

1 **Authentication of ready-to-eat anchovy products sold on the Italian market by BLAST**
2 **analysis of a highly informative cytochrome b gene fragment**

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27 **Abstract**

28 In this study, 111 ready-to-eat anchovy products were collected on the Italian market. The products
29 were molecularly identified through a BLAST analysis of a highly informative *cytb* fragment
30 amplified by a newly designed primer pair for the genus *Engraulis spp.* and the mislabelling rate was
31 assessed. In addition, the labels were analysed in the light of the current EU law. Despite only one
32 mislabelling case was observed (mislabelling rate 0.9%), which involved the substitution of the
33 European anchovy (*Engraulis encrasicolus*) with the low-valuable Peruvian/Chilean anchoveta
34 (*Engraulis ringens*), the molecular technique developed in this study was proved as suitable tool for
35 detecting species in processed anchovy products. It could be therefore applied to carry out more
36 extensive EU survey aimed at evaluating the mislabelling rate of such products, still poorly covered
37 by a targeted and clear legislation system.

38 **Keywords**

39 Anchovy; *Engraulis spp.*; anchoveta; species identification; *cytb*; mislabelling

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53 1. Introduction

54 Consumer protection throughout the single market is one of the EU policy benchmarks and the
55 obligation they are appropriately informed about the food they consume has been fixed as key point
56 by the General Food Law (Regulation (EC) No 178/2002). In this respect, the labelling of foodstuff
57 represents a basic tool for guaranteeing traceability throughout the production and distribution chain,
58 from the raw material supplier to the end-consumer. The Regulation (EU) No 1169/2011 on the
59 provision of food information to consumers has been precisely issued to achieve a high level of health
60 protection for consumers and to guarantee their right to information. Furthermore, such provisions
61 consider the well-known food fraud phenomena that over the years have undermined consumer
62 confidence and damaged the whole EU food supply chain and contribute to tackle this problem
63 (European Parliament Resolution 2013/2091).

64 Food fraud is committed when food is illegally placed on the market with the intention of deceiving
65 the customer, usually for financial gain (FAO, 2018). Intentional mislabelling and species substitution
66 in seafood products, which generally occur when low-value or less-desirable fish species are swapped
67 for more expensive varieties, are among the most reported kind of frauds at international level. Such
68 practices are favoured by the fact that seafood chain is often long and complex, involving many food-
69 business operators and an extremely wide range of species (i.e. 1200 different species being marketed
70 in Europe). In addition to financially damage consumers, frauds may constitute an important global
71 threat to sustainable fisheries as encouraging activities of Illegal, Unreported and Unregulated (IUU)
72 fishing (Helyar et al., 2014; Petrossian, 2015). Health issues may even occur if poisonous species
73 accidentally enter in the seafood chain (Giusti et al., 2016).

74 Available literature reported several mislabelling cases involving different kinds of products
75 marketed within the EU in the last decade (Garcia-Vazquez et al., 2010; Pardo, Jiménez, & Pérez-
76 Villarreal, 2016; Sotelo et al., 2018), even though an apparent reduction was observed during the last
77 years. This trend was mainly attributed to the recent efforts in EU legislation that have played a
78 pivotal role in shaping a more transparent market (Mariani et al., 2015). In this respect, in 2013

79 specific dispositions on seafood products labelling were provided by the Regulation (EU) No
80 1379/2013 on the common organisation of the markets in fishery and aquaculture products.
81 Mandatory consumer information should include (a) the commercial designation of the species and
82 its scientific name, (b) the production method (“caught”, “caught in freshwater” or “farmed”), (c) the
83 area where the product was caught or farmed, and the category of fishing gear used in capture of
84 fisheries, (d) whether the product has been defrosted and (e) the date of minimum durability, where
85 appropriate. However, as highlighted by D’Amico, Armani, Gianfaldoni, & Guidi (2016), the
86 exclusion of some kinds of prepared and processed products from the application of [this Regulation](#)
87 [\(EU\) No 1379/2013](#) represents a significant shortcoming. Basically, these products only fall under
88 the field of application of the Regulation (EU) No 1169/2011, which factually excludes the mandatory
89 information above listed for seafood products. For this reason, with the Resolution No 2016/2532,
90 the European Parliament encouraged the Member States, in the context of voluntary labelling, to state
91 all available information that enables the consumer to make an informed choice (European Parliament
92 Resolution No 2016/2532).

