
Condition states in anchovy (*Engraulis encrasicolus*) and sardine (*Sardina pilchardus*) revealed by energy and proximate composition relationships

Favreau Aurélien ^{1,*}, Doray Mathieu ², Spitz Jérôme ^{3,4}, Le Mestre Sophie ¹, Huret Martin ¹

¹ DECOD, L'Institut Agro, IFREMER, INRAE, Plouzané, France

² DECOD, L'Institut Agro, IFREMER, INRAE, 44000 Nantes, France

³ Observatoire Pelagis, UAR 3462, La Rochelle Université/CNRS, La Rochelle, France

⁴ CEBC, UMR 7273, La Rochelle Université/ CNRS, La Rochelle, France

* Corresponding author : Aurélien Favreau, email address : aurelien.favreau@ifremer.fr

Abstract :

Energy content has long been proposed as a fundamental, integrated, and reliable indicator of the condition of individuals as it reflects past bioenergetics and influences future life-history traits. There is a direct biochemical link between energy density and body composition described by four main compounds in fish (protein, lipid, ash, and water), with proteins and lipids being the sources of energy. If relationships between water content, or lipid content, and energy density have been well described in relative terms, the absolute mass variations in the proximate composition have been overlooked and thus their interpretation is often equivocal. In our study, based on a large and unique dataset on the proximate composition and energy density of anchovy (*Engraulis encrasicolus*) and sardine (*Sardina pilchardus*) from sampling in the Bay of Biscay and the English Channel, we aimed to better explain the patterns between water content and other proximate components or energy density, based on the dynamics of proteins, lipids, and water absolute masses. For the first time, we defined good, intermediate, and poor condition states in wild fish, based on water content, corresponding to the different dynamics of lipids and proteins in the metabolism of individuals. Anchovy and sardine exhibited remarkably similar patterns of variation in the compounds and in the limits between the condition states with respect to water content. Those patterns revealed that water mass remained constant for a given fish size whatever its condition state, and that variability in water content only resulted from the variation in lipid and protein masses. Furthermore, the differential dynamics of proteins and lipids, with only lipids mobilized in the good condition state, only proteins in the poor condition state, and both proteins and lipids in the intermediate condition state, elucidates the nonlinear pattern observed in the relationship between energy density and water content. Overall, our results highlight the importance of monitoring the intraspecific variations in water content to predict the proximate composition and energy content in small pelagic fish and better assess individual and population conditions in changing ecosystems as well as to better parameterize bioenergetic models.

Keywords : Bay of Biscay, condition, energy, lipid, protein, small pelagic fish, water content

42 **1. Introduction**

43 Energy content has long been proposed as a fundamental, integrated and reliable indicator of the
44 condition of individuals (Molnár *et al.*, 2009), populations (Pagano *et al.*, 2018), and ecosystems
45 (Benoit-Bird, 2004; Spitz and Jouma'a, 2013; Pothoven and Fahnenstiel, 2014). Energy content is
46 key to the understanding of the observed variability in animal life history traits such as growth,
47 reproduction and mortality and is biochemically linked to body composition. As such, body
48 composition has been studied for a long time in both farmed and wild mammals (Moulton, 1923;
49 Noren *et al.*, 2003), birds (Eits *et al.*, 2002; Rivera-Torres *et al.*, 2011; Guglielmo, 2018) and fish
50 (Groves, 1970), for its large significance in various aspects of individual physiology, biology and
51 ecology. Body composition mostly varies in response to feeding (Gerking, 1955; Bonvini *et al.*,
52 2015) and ontogeny (Wuenschel *et al.*, 2006; Bakhtiarvandi *et al.*, 2014). It can give insights into
53 the well-being or condition of the animal through the amount of lipid or protein reserves (McCue,
54 2010).

55 Lipids are the most variable component within an animal's body, serving as the primary energy
56 reserve to be catabolised (McCue, 2010). Their quantity is regarded as a good indicator of an
57 organism's condition (Lloret, 2014) and can vary significantly, not only between individuals, but
58 also throughout an individual life cycle, due to both internal factors, reproductive status and
59 ontogeny, and environmental variations, ranging from 2% to 20% in fish for example (Lloret,
60 2014). Proteins usually represent the dominant component of animals weight, after water, with their
61 quantity being relatively stable (approximately 20% of wet mass in fish, Lloret, 2014), although
62 they can be catabolised to provide energy when lipids are depleted (Savitz, 1971; Black and Love,
63 1986). While both proteins and lipids serve as sources of energy, proteins primarily contribute to
64 structural components, whereas lipids represent mobilisable energy reserves. Ash is a highly stable
65 component, as it constitutes inorganic molecules (scales, skeleton) and is strongly correlated with
66 the organism's length (McComish et al., 1974). In fish studies, four components are typically
67 considered: water, ash, protein and lipid. Carbohydrates are generally not considered, as they
68 constitute a very small proportion of wet mass (<0.14%; Brett and Groves, 1979). As water and ash
69 are not energetic, energy content is intrinsically linked to protein and lipids content and can then be
70 assessed using the energy density of these components (Breck, 2008, 2014; Schloesser and Fabrizio,
71 2015).

