23.15	−3.305**
<sup>56</sup> Fe	0	2135.01	1026.40	805.91	0	8100.84	1796.06	1353.90	−2.723**
<sup>59</sup> Co	3.90	15.43	11.95	2.12	9.18	52.83	15.01	8.78	−0.925
<sup>60</sup> Ni	67.69	162.91	133.17	17.76	91.18	518.64	152.11	80.31	−1.137
<sup>65</sup> Cu	11.32	19.41	14.19	2.33	9.50	64.91	17.82	11.24	−0.728
<sup>66</sup> Zn	2.32	37.59	9.61	8.49	4.60	167.41	27.95	32.52	−4.109**
<sup>88</sup> Sr	7879.08	63016.06	48639.24	9427.77	19947.51	134430.22	38445.14	20938.70	−4.215**
<sup>107</sup> Ag	0	0.87	0.07	0.15	0	1.25	0.18	0.31	−1.518
<sup>137</sup> Ba	11.49	82.27	44.85	16.54	40.04	130.14	61.25	18.23	−3.972**
<sup>208</sup> Pb	0.33	26.28	4.85	6.48	0.51	3.30	1.49	0.61	−1.425

\* Significant difference *p* value <0.05\*\* Significant difference *p* value <0.01

*p* = 0.001), <sup>39</sup>K (*Z* = −5.988, *p* = 0.001) and <sup>55</sup>Mn (*Z* = −5.971, *p* = 0.001) in otoliths differed significantly between wild and farmed sea bass (Fig. 2; Table 1). Concentrations of <sup>24</sup>Mg (*Z* = −3.154, *p* = 0.002), <sup>27</sup>Al (*Z* = −2.175, *p* = 0.030), <sup>39</sup>K (*Z* = −3.093, *p* = 0.002), <sup>55</sup>Mn (*Z* = −3.305, *p* = 0.001), <sup>56</sup>Fe (*Z* = −2.723, *p* = 0.006), <sup>66</sup>Zn (*Z* = −4.109, *p* = 0.001), <sup>88</sup>Sr (*Z* = −4.215, *p* = 0.001), and <sup>137</sup>Ba (*Z* = −3.972, *p* = 0.001) were significantly different between wild and farmed sea bream otoliths (Fig. 3; Table 1). The rest of the trace elements analysed did not showed any significant difference (*p* < 0.05) between fish origins for both fish species (Figs. 2, 3; Table 1). Considering the 13 selected trace elements, canonical discriminant analysis for sea bass otoliths showed that the 87.9 % of total variance among sampling groups was explained by two main axes (Fig. 4a). Among the elements in sea bass otoliths, concentrations of <sup>55</sup>Mn, <sup>39</sup>K, and <sup>137</sup>Ba were the most contributors for discrimination between wild and farmed sea bass individuals (Fig. 4a). Concentrations of <sup>60</sup>Ni, <sup>56</sup>Fe, <sup>65</sup>Cu, <sup>66</sup>Zn, <sup>59</sup>Co, <sup>27</sup>Al, and <sup>107</sup>Ag were the most contributors on