93 Semi-preserved anchovies are traditionally consumed within EU. Spain and Italy, the second and
94 the fourth anchovy world producers respectively, are even the major EU consumers, covering alone
95 the 71% of the total EU consumption (EUMOFA, 2018). In Italy, anchovies are mainly consumed in
96 form of ready-to-eat products, i.e. salted, marinated or in oil. Except for salted products, that fall
97 within the scope of Regulation (EU) No 1379/2013, tracing back the anchovy species used in
98 marinated or in oil products may prove tricky since, as detailed above, no information on the scientific
99 name and the catching area should be mandatory reported on the label of such products, except in
100 form of voluntary claims.

101 Recently, Velasco, Aldrey, Pérez-Martín, & Sotelo (2016), which applied DNA-based methods
102 for assessing the labelling accuracy of Spanish semi-preserved anchovies’ products, highlighted a
103 mislabelling rate higher than 15% and reported the Argentine anchovy (*Engraulis anchoita*) as the
104 most substituted species. [Cytb gene has been proved to be able to differentiate species belonging to](#)

105 the Engraulidae family (Santacarla et al., 2006; Velasco et al., 2016). In particular, it was *Cytb* was
106 chosen as molecular marker as reported as more suitable to discriminate some *Engraulis spp.* respect
107 to the cytochrome oxidase I (COI) gene (Jérôme et al., 2008).

108 In this study, a PCR primer pair was designed for amplifying a highly informative fragment,
109 proved as polymorphic among *Engraulis spp.*, from the mitochondrial cytochrome b (*cytb*) gene of
110 processed anchovies. Then, a BLAST analysis was performed for molecularly identifying ready-to-
111 eat anchovy's products sold on the Italian market. In addition, the labelling accuracy of the products
112 was assessed. This study was proposed both as a survey for assessing which anchovy species are
113 mainly used for manufacturing commercial products sold within the Italian market and as a useful
114 tool for properly detecting mislabelling accidents involving this seafood category.

115 **2. Materials and Methods**

116 **2.1 Sampling**

117 A total of 111 ready-to-eat products (RTEs) made of anchovies, belonging to three different
118 categories "salted" (n= 30), "marinated" (n=31) and "in oil" (n=50), were sampled in Tuscany
119 (Northern Italy), at different points of sale of large-scale retail distribution (Table 1). A convenience,
120 non-probabilistic sampling was conducted, structured to include a proportional number of products
121 per type, according to the market supply and the brands variety.

122 **2.2 Label analysis**

123 Firstly, as already performed by Velasco, Aldrey, Pérez-Martín, & Sotelo (2016), the label
124 accuracy was evaluated in the light of the mandatory information required by Regulation (EU) No
125 1169/2011. In detail, for each product, the presence of the commercial denomination, the ingredient
126 list, the net and drained weight, the conservation instructions, the best before date, the company name
127 or code and the batch number was assessed. Only for salted products, the presence of mandatory
128 information required by Regulation (EU) No 1379/2013 (the commercial designation of the species
129 and its scientific name, the production method, the area where the product was caught, and the
130 category of used fishing gear and the date of minimum durability) was also verified. For marinated

131 and in oil RTEs voluntary claims on the species scientific name and on the catching area were
132 considered when reported.

133 **2.2 Molecular analysis**

134 **2.3.1 Total DNA extraction and evaluation.** Total DNA extraction was performed starting from
135 ~100 mg of tissue following the protocol described by Armani et al. (2014). The quality and quantity
136 of the DNA from each sample were determined with a NanoDrop ND-1000 spectrophotometer
137 (NanoDrop Technologies, Wilmington, DE, US). One microgram of DNA was electrophoresed on
138 1% agarose gel Gelly-PhorLE (Euroclone, Wetherby, UK), stained with GelRed™Nucleid Acid Gel
139 Stain (Biotium, Hayward, CA, USA), and visualized via ultraviolet transillumination. DNA fragment
140 size was estimated by comparison with the standard marker SharpMass™50-DNA ladder and Sharp-
141 Mass™1-DNA ladder (Euroclone S.p.A-Life Sciences Division, Pavia, Italy). Each DNA sample was
142 stored at -20°C until further analysis.