72 In nature, all animals can face periods of food deprivation, and some, such as fish, can cope with
73 long periods of starvation (McCue, 2010). During these periods, proximate composition varies,
74 eventually reaching energy depletion. For an animal with plenty of reserves, food deprivation is
75 usually followed by three phases for birds (Le Maho *et al.*, 1981; Robin *et al.*, 1988; Groscolas and
76 Robin, 2001), mammals (Cherel *et al.*, 1988, 1991, 1992; Caloin, 2004) and fish (Bar, 2014). Phase
77 1 is short and characterised by the mobilisation of protein and/or lipids to produce energy, phase 2
78 lasts longer and involves lipids catabolisation, until phase 3 where lipids are depleted and proteins
79 become the main source of energy (Bar, 2014). Studying the effect of starvation on animals has
80 been mainly assessed in controlled environments, following individuals in longitudinal experiments.

81 Identifying condition states and understanding the respective contributions of lipids and proteins
82 when reserves are catabolised in wild organisms remains challenging. The scarcity of extensive in-
83 situ datasets documenting fish proximate composition and energy density, over a wide range of fish
84 condition, has limited our understanding of the variations in wild fish condition.

85 Water content (or dry mass content), which shows a strong relationship with energy density ($R^2 >$
86 0.90) for many taxa (Ciancio et al., 2007), provides a quick and reliable method for assessing
87 energy density. This relationship is mainly related to the catabolism of lipids, which once mobilised
88 lead to a relative increase of water content and a decrease in energy density. For fish, numerous
89 studies have explored the correlation between water content (or dry mass content) and energy
90 density (Hartman and Brandt, 1995; Tirelli et al., 2006; Dubreuil and Petitgas, 2009; Gatti et al.,
91 2018). However, Hartman and Brandt, (1995) and Gatti *et al* (2018) have shown that this
92 relationship was nonlinear. This non linearity was discussed with respect to size and ontogeny
93 effects (Wuenschel et al., 2006). Although energy density and proximate composition have been
94 extensively studied in fish, their interdependent variations still remain misunderstood. We here
95 suggest that the usual representation of proximate composition, water content and energy density in
96 relative terms, has maintained confusion in the interpretation of the absolute variation of these
97 variables along a gradient of fish condition. Specifically, a change in the relative value of one
98 component will inevitably affect the relative value of another component, even if the absolute term
99 remains constant. Unraveling the patterns in their absolute variations would therefore benefit the
100 community.

101 Small pelagic fish (SPF) are high-fat, high-energy forage species, that play a key role in pelagic
102 ecosystems, transferring energy from plankton to higher trophic levels (Cury, 2000). In temperate
103 shelf ecosystems, such as the Bay of Biscay or the English channel, SPF energy density displays
104 large variations, with low energy density in winter and spring after overwintering and high energy
105 density in summer and autumn during and after the feeding season (Spitz and Jouma'a, 2013; Gatti
106 *et al.*, 2018). Because farming small pelagic fish remains challenging (but see Bandarra et al, 2018

107 Queiros *et al.*, 2019, 2021), no SPF proximate composition has been studied in vivo. These fish
108 species are hence absent from existing reviews of experimental studies exploring strategies for
109 dealing with starvation and biochemical processes at stake (McCue, 2010; Bar, 2014).

110 Our study was based on a large and unique dataset of proximate composition and energy density of
111 European anchovy *Engraulis encrasicolus* (Linnaeus 1758), hereafter anchovy, and European
112 sardine *Sardina pilchardus* (Walbaum 1792), hereafter sardine, sampled in the Bay of Biscay and
113 the English Channel. We aimed to provide new insights on the link between proximate composition,
114 state in fish condition and associated energy density. First, we characterised the patterns of variation
115 of protein and lipid contents over the observed range of water content, to derive distinct condition
116 states. Second, we tested the effect of internal (reproductive status and length) and external (season
117 and region) factors on the individuals condition state. Third, we checked if the water mass in
118 absolute terms was varying across the different condition states in response to the variations of
119 proximate composition. Fourth, we assessed the respective contributions of proteins and lipids to
120 energy supply in each condition state, when energy has to be mobilised. Finally, we further
121 investigated the non linearity in the well-known relationship between energy density and water
122 content, in light of lipid and protein content variations.