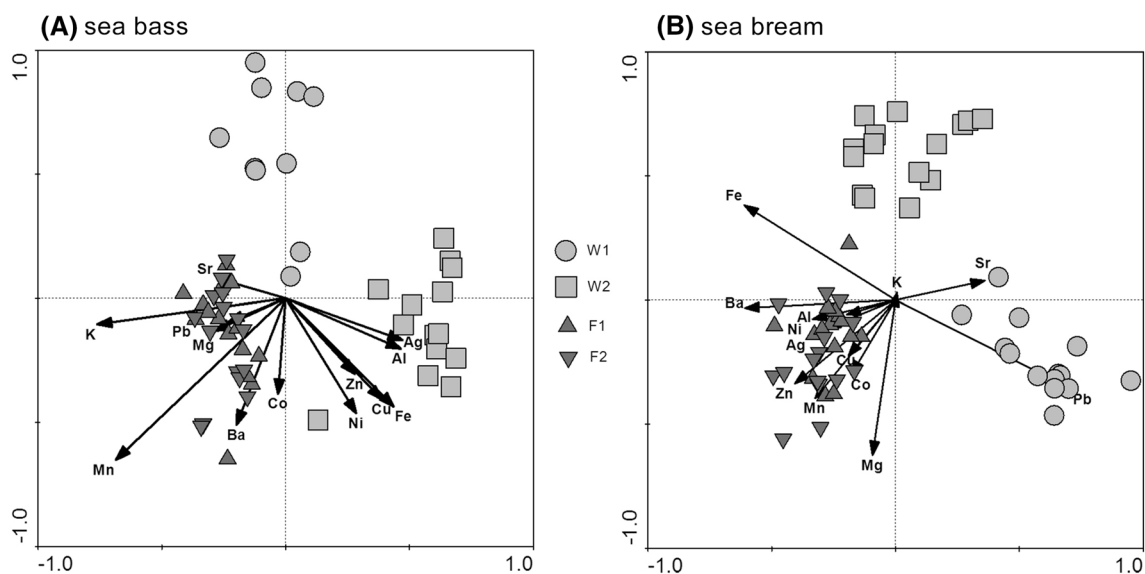


**Fig. 2** Bar plots (showing standard error in *whiskers*) comparing the trace element concentrations (µg g<sup>-1</sup>) in otoliths of wild and farmed sea bass. Significant differences from non-parametric Mann-Whitney *U* tests are shown with *asterisks*. \*Significant at the 0.05 level; \*\*significant at the 0.01 level



**Fig. 3** Bar plots (showing standard error in *whiskers*) comparing the trace element concentrations (µg g<sup>-1</sup>) in otoliths of wild and farmed sea bream. Significant differences from non-parametric Mann-Whitney *U* tests are shown with *asterisks*. \*Significant at the 0.05 level; \*\*significant at the 0.01 level





**Fig. 4** Scatter plot from canonical discriminant analysis for trace elements in otoliths of wild and farmed sea bass and sea bream groups. W1: group 1 of wild fish; W2: group 2 of wild fish; F1: group 1 of farmed fish; F2: group 2 of farmed fish. **a** Sea bass: axe X (PC1) explained the 51.5 % of total variance; axe Y (PC2) explained the 36.4 % of the total variance. **b** Sea bream: axe X (PC1) explained the 37.2 % of total variance; axe Y (PC2) explained the 36.3 % of the total variance

variations between wild sea bass sampled stocks, but no variations were found between farmed stocks (Fig. 4a). Cross-validation classification regarding trace elements profiles in sea bass otoliths showed that the 90.7 % of sampled individuals were correctly classified according to their wild or farm origin, while only the 61.1 % were correctly classified within sea bass sampling groups (Table 2). For sea bream otoliths, canonical discriminant analysis showed that the 73.5 % of total variance among sampling groups was explained by two main axes (Fig. 4b). Among the elements in sea bream otoliths, concentrations of  $^{55}\text{Mn}$ ,  $^{66}\text{Zn}$ ,  $^{88}\text{Sr}$ , and  $^{137}\text{Ba}$  were the most contributors on variations among wild and farmed individuals (Fig. 4b). Concentrations of  $^{56}\text{Fe}$ ,  $^{208}\text{Pb}$ ,  $^{24}\text{Mg}$ , and  $^{137}\text{Ba}$  in sea bream otoliths were the most contributors on differences between wild stocks, while no variations were detected between farmed stocks (Fig. 4b). Cross-validation classification regarding trace elements profiles in sea bream otoliths showed that the 96.6 % of sampled individuals were correctly classified according to their wild or farm origin, while the 89.8 % were correctly classified within sampling groups (Table 2).