143 **2.3.2 PCR Primer pair design.** DNA sequences of the complete mitochondrial cytochrome b (*cytb*)
144 gene belonging to the *Engraulis spp.* reported as valid on the official finfishes' database
145 (www.fishbase.org) were retrieved from NIH genetic sequence database GenBank®
146 (<https://www.ncbi.nlm.nih.gov/genbank/>). In detail, at least twenty sequences from the species *E.*
147 *australis*, *E. encrasicolus* and *E. japonicus* and only the ~~unique~~-available sequences from *E. albidus*
148 (n=1), *E. anchoita* (n=2), *E. eurystole* (n=1), *E. mordax* (n=3) and *E. ringens* (n=1) were retrieved.
149 When possible, co-specific sequences belonging to specimens from different geographical areas were
150 retrieved, given the reported intra-specific heterogeneity of *Engraulis spp.* (Jérôme et al., 2008), and
151 especially the considerable number of haplotypes within both *E. encrasicolus* and *E. japonicus*
152 (Magoulas, Castilho, Caetano, Marcato, & Patarnello, 2006; Yu et al., 2005). No sequences were
153 available for *E. capensis* that, although reported as valid species on Fishbase (www.fishbase.org),
154 was instead considered as unaccepted by the World Register of Marine Species
155 (www.marinespecies.org) and classified as a synonym of *E. encrasicolus*. All the retrieved sequences
156 were aligned in with Clustal W in BioEdit version 7.0.9 (Hall, 1999). The primer pair ANCH-531_F

157 (5'-GTTCTTYGCCTTCCACTTCYT-3') and ANCH-1059_R (5'-
158 YACTTGRCCAATAATAATGAATGG-3') was designed to amplify a 484 bp fragment according
159 to the parameters proposed by Giusti et al. (2016), and especially a) considering the level of DNA
160 degradation observed by electrophoresis (section 2.3.1), b) avoiding primer mismatches in critical
161 position, c) taking into consideration the inter-species variability of the selected DNA fragment.

162 *2.3.3 DNA amplification and sequencing.* DNA samples obtained from all RTEs were amplified
163 with the following PCR protocol: 20 µl reaction volume containing 2 µl of a 10X buffer
164 (BiotechRabbit GmbH, Berlin, Germany), 100 mM of each dNTP (Euroclone Spa, Milano), 200 nM
165 of forward primer, 200 nM of reverse primer, 1.0 U PerfectTaq DNA Polymerase (BiotechRabbit
166 GmbH, Berlin, Germany), 100 ng of DNA and DNase free water (Euroclone Spa, Milano). The
167 following cycling program was applied: denaturation at 95 °C for 3 min; 40 cycles at 95 °C for 30 s,
168 55°C for 30 s, and 72°C for 30 s; final extension at 72°C for 7 min. Five microliters of each PCR
169 products were checked by gel electrophoresis on a 2% agarose gel. The amplification of fragments
170 of the expected length was assessed by making a comparison with the standard marker SharpMass™
171 50-DNA ladder (Euroclone Spa, Milano) and the concentration of PCR products by making a
172 comparison with the intensity of the bands of the DNA ladder. A concentration of 10 ng/ml was used
173 as a threshold to destine the samples to the following sequencing phase. All the PCR products were
174 purified with EuroSAP PCR Enzymatic Clean-up kit (EuroClone Spa, Milano) and sequenced by the
175 Experimental Institute of Zooprophyllaxis of Piedmont, Liguria and Aosta Valley (Turin, Italy).