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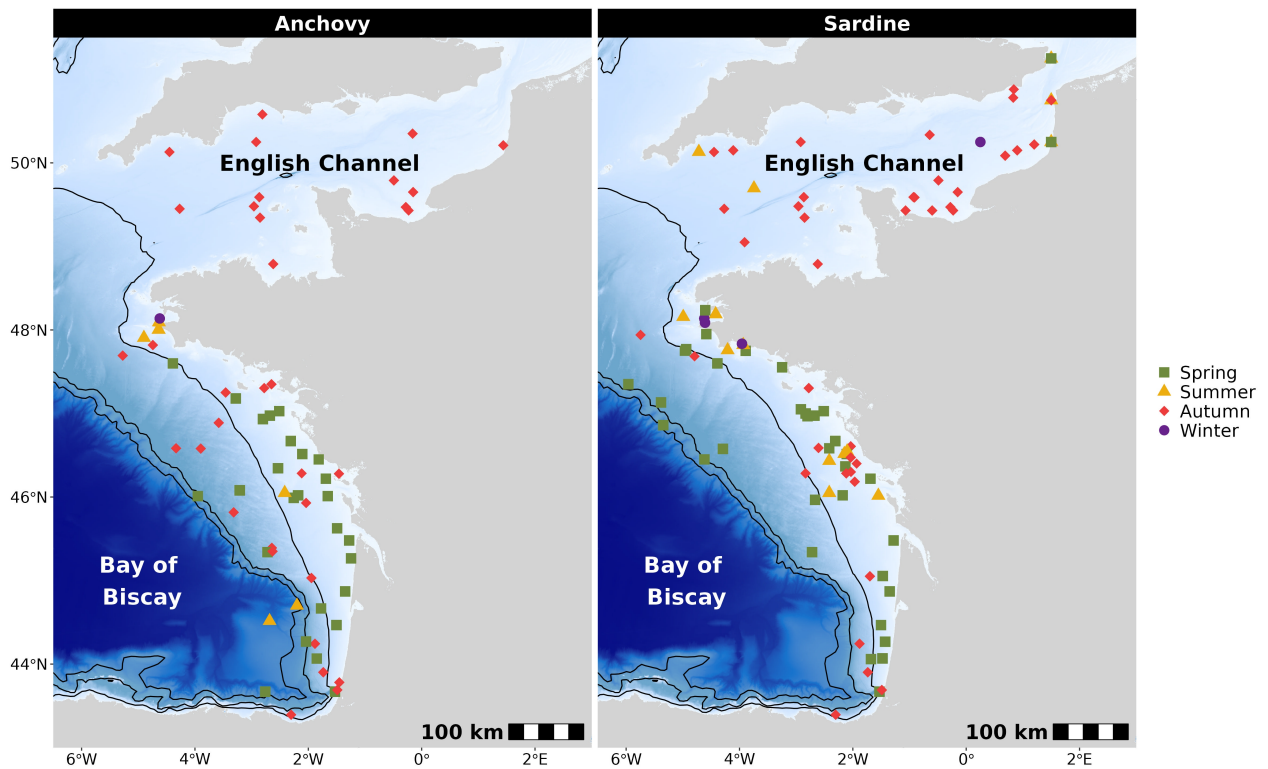
124 **2. Material and methods**

125 *2.1. Study area and sampling*

126 Juvenile and adult anchovy and sardine were collected from 2014 to 2017 in the Bay of Biscay and
127 the English Channel (Figure 1). All the data used in this study were made available in Huret *et al.*
128 (2024). Fish were sampled during the PELGAS (Doray *et al.*, 2000; 2018b), EVHOE (Laffargue *et*
129 *al.*, 1987), CGFS (Giraldo *et al.*, 1988) and CAMANOC (Travers-Trolet and Verin, 2014) scientific
130 surveys conducted by IFREMER on the RV “Thalassa”, during the JUVENA survey conducted by
131 AZTI (Boyra *et al.*, 2003), and from commercial landings within the European Data Collection

132 Framework (DCF). The PELGAS integrated pelagic survey monitors small pelagic fish and their
133 ecosystem in spring, while EVHOE bottom trawl survey evaluates demersal fish communities in
134 autumn, both in the Bay of Biscay. The CGFS and CAMANOC bottom trawl surveys evaluate
135 demersal fish communities and their environment in the English Channel in autumn. Despite their
136 focus on demersal fish communities, EVHOE, CGFS and CAMANOC surveys catch small pelagic
137 fish in sufficient numbers to get samples representative of the fish condition in autumn (Gatti et al,
138 2018). JUVENA survey monitors the abundance of juvenile anchovy in early autumn in the Bay of
139 Biscay. Samples from commercial landings were obtained as part of the national CAPTAIN
140 research project.

141 A total of 503 anchovies and 976 sardines sampled between 2014 and 2017 were used in this study
142 (Table 1). Sampling locations were representative of the species core distribution areas in the Bay
143 of Biscay and the English Channel (Figure 1). Each fish was individually measured to the nearest
144 tenth of a centimeter and weighed to the nearest tenth of a gram. Maturity stages were determined
145 following ICES guidelines (ICES, 2008) based on macroscopic gonads observation and using a six-
146 stage key as follows: stages 1 & 2 indicate immature and developing individuals, stages 3–5
147 indicate three steps of increasing gonad development and the spawning period (stage 3: partial pre-
148 spawning; stage 4: spawning (hydrated); stage 5: partial post-spawning), and stage 6 features the
149 final post-spawning period. Fishes characterised by maturity stages 3, 4 or 5 were considered as
150 being in an active reproductive period as opposed to fishes in stages 1, 2 or 6. These measurements
151 were taken either at sea or later in the laboratory. A wide range of body lengths (4.7 – 19.1 cm for
152 anchovy and 8 – 26.8 cm for sardine) and masses (0.5 – 50.2 g for anchovy and 3.4 – 162.2 g for
153 sardine) was available, representing the size spectrum of the two species across juvenile and adult
154 stages. The collected fish were frozen individually at -20°C before laboratory processing.