## Discussion

The otolith grows throughout the life of the fish and the addition of new material will cause the average composition of the otolith to change with time and environment characteristics (Veinott and Porter 2005). Our data support that differences between wild and farm environments can be detected through the variations on trace elements compositions in sea bass and sea bream otoliths. In agreement, specific trace elements incorporated into the growing surface of the fish otolith reflect the physical and chemical characteristics of the ambient water, although not necessarily in a simplistic manner, the otolith elemental composition can serve as an environmentally induced tag of groups of fish (Campana et al. 2000). However, before an otolith elemental concentration can be applied as a biological tracer of stock mixing, it must be shown to differ among stocks or geographic locations (Campana et al. 2000). Regarding the trace elements of sea bass otoliths in this study,  $^{24}\text{Mg}$ ,  $^{39}\text{K}$ , and  $^{55}\text{Mn}$  concentrations were detected significantly higher in farmed fish otoliths than in wild ones. The previous studies showed significant differences between the farmed and wild origin in muscle and liver of sea bass for those specific elements, but also showed considerable divergence among the results, either accordance, discrepancies, or no significant results (Orban et al. 2002; Monti et al. 2005; Santaella et al. 2007; Fuentes et al. 2010; Mnari et al. 2010). However, a wider range of trace elements were found on the previous



**Table 2** Predicted group membership (%) by discriminate analysis classification with the cross-validation testing procedure, for the trace elements signature on wild and farmed sea bass and sea bream

	Wild		Farm		N
	W1	W2	F1	F2	
Sea bass					
Wild					
W1	90	0	10	0	10
W2	0	85.7	0	14.3	14
Total W	83.3		16.7		24
Farm					
F1	0	0	53.3	46.7	15
F2	6.7	0	66.7	26.7	15
Total F	3.3		96.7		30
Sea bream					
Wild					
W1	100	0	0	0	15
W2	0	93.3	6.7	0	14
Total W	96.6		3.4		29
Farm					
F1	6.7	6.7	86.7	0	15
F2	0	0	20	80	15
Total F	3.3		96.7		30

Sea bass: correctly classified within sampling groups: 61.1 %; correctly classified within origin groups: 90.7 %. Sea bream: correctly classified within sampling groups: 89.8 %; correctly classified within origin groups: 96.6 %

studies with significant differences on muscle and liver of sea bass from different origins, such as  $^{11}\text{Na}$ ,  $^{13}\text{Al}$ ,  $^{20}\text{Ca}$ ,  $^{29}\text{Cu}$ ,  $^{33}\text{As}$ ,  $^{34}\text{Se}$ ,  $^{48}\text{Cd}$ ,  $^{56}\text{Fe}$ ,  $^{66}\text{Zn}$ , and  $^{82}\text{Pb}$  (Alasalvar et al. 2002; Orban et al. 2002; Monti et al. 2005; Fernandes et al. 2007; Santaella et al. 2007; Ferreira et al. 2010; Fuentes et al. 2010; Mnari et al. 2010). For sea bream otoliths analysed in this study, concentrations of  $^{24}\text{Mg}$ ,  $^{27}\text{Al}$ ,  $^{39}\text{K}$ ,  $^{55}\text{Mn}$ ,  $^{56}\text{Fe}$ ,  $^{66}\text{Zn}$ , and  $^{137}\text{Ba}$  were significantly higher in farmed fish, while  $^{88}\text{Sr}$  was detected with higher concentrations in wild sea bream. The previous studies on sea bream muscle did not find significant differences on the trace elements analysed (Carpene et al. 1998; Minganti et al. 2010), except for  $^{33}\text{As}$  and  $^{80}\text{Hg}$  that higher concentrations were detected in both cases on wild individuals compared with farmed ones (Minganti et al. 2010). Divergent results might be explained by the existence of interannual variability in elemental composition of otoliths which may confound spatial interpretations and origin determinations (Gillanders 2002).