176 The molecular analysis was carried out avoiding contaminations.

177 *2.3.4 BLAST analysis: species identification and mislabelling assessment.* The obtained sequences
178 were analysed using Clustal W in Bio Edit version 7.0.9 (Hall, 1999). Fine adjustments were manually
179 made after visual inspection. All the sequences were used to run a BLAST analysis on GenBank
180 (<https://blast.ncbi.nlm.nih.gov/Blast.cgi>). A top match with a sequence similarity of at least 99%
181 associated to 100% query coverage was used to designate potential species identification. Outcomes
182 from this phase were compared to data obtained from RTEs labelling analysis. The mislabelling rate

183 in relation to the scientific denomination was calculated according to the product category: on (a)
184 salted products declaring the species scientific name (as mandatory information) (b) marinated/in oil
185 products voluntary reporting the species scientific name and (c) marinated/in oil products not
186 reporting the species scientific name but voluntary reporting geographical catching areas that could
187 be unequivocally associated to a unique (or few) species.

188 **3. Results and Discussion**

189 ***3.1 Labelling compliance with EU legislation and voluntary claims assessment***

190 Given the higher brands diversity and the stronger presence at Italian retail level, certainly related
191 to a major consumers' demand, in oil RTEs were the most representative of the sampling (N=50).

192 *3.1.1 Labelling compliances with Regulation (EU) No 1169/2011.* 100% of the collected products
193 were compliant with the disposition provided by ~~the cited r~~Regulation (EU) No 1169/2011, as all the
194 mandatory information were reported on the label and no differences were observed among the
195 distinct categories of products. In all the RTEs, the name of the product was associated to the
196 commercial designation “anchovies” (“acciughe” in Italian), sometimes along with the wording
197 “Mediterranean” (5/111), “Adriatic” (9/111), “Sicilian” (5/111), “Spanish” (5/111), “Cantabrian”
198 (1/111) or “Chilean” (2/111), attributed regardless to the product category (Table 1). Of the whole,
199 the absolute labels compliance with Regulation (EU) No 1169/2011 observed in this study was in
200 accordance with that from Spanish anchovy products analysed by Velasco, Aldrey, Pérez-Martín, &
201 Sotelo (2016), confirming that such law is well obeyed by EU FBOs.

202 *3.1.2 Labelling compliances with Regulation (EU) No 1379/2013.* The label analysis, uniquely
203 performed on the salted RTEs, showed that the mandatory information on the commercial designation
204 and the scientific name of the species, as well as the product's minimum durability, were correctly
205 provided by 100% of the samples. ~~The Article 35 of the considered Regulation especially disposes As~~
206 ~~that~~ the mandatory information on commercial designation and related scientific name should match
207 with those reported on official lists drew up and published by each Member State ~~(Article 35);~~
208 ~~therefore,~~ the term “anchovy” (“acciuga” in Italian) should be uniquely related to the species *E.*

209 *encrasicolus* for products sold on the Italian market according to the list of Ministerial Decree No
210 19105 of September the 22th, 2017. In this respect, as *E. encrasicolus* was the species declared in all
211 the salted RTEs, an actual compliance with the law was observed (Table 1). The catching area was
212 present in all the products, but in 13 out of 30 samples (43.3%) the name in writing of the FAO sub-
213 area or division, mandatorily introduced by ~~the cited r~~Regulation (EU) No 1379/2013, lacked (Table
214 1). Five samples (from RTE-22 to RTE-26) indicated two distinct catching areas (FAO 37 and FAO
215 34) for the same product. Anyway, in both the declared areas, *E. encrasicolus* is the only present
216 species (www.fishbase.org). Finally, for 23.3% of the samples (7/30) no information on both the
217 production method and the fishing gear were provided. Such latter products belonged to the same
218 brands that also reported the incorrect catching area in the label. So, 13 out of 30 salted RTEs (43.3%)
219 was overall not fully compliant with the disposition of Regulation (EU) No 1379/2013.