155

156 *Figure 1: Sampling locations and associated seasons, for anchovy (left) and sardine (right) from the*
 157 *six data sources considered: PELGAS (spring in the Bay of Biscay), JUVENA and EVHOE (autumn*
 158 *in the Bay of Biscay) and CGFS/CAMANOC (autumn in the English Channel) surveys, as well as*
 159 *commercial landings (CAPTAIN project, several seasons and both regions). Black lines represent*
 160 *isobaths of 100, 200 and 1000 meters.*

161

162 2.2. Water content

163 In the laboratory, fish were unfrozen, ground and freeze-dried for at least 48 hours. Water content
 164 of the entire fish was determined from dry mass (M_D) and wet mass (M_W) ratio. Then, fish were
 165 ground again to obtain fine homogeneous dry powder for subsequent analysis. In this study, the
 166 term “content” refers to the relative proportion of the component on the total fish mass. It will be
 167 used for ash, lipid, protein and water components.

168 2.3. Proximate composition analysis

169 Ash content was determined gravimetrically by combusting dried tissue in a muffle furnace at
170 550°C for six hours. Lipids and proteins were analysed by a certified laboratory (Laboccea,
171 Plouzané, France). Protein content was estimated using the Kjeldahl method (Sáez-Plaza *et al.*,
172 2013). It consists in first determining the quantity of nitrogen contained in the sample, and to
173 convert it into protein content using a conversion factor (6.25 here, Sáez-Plaza *et al.*, 2013). Lipid
174 content was determined by hydrolysis, using petroleum ether as an organic solvent. Carbohydrates
175 represent less than 1% of fish mass and were thus neglected as in previous studies (Brett and
176 Groves, 1979; Craig, 1977; Craig et al, 1978). Protein, lipid and ash content did not exactly sum to
177 1 in dry weight (DW, anchovy: mean = 0.91, sd = 0.04; sardine: mean = 0.90, sd = 0.04). This
178 discrepancy may arise from residual water, measurement uncertainties, or to a lesser extent the
179 exclusion of carbohydrates. Body component contents have been normalised by dividing each
180 component by the sum of lipids, proteins and ash content, to sum to one, enabling comparisons
181 between fishes, assuming proportional errors across the components (Breck, 2014). Values of water,
182 protein, lipids and ash masses were calculated by multiplying the fish body mass by the normalised
183 percentages of fish components. A total of 104 and 116 proximate composition analyses were
184 performed for anchovy and sardine, respectively (Table 1).

185 2.4. Energy density

186 Energy density measurements were performed following the protocols of Dubreuil and Petitgas,
187 (2009) and Spitz and Jouma'a, (2013). Two subsamples of fish powder were placed in an adiabatic
188 bomb calorimeter (IKA C-4000 adiabatic bomb calorimeter, IKA-WerkeGmbH & co. KG) for
189 energy measurements. The energy density (ED, kJ.g⁻¹ dry mass) was determined by measuring the
190 heat released through the combustion of a small subsample, approximately 200 mg. If the
191 coefficient of variation between the two measurements exceeded 3%, a third measurement was
192 made. Finally, ED subsamples measurements were averaged and assigned to each individual fish.

193 Energy density analyses were conducted on 503 individuals for anchovy and 976 individuals for
 194 sardine (Table 1).

195 *Table 1: Sampling design: data sources, region, season, number of samples per species for energy*
 196 *density and proximate composition analysis.*

Data sources	Region	Season	Energy density		Proximate composition	
			Anchovy	Sardine	Anchovy	Sardine
PELGAS	Bay of Biscay	Spring (May)	186	204	17	17
EVHOE	Bay of Biscay	Autumn (Oct-Nov)	130	91	31	12
CGFS/ CAMANOC	English Channel	Autumn (Sept-Oct)	91	164	24	25
JUVENA	Bay of Biscay	Autumn (Sept)	22	0	0	0
CAPTAIN	Bay of Biscay, English Channel	All	74	517	32	62
Total			503	976	104	116

197

198 2.5. Statistical analysis

199 The contents of proximate components, namely lipids, proteins and ash, were used as response
 200 variables and water content as the explanatory variable (Breck, 2008, 2014) in segmented linear
 201 models (Muggeo, 2008). Segmented linear models (SLM) are linear regression models where the
 202 relationships between the response and one or more explanatory variables are piecewise linear,
 203 represented by two or more straight lines connected at unknown values, referred as breakpoints.
 204 SLM were used to identify breakpoints delimiting intervals in the relationship between proximate
 205 components and water content. These intervals were assumed to represent fish condition states,
 206 defined by thresholds in water content.