Thus, elemental signatures of Mg, Mn, Sr, or Ba may indicate migratory patterns (coastal and estuarine areas) of wild fish compared with stabled farmed sea bass and sea bream, but also differences on temperature and oxygen conditions among fish groups throughout their lives (Elsdon and Gillanders 2004; Gillanders 2005; Limburg et al. 2015). However, the use of combined elemental signatures is known to allow for greater distinction among fish than only one element, being preferable the use all elements at once, e.g., a multivariate elemental composition (Campana et al. 2000). The statistical evaluation of the data showed that differences in otolith chemistry enabled discrimination and retrospective assignment of fish to their wild or farm origin with high accuracy for both sea bass (90.7 % of individuals correctly classified) and sea bream (96.6 % of individuals correctly classified). Despite the fact that most of the previous studies gave contrasting results, most were also able to distinguish wild and farmed fish with great accuracy through a multivariate approach that accounted for a wide range of elements, indicating that the trace elemental profile might be more appropriate than the presence or absence of a specific quantity of an element (Arechavala-Lopez et al. 2013a). Multivariate approaches have been successfully applied for discriminating the wild and farm origin through trace elements in otoliths of other fish species (Gillanders and Joyce 2005; Veinott and Porter 2005; Zitek et al. 2010; Perrier et al. 2011), but also on sea bass and sea bream using other physical and chemical parameters, such as body





morphology, otolith shape, lipids and fatty acids compositions, stable isotopes, genetic markers, etc. (Arechavala-Lopez et al. 2013a; Šegvić-Bubić et al. 2014). Many of these parameters were highly recommended to identify escaped individuals from farms within wild stocks. However, it remains unknown for how long these parameters persist post-escape, which will influence their accuracy (Arechavala-Lopez et al. 2013a). Either physical parameters or biochemical components might change on escaped fish along the time, since they are directly influenced by the surrounding environment. However, diet has no significant effect on the incorporation of elements into fish otoliths (Marohn et al. 2009); hence, the main cause of variations of those parameters in a short-medium term might be any modification on habitat characteristics. The use of molecular genetic markers is likely to be the most suitable, non-destructive, and highly informative tool for genetic discrimination of wild and farmed fish, despite the heterogeneous mixed gene pool in wild sea bass and sea bream populations in the Mediterranean (Arechavala-Lopez et al. 2013a; Šegvić-Bubić et al. 2014). Unfortunately, genetic techniques are among the most expensive and time-consuming techniques currently available, and require specific knowledge making them unavailable to many sectors (Arechavala-Lopez et al. 2013a). Nevertheless, unique otolith fingerprints do not require genetic diversity among fish groups or populations, therefore combining chemical, morphological, and genetic data could enhance the discriminatory power of each technique.

Analysis of the chemical composition of otoliths has also been widely used as a technique to differentiate among fish stocks within the same species, reflecting differences in the chemical composition of the individual fish habitat or geographic-associated fish origin (Campana 1999; Thresher 1999). Gillanders et al. (2001) suggested that it is difficult to detect substantial differences in trace elemental signatures among different populations of wild fish in the Mediterranean, since they move among various coastal habitats. However, our results showed a clear discrimination between wild stocks for sea bass (85.7–90 % of wild fish correctly classified) and for sea bream (93.3–100 % of fish correctly classified). On the contrary, it was not possible to distinguish with accuracy between farmed stocks for both species. These results might reflect the proximity between farms with similar water chemistries in both areas, but also the similarities among fish culture techniques in the Mediterranean off-shore farms, in terms of net-pen environments, stocking densities, and feeding regimes. Thus, the analysis of trace elements composition in otoliths of sea bass and sea bream can be suggested as a useful method to distinguish between wild fish stocks from different geographical areas, but also to identify escaped fish from farms in the wild. Further researches are necessary to better understand the potential use of the trace elements in otoliths on fish ecology and populations dynamic studies (e.g., migrations, reproduction and growth studies of farmed, escaped, and wild fish), as well as to evaluate their applicability on sustainable coastal aquaculture and fisheries management.

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