220 3.1.3 *Voluntary claims*. For the marinated and in oil RTEs, voluntary claims on the species
221 scientific name and/or on the catching area were reported in more than half of the samples (58%).
222 ~~This percentage is associated to the operate of~~ EU FBOs ~~are~~ responding well to the European
223 Parliament Resolution No 2016/2532 that encouraged the Member States, in the context of voluntary
224 labelling, to state all available information that enables the consumer to make an informed choice. In
225 detail, the scientific name was provided in 44.4% (36/81) of the samples, including 66% (33/50) of
226 in oil products and only 9.7% of marinated products (3/31), and the catching area was provided in
227 38.3% of the samples (31/81) including 38% (19/50) of in oil products and 38.7% (12/31) of
228 marinated products. *E. encrasicolus* was declared in 94.4% (34/36) of the product reporting the
229 species name, while only 2 in oil RTEs (RTE-102 and RTE-103) reported the presence of *E. ringens*
230 (Table 1).

231 Of the remained 45 samples not declaring the species, 10 however reported a catching area
232 possibly related to both *E. encrasicolus* and *E. albidus* (FAO area 27 and FAO area 37) while 1
233 sample (RTE-90) reported a catching area clearly related to *E. encrasicolus* (FAO area 27) (Fig. 1).
234 However, it should be noted that the taxonomic status of *E. albidus*, described from the Mediterranean

235 in 2004 (Borsa, Collet, & Durand, 2004) and currently considered as accepted in the World Register
236 of Marine Species (www.marinespecies.org), is questioned by some scientists, specifically suggesting
237 that this may be an eco-morph of *E. encrasicolus* (<http://www.iucnredlist.org/details/18124888/0>).
238 Such samples were anyway considered as composed by *E. encrasicolus*/*E. albidus*, or only *E.*
239 *encrasicolus* in the case of RTE-90.

240 **3.2 Molecular analysis**

241 **3.2.1 Evaluation of total DNA fragmentation and primer design.** The total DNA extracted from
242 the RTEs showed a certain degree of fragmentation, with an electrophoretic pattern hardly visible
243 above 500 bp. Such pattern is typically associated to degraded DNA from processed products packed
244 as cans, tins, jars or tubes having experienced different level of processing (smoking, salting etc.) and
245 also containing multiple additives, preservatives and flavours that may affect the quantity and quality
246 of the DNA (Shokralla, Hellberg, Handy, King, & Hajibabaei, 2015). In this respect, the analysis of
247 processed products has been generally performed by the means of molecular techniques focused on
248 relatively short DNA fragments as genetic marker (Armani et al., 2015a; Meusnier et al., 2008;
249 Shokralla, Hellberg, Handy, King, & Hajibabaei, 2015). Such approach was even proposed for
250 applying next generation meta-barcoding technologies on highly processed complex food matrices
251 (Giusti, Armani, & Sotelo, 2017).

252 For the RTEs analysis, the primer pair ANCH-531_F/ANCH-1059_R was thus designed for
253 amplifying a DNA fragment shorter than 500 bp especially avoiding mismatches within the first three
254 bases near the 3' end since mismatches in this position may affect PCR more dramatically than
255 mismatches located internally or at 5' end (Armani et al., 2016) (Fig. 2). Of course, the high inter-
256 species variability of the molecular marker is a basic prerequisite for ensuring the analysis success.
257 In this respect, ten years ago, Jérôme et al. (2008) already tested the three mitochondrial genes (*16S*,
258 *COI*, and *cytb*) for their suitability to differentiate and authenticate *Engraulidae* species. Even though
259 it was proved that these markers can equally be used, *cytb* appeared slightly more variable between
260 *E. japonicus* and *E. encrasicolus*, strongly distinguished with the maximum bootstrap value of 100%.

261 This is of great importance considering that in a previous study aimed at identifying fish species in
262 ethnic food (Armani et al. 2015b) a low discrimination power of the *COI* gene between *E. japonicus*
263 and *E. encrasicolus* was highlighted. In addition, *Cytb* was proved as suitable marker for
264 distinguishing anchovy species even in the study of Santaclara, Cabado, & Vieites (2006) and, more
265 recently, in the study of Velasco, Aldrey, Pérez-Martín, & Sotelo (2016), even though not all the
266 currently valid eight species belonging to *Engraulis* genus were tested.