207 Multinomial logistic regression (Starkweather and Moske, 2011) analysis were used to examine the
 208 influence of both internal (length and reproductive status) and external factors (season and region)
 209 on the condition states, the response variable comprising three categories: poor, intermediate, and

210 good. This regression method extends the binomial logistic regression to situations where the
211 dependent variable is nominal with more than two categories. The intermediate state was chosen as
212 the reference category, representing the state with the most available data, while good and poor
213 states were the alternative categories. The optimal model for each species was determined through
214 stepwise analysis based on the Akaike Information Criterion (AIC). Odds ratios (ORs) with 95%
215 confidence intervals (CI) and associated p-values were used to display the results. The OR
216 represents the likelihood of the independent variable to belong to a specific category relative to the
217 reference category. The OR is therefore greater than 1 if the individual has a higher probability to
218 be in the considered category than in the reference category, and less than 1 if it has a lower
219 probability to be in the considered category.

220 Breck (2008) showed that proteins are closely linked to water content. The loss of protein during
221 starvation was shown to be accompanied by water losses. Thus, we investigated the effect of the
222 content in proximate components other than water on the water mass, independently of fish length,
223 using the following multiple linear regression model:

$$224 \quad W \sim L + p + l + a + \epsilon (1)$$

225 With W the mass of water in gram, L the fish length in cm, p , l and a the protein, lipids and ash
226 contents, respectively and ϵ the residuals term.

227 Log-log linear relationships were fitted with proximate components masses as response variables,
228 and water mass as explanatory variable, for each condition state determined from the segmented
229 regression, for anchovy and sardine. Additionally, a covariance analysis (ANCOVA) was applied to
230 assess if the relationships between water mass and proximate component mass differed among
231 condition states.

232 The respective contributions and their associated uncertainties of lipids and proteins to energy
233 mobilisation were calculated using the variation of each component (i.e. regression slopes) within
234 the different intervals (i.e. condition state). Considering the higher energy content of lipids

235 compared to proteins, we assessed energy contributions in terms of both mass and energy. Usually,
236 energy density of lipids ranges from 34.7 to 39.5 kJ.g⁻¹ (Schloesser and Fabrizio, 2015), and energy
237 density of proteins is taken as 23.6 kJ.g⁻¹ (Paine, 1971; Beamish et al, 1975; Brett and Groves,
238 1979). Consequently, we used the value of 23.6 kJ.g⁻¹ for protein and 38.5 kJ.g⁻¹ for lipids, based on
239 the measurement of mesenteric fat of sardine using bomb calorimetry (n=4, CV = 1.2%) which fits
240 in the range of the literature. Calculations of the respective contributions based on the regressions
241 are detailed in appendix A.

242 Finally, the relationship between energy density and water content was further investigated. The
243 former was used as the response variable and the latter as the explanatory variable using two linear
244 regression models. The standard one was fitted to all individual data together. An alternative model
245 was a segmented linear model between water content and energy fitted over each condition state.

246 Linear models goodness of fit was assessed using classical diagnosis tools (coefficients of
247 determination (R²), residuals Q-Q plots, residuals versus fitted values plots, standard error and
248 confidence interval at 95% of regression coefficients). Statistical tests were performed at a
249 significance level of 0.05. Analyses were performed using the statistical R software (R Core Team,
250 2023). Segmented regression analysis was performed using the package *segmented* (Muggeo,
251 2008). Multinomial logistic regression models were performed using the function *multinom* in
252 package *nnet*.

253 2.6. *Ethical statement*

254 Ethical review and approval were not required for this study because it did not involve animal
255 experimentation or harm. No animals were involved in the conduct of this study as the analysis was
256 done on previously collected data on wild-caught fish that were sampled within the European Data
257 Collection Framework.

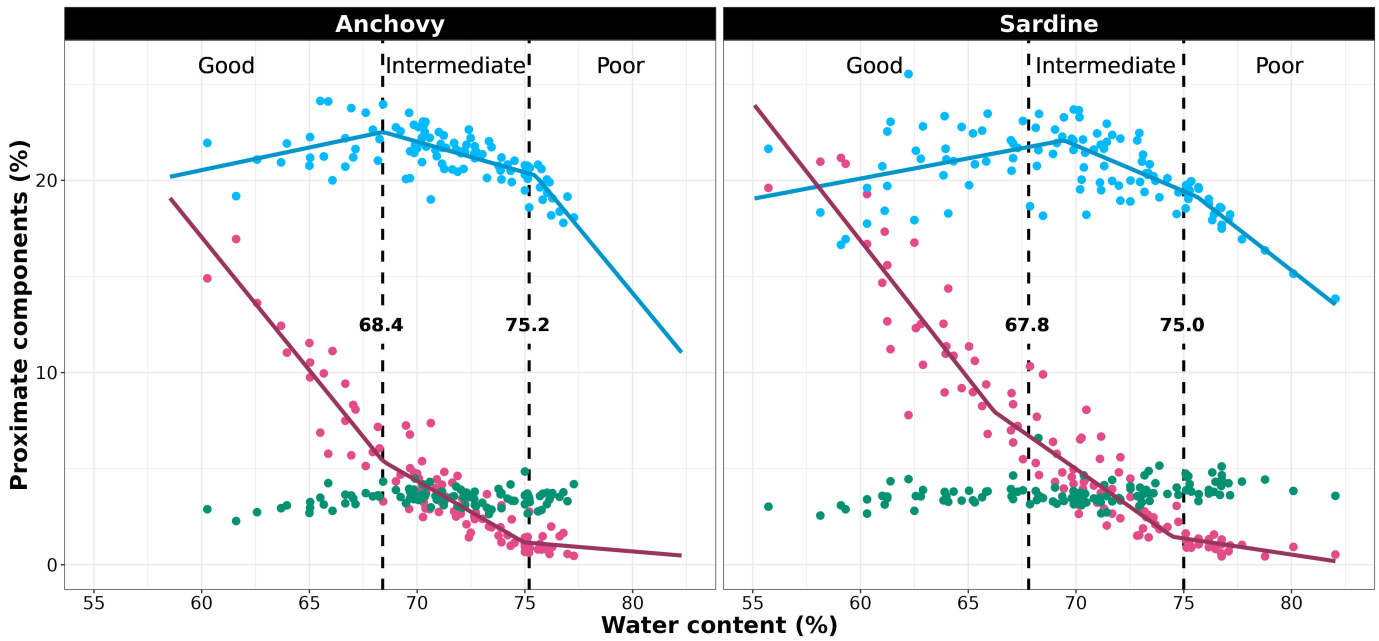
258 **3. Results**

259 *3.1. Fish condition states from proximate composition*

260 Strong relationships were found between contents of water and proximate components, namely
261 proteins and lipids (Figure 2). The segmented regression analysis identified two breakpoints,
262 characterising a change in the regression slopes between water content and contents in proximate
263 components for both anchovy and sardine (Figure 2, Table 2 and Table A1). Variation in water
264 content explained a great proportion of lipid content variation in the segmented regression (R^2 equal
265 to 0.90 and 0.91 for anchovy and sardine, respectively). Water content variation explained a lower
266 proportion of protein variation (R^2 equal to 0.52 and 0.50 for anchovy and sardine, respectively).
267 Six among eight breakpoints were significant. Only the second lipid breakpoint of anchovy and the
268 second protein breakpoint for sardine had p-values higher than 0.05 (0.32 and 0.12, respectively,
269 Table 2). Despite this non significance, these breakpoints have been considered in the rest of the
270 results as their complementary breakpoints (2nd protein breakpoint of anchovy and 2nd lipid
271 breakpoints for sardine), as well as their counterparts in the other species, are significant, which
272 suggests two changes in the pattern of proximate composition against water content that appeared
273 interesting to further investigate in our study.

274 For anchovy and sardine, the first lipid breakpoint occurred respectively at 68.4% and 66.2% of
275 water content, and the second at 75% and 74.5%. Similarly, the first protein breakpoint occurred
276 respectively at 68.4% and 69.4% of water content, and the second at 75.5% and 75.6%. For each
277 species, the two protein and lipid breakpoints were close, the maximum difference being 3.2% of
278 water content between the first breakpoints for sardine. Considering these similarities, and in order
279 to isolate three distinct composition intervals for each species, we used the average breakpoints
280 calculated over the corresponding protein and lipid breakpoints as common breakpoints, for each
281 species (Table 2). Based on these average breakpoints we identified three condition states, which
282 exhibited remarkable similarities between the two species (Figure A1, Table 2). In the first state,

283 fish water content was lower than 68.4% and 67.8% for anchovy and sardine, respectively, and lipid
284 content was higher than 5% in wet mass for both species. Here, we assumed that the higher the lipid
285 content, the better the condition state. Within this 'good condition' state, lipid content strongly
286 decreased as water content increased, while protein content slightly increased. The second state
287 extended from 68.4% to 75.2% and from 67.8% to 75% in water content for anchovy and sardine,
288 respectively. Both lipid and protein contents decreased with increasing water content, albeit at
289 different rates in this 'intermediate condition' state. In the third state, fish water contents were
290 above 75.2% and 75% for anchovy and sardine, respectively. Lipid content reached a minimal value
291 "threshold" of ca. 1% suggesting exhaustion of lipid reserves, while protein levels significantly
292 decreased. Fish were considered in a 'poor condition' state.



293 Figure 2: Observed contents of proteins (blue dots), lipids (purple dots) and ash (green
 294 dots) in anchovy and sardine body composition, as a function of water content. Solid
 295 lines: segmented regression models for proteins (blue lines), and lipids (purple lines).
 296 The dotted vertical lines represent the mean values of the breakpoints identified from
 297 proteins and lipids segmented regressions.

298
 299 Table 2: Breakpoint values for lipids and proteins, defining condition states in anchovy
 300 and sardine. Slope coefficient, associated standard error and confidence intervals for
 301 each segment are presented in Table A1