267 In this study, the 484 bp *cytb* fragment amplified by our primer pair showed an *Engraulis spp*
268 inter-species variability that was among all the *Engraulis spp.* comparable, and in many cases even
269 higher, to that of the complete mitochondrial *cytb* sequence (Fig. 3). ~~especially regarding for the~~
270 species most used for manufacturing semi-preserved products (*E. encrasicolus*, *E. anchoita*, and *E.*
271 *ringens*). In particular, the fragment shows a higher variability respect to the complete gene also in
272 discriminating the species *E. encrasicolus* and *E. japonicus* that were reported as the most
273 phylogenetically closer by Jerome et al., 2008. In addition, the design of primers pairs specifically
274 intended to some species could even increase the amplification rate in processed products.

275 **3.2.2. Species identity assessment.** All the analysed products were successfully extracted obtaining
276 a total DNA of good quality. All the samples produced at least one amplicon suitable for sequencing
277 and one readable sequence. On the whole, most of the RTEs (102/111, 91.9%) were molecularly
278 identified as *E. encrasicolus*, with ID values of 99% (35.3% of the cases) or 99-100% (64.7% of the
279 cases), regardless the catching area (where it was provided) (Table 1). This difference between ID
280 values may further prove the actual presence of a number of haplotypes within this species already
281 showed by Magoulas, Castilho, Caetano, Marcato, & Patarnello (2006). The remaining 9 RTEs (all
282 in oil products) were instead undoubtedly identified as *E. ringens* with ID value 99-100% (Table 1).

283 **3.2.3 RTEs mislabelling rate.** In this study, criteria for assessing RTEs mislabelling were related
284 to the eventual discrepancy between the species declared on the product label and that actually
285 recovered through the molecular analysis. This eventuality could in fact be related to the voluntary
286 species substitution phenomenon that often occurs where low-value or less-desirable fish species are

Commented [AG1]: Commento di Carmen:

From my point of view you have obtained very homogenous identification results, in our case we had higher number of species coming from S.America. I tend not to use Blast as single identification tool. My suggestion is to compare with results with genetic distance approach

287 swapped for more expensive varieties (FAO, 2018). Mislabelling rate was calculated on 77 RTEs,
288 including all the salted RTEs (N=30), 12 marinated RTEs and 35 in oil RTEs, according to the
289 parameters described in section 2.5. The analysis revealed only one case of mislabelling (mislabelling
290 rate 0.9%), involving 1 in oil product (RTE-91) voluntarily labelled as “*E. encrasicolus*”/FAO 37.2.1,
291 that was instead proved *E. ringens*. This case can be fully considered as an intentional species
292 substitution, as the species *E. ringens* typically inhabits southeast Pacific Ocean (Fig. 1), and it could
293 not have been therefore erroneously by-caught with the con-generic species.

294 It was instead not possible to evaluate the mislabelling rate in the remaining marinated/in oil RTEs.
295 Differently from other categories of preserved products, such as tuna and bonito whose market
296 standards were opportunely fixed (Council Regulation (EEC) No 1536/92), labelling of ready-to-eat
297 anchovies currently presents objective legislative gaps. In fact, no dispositions on which species
298 should be standardly used in marinated or in oil products labelled as “anchovy/anchovies” were laid
299 down at European level yet. Some Member States currently refer to national legislation, such as in
300 the case of Spain, where the semi-preserved products labelled as “achovy” (“anchoa”) must be made
301 only with *E. encrasicolus* (Spanish Royal Decree, 1984). Even the Italian legislation reports a Royal
302 Decree of 1927 stating that all canned products labelled as “anchovy” should be made with *E.*
303 *encrasicolus* (Italian Royal Decree, 1927). However, this old disposition is poorly accurate with
304 regard to the type of products included in the application field and, given the market developments
305 and growth since then, it may be actually considered as anachronistic. Except for that decree, any
306 related disposition has been provided. Not even the international Codex standards may be adopted,
307 as only referring to all commercial species of fish belonging to the family Engraulidae that have been
308 salted, boiled and dried and not covering products which have undergone an enzymatic maturation in
309 brine (CODEX STAN 236-2003). Actually, some information can be uniquely extrapolated from EU
310 codes from combined nomenclature, where the definition “prepared and preserved anchovy” covers
311 all the *Engraulis spp.* and even other anchovy species (*Anchovia macrolepidota*, *Lycengraulis*
312 *grossidena*) (EUMOFA, 2018). However, as already mentioned, this condition is not supported by a

313 concrete legislation system. This limit unavoidably leads to concerns such as the effective higher
314 possibility that cases of replacement with low-value species occur, as well as the concrete
315 impossibility for consumers to make informed choices on the products they purchase.