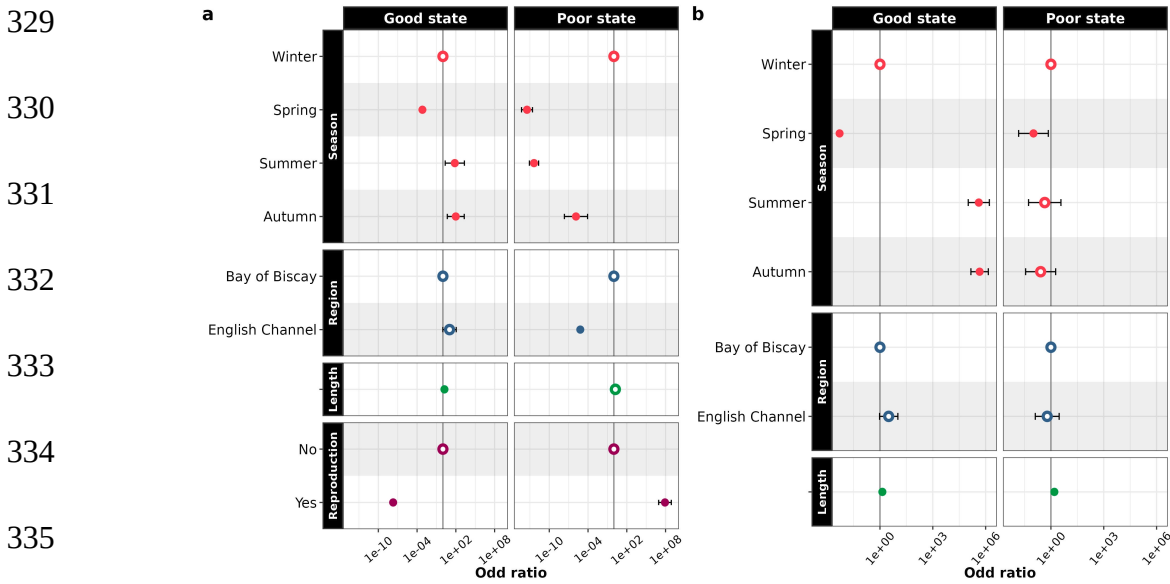
	Anchovy						Sardine							
	Protein		Lipid		Mean	Protein		Lipid		Mean				
	Value	P-value	R ²	Value	P-value	R ²	Value	Value	P-value	R ²	Value	P-value	R ²	Value
Breakpoint 1	0.684	3.3.10 ⁻⁵	0.52	0.684	1.6.10 ⁻⁴	0.90	0.684	0.694	6.7.10 ⁻⁵	0.50	0.662	3.4.10 ⁻³	0.91	0.678
Breakpoint 2	0.755	2.3.10 ⁻²	0.52	0.750	3.2.10 ⁻¹	0.90	0.752	0.756	1.2.10 ⁻¹	0.50	0.745	2.6.10 ⁻³	0.91	0.750

303 3.2. *Effect of length, season, reproductive status and region on the condition state*

304 Through multinomial logistic regression analyses, we explored the impact of season, region, length
305 and reproductive status on the condition states of anchovy and sardine, with the intermediate
306 condition state serving as the reference category (Figure 3, Table A2, Table A3). For anchovy, the
307 optimal model, as determined by the AIC, included all variables (season, reproductive status,
308 region, and length). In contrast, for sardine, the reproductive status variable was excluded from the
309 best model, while season, region, and length variables were retained.

310 For anchovy, seasonal variations significantly influenced its condition state (Figure 3). During
311 summer [OR = 6.91.10¹, 95% CI = 2.27 - 2.10.10³] and autumn, they were more likely to be in good
312 condition than in winter, whereas in spring, the likelihood decreased. Conversely, for all seasons,
313 the likelihood of being in a poor condition was lower than in winter. Similar patterns are observed
314 for sardine, despite non significant ORs for summer and autumn, for poor condition state.
315 Regarding region, both species exhibited comparable patterns although significance was not always
316 verified. Anchovy and sardine from the English Channel were more likely to be in good condition
317 (p-value = 0.052 for anchovy and 0.061 for sardine), and less likely to be in poor condition
318 compared to those from the Bay of Biscay (p-value < 0.001 for anchovy and p-value = 0.54 for
319 sardine). Both species also displayed size-related patterns (Figure 3). Larger sardine were more
320 likely to be in good (OR > 1 and p-value < 0.001) and poor (OR > 1 and p-value < 0.001) condition
321 state than smaller sardine. Results are similar for anchovy with larger individuals more likely to be
322 in good (OR > 1 and p-value < 0.05) and poor (OR > 1 but p-value > 0.05) condition state than
323 smaller ones. Graphical representation supports these results (Figure A2). For anchovy, being in
324 good condition was unlikely during reproduction, whereas being in poor condition was more likely
325 (Figure 3). ORs, associated 95% CIs and p-values are available in Table A2 and Table A3. A
326 graphical representation of individual condition states for anchovy and sardine considering each
327 explanatory variable is provided in Figure A2.

328



336 *Figure 3: Summary of multinomial logistic regressions for anchovy (a) and sardine (b). Odds ratios*
 337 *and 95% confidence intervals for explanatory variables associated with condition states. The inter-*
 338 *mediate condition state served as the reference group. Filled dots indicate significance (p-value <*
 339 *0.05) and empty dots indicate non-significance (p-value > 0.05).*

340 3.3. Variation in water mass

341 The influence of proximate component content on water mass independently of fish length was
 342 explored using a multiple linear regression (Eq.1) (Table 3). First, length had a significant effect on
 343 water mass (p-value < 2.10^{-16}) for both species, explaining 88% of the variance for anchovy and
 344 94% for sardine. Second, protein or lipid content had no significant effect on water mass (p-value >
 345 0.05) for both anchovy and sardine. Third, ash content had a significant effect on water mass for
 346 anchovy (although explaining only 2.21% of the variance) but not for sardine.