316 **3.3 Anchovy products in the context of the EU market: local species exploitation and** 317 **entrance of non-indigenous species**

318 *E. encrasicolus*, or European anchovy is a pelagic species distributed in the Eastern North and
319 Central Atlantic, Mediterranean, Black and Azov seas, as well as the Coast of West Africa to Angola
320 (EUMOFA, 2018) (Fig. 1). Given its abundance in such areas, it is the mainly caught, processed and
321 consumed species in EU (data also confirmed by the outcomes from this study). A report by the EU
322 Market Observatory for Fisheries and Aquaculture products (EUMOFA), specifically dedicated to
323 the anchovy market and providing data of the year 2015, states that 38% of anchovies consumed in
324 form of processed ready-to-eat products in Italy (38% of the total anchovy consumption) belong to
325 small processing industry relying on locally caught species flanked by large scale production
326 (EUMOFA, 2018). In both cases, *E. encrasicolus*, as inhabiting European waters, is mainly used.
327 However, large scale production even relies on extra-EU imports, and involving other *Engraulis spp.*
328 such as anchoveta (*E. ringens*), Argentine anchovy (*E. anchoita*) and Japanese anchovy (*E.*
329 *japonicus*). *E. ringens*, the other species found in the products analysed in this study is a pelagic fish
330 in the south-eastern Pacific Ocean, regularly caught by purse seiner vessels all along the Peruvian
331 coast including the north of Chile (Fig. 1). It represents the most caught fish in the world history in
332 terms of volume (FAO, 2016) and the Peruvian northern-central stock currently supports the single
333 most important mono-specific fishery in the world (cedepesca.net). It should be however noted that
334 the vast majority of the catch does not go for human consumption but is reduced to fishmeal and fish
335 oil and exported, primarily for aquaculture and animal feed (FAO, 2016). However, efforts in
336 encouraging the local anchovy value by addressing the Peruvian processing industry to manufacture
337 products for human consumption have been made in the last years. For instance, in July 2012,
338 “Compañía Americana de Conservas”, the major Peruvian society of fishery industry, signed a

339 framework collaboration agreement with CeDePesca for the implementation of a fishery
340 improvement project targeting the portion of the fishery fishing for direct human consumption
341 (<http://cedepesca.net/promes/small-pelagics/peruvian-anchovy-direct-human-consumption/>). Other
342 examples of private and public initiatives were reported in a recent report by *The Marine Ingredients*
343 *Organization* (IFFO, 2017). However, data from that report showed a still low amount of exported
344 product volume. This aspect may explain the low presence of this products within EU market and
345 therefore the low percentage of *E. ringens* RTEs respect to *E. encrasicolus* RTEs in this study.

346 **Conclusions**

347 This study confirms how the use of simple and cost-effective DNA-based analytical techniques,
348 allow to properly identify fish species and thus support traceability in the seafood supply chain. It
349 also furnishes encouraging results showing a very low mislabelling rate of ready-to-eat anchovy
350 products. This is particularly interesting considering these products are characterized by a high price
351 on the Italian market and could thus represent an optimum target to perpetrate voluntary frauds.
352 Overall, this survey confirms previous studies reporting a reduction of misdescription incidents in the
353 EU highlighting how the market can positively respond to policies intended to regulate the seafood
354 sector.

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462

463 **Figure captions**

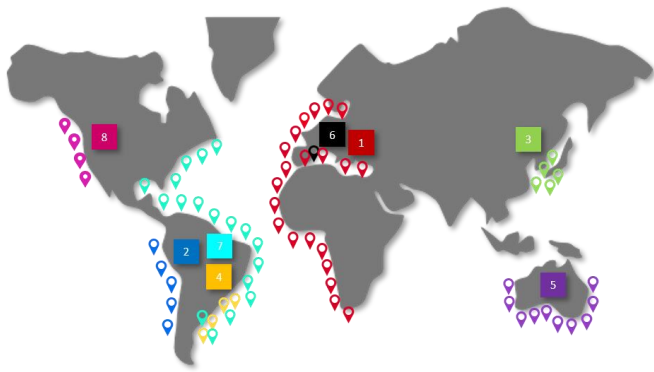
464 **Fig. 1:** Native distribution map for *Engraulis* species (source: www.fishbase.org). 1: *E.*
 465 *encrasicolus*; (European anchovy); 2: *E. ringens* (Anchoveta); 3: *E. japonicus* (Japanese
 466 anchovy); 4: *E. anchoita* (Argentine anchovy); 5: *E. australis* (Australian anchovy); 6: *E.*
 467 *albidus* (White anchovy); 7: *E. eurystole* (Silver anchovy); 8: *E. mordax* (Californian
 468 anchovy).

469 **Fig. 2:** Alignment between the primer pair ANCH-531_F/ANCH-1059_R projected in this study
 470 and the cytb sequences retrieved from GenBank; mismatches were highlighted in grey.

471 **Fig. 3:** Inter-species variability analyses conducted in MEGA 6 (Tamura, Stecher, Peterson, Filipiński,
 472 & Kumar, 2013) using the Kimura 2-parameter model (Kimura, 1980) on **a**) 484 bp fragment
 473 of this study and **b**) complete cytb gene. The analysis involved the following nucleotide
 474 sequences randomly selected among those used for primer projecting: *E. albidus* (MG958167);
 475 *E. anchoita* (JQ012416); *E. australis* (KJ007734); *E. encrasicolus* (JQ716614); *E. eurystole*
 476 (JQ012427); *E. japonicus* (KJ007662); *E. mordax* (JQ012421); *E. ringens* (JQ012426).

477

478 **Fig. 1:**



479

480 **Fig. 2:**

ANCH-531_F	ANCH-1059_R (reverse complement)
GTTCPTYGCCTTCCACTTCYT	CCATTCATTATTATGGYCAAGTR
***** <i>E. encrasicolus</i>	***** <i>E. encrasicolus</i>
***** <i>E. japonicus</i>	***** <i>E. japonicus</i>
***** <i>E. albidus</i>	***** <i>E. albidus</i>
ⓐ***** <i>E. anchoita</i>	**ⓐ***** <i>E. anchoita</i>
***** <i>E. australis</i>	***** <i>E. australis</i>
ⓐ***** <i>E. mordax</i>	**ⓐ***ⓐ*ⓐ***** <i>E. mordax</i>
ⓐ***** <i>E. ringens</i>	**ⓐ***** <i>E. ringens</i>

481

482 **Fig. 3**

a)

	1	2	3	4	5	6	7	8
1. <i>E. albidus</i>								
2. <i>E. anchoita</i>	0.22063							
3. <i>E. australis</i>	0.05440	0.24473						
4. <i>E. encrasicolus</i>	0.03193	0.23852	0.05659					
5. <i>E. eurystole</i>	0.01255	0.22063	0.06369	0.04080				
6. <i>E. japonicus</i>	0.05915	0.24510	0.02755	0.05211	0.06382			
7. <i>E. mordax</i>	0.20146	0.24433	0.19542	0.20701	0.20146	0.19856		
8. <i>E. ringens</i>	0.23271	0.12723	0.22935	0.22034	0.23271	0.22063	0.24433	

b)

	1	2	3	4	5	6	7	8
1. <i>E. albidus</i>								
2. <i>E. anchoita</i>	0.22272							
3. <i>E. australis</i>	0.03665	0.23407						
4. <i>E. encrasicolus</i>	0.02646	0.22679	0.03971					
5. <i>E. eurystole</i>	0.01064	0.22132	0.03975	0.02746				
6. <i>E. japonicus</i>	0.04290	0.23713	0.02249	0.04182	0.04394			
7. <i>E. mordax</i>	0.18741	0.21489	0.18609	0.18465	0.18085	0.18359		
8. <i>E. ringens</i>	0.21561	0.09386	0.21149	0.20862	0.21423	0.21026	0.21752	

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