347 *Table 3: Results of the multiple regression of water mass as a function of proximate component*
 348 *content and length, for anchovy and sardine.*

	Anchovy			Sardine		
	Coefficient	p-value	Explained variance	Coefficient	p-value	Explained variance
Intercept	-23.0 ± 3.5	$3.6 \cdot 10^{-9}$		-56.0 ± 8.0	$2.2 \cdot 10^{-10}$	
Length	3.0 ± 0.1	$< 2.10^{-16}$	88%	5.6 ± 0.2	$< 2.10^{-16}$	94%
Protein	3.6 ± 13.5	0.80	0.002%	-30.0 ± 28.3	0.29	0.06%

Lipids	9.8 ± 6.0	0.10	1.44%	-3.7 ± 11.4	0.74	0%
Ash	-234.1 ± 44.8	9.9.10 ⁻⁷	2.21%	-67.3 ± 105.0	0.52	0.02%

349

350 3.4. Relationships between mass of water and proximate components

351 Strong log-log linear relationships were found between protein and water masses, as well as
 352 between ash and water masses, for anchovy and sardine (Table A4). R² were similar for protein
 353 (0.93 and 0.95 for anchovy and sardine, respectively) but differed for ash (0.89 and 0.96 for
 354 anchovy and sardine, respectively). Lipid mass exhibited higher variability for a given water mass,
 355 with a low associated R² (< 0.3) for both species.

356 The relationships between the mass of lipids, of proteins, of ash and that of water were investigated
 357 for each of the three condition states (Figure 4). Log-log linear model relating lipid to water masses,
 358 segmented by condition state, displayed higher explained variance (R² > 0.7) for individuals in good
 359 state for both anchovy and sardine (Table A4). Significant differences in intercepts were found
 360 between condition states in the segmented linear models of protein and water masses and lipids and
 361 water mass (all P-values < 0.001 for anchovy and sardine in ANCOVA results), but not in the
 362 model relating ashes to water (ANCOVA P-values = 0.21 for anchovy and 0.36 for sardine, Figure
 363 4, Table 4). The interaction between the condition state and water mass had also a significant effect
 364 on proteins and lipids (all ANCOVA P-values < 0.001 for anchovy and sardine) but not on ashes
 365 (ANCOVA P-value = 0.16 for anchovy and 0.14 for sardine, Table 4).

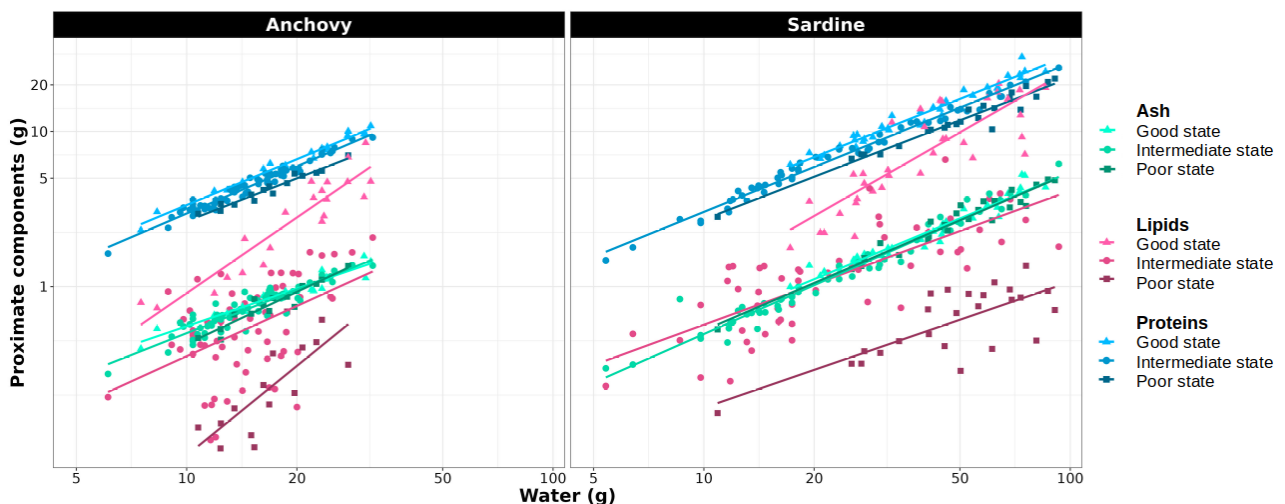
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371 *Figure 4: Observed mass (y-axis) of proteins (blue dots), lipids (purple dots) and ash (green dots)*
 372 *in anchovy and sardine body composition, as a function of water mass (x-axis). Solid lines:*
 373 *regression models for the three condition states for proteins (blue lines), lipids (purple lines) and*
 374 *ash (green lines).*

375 *Table 4: P-values of ANCOVA testing the effect of the condition state on the proximate components*
 376 *mass versus water mass relationship for anchovy and sardine.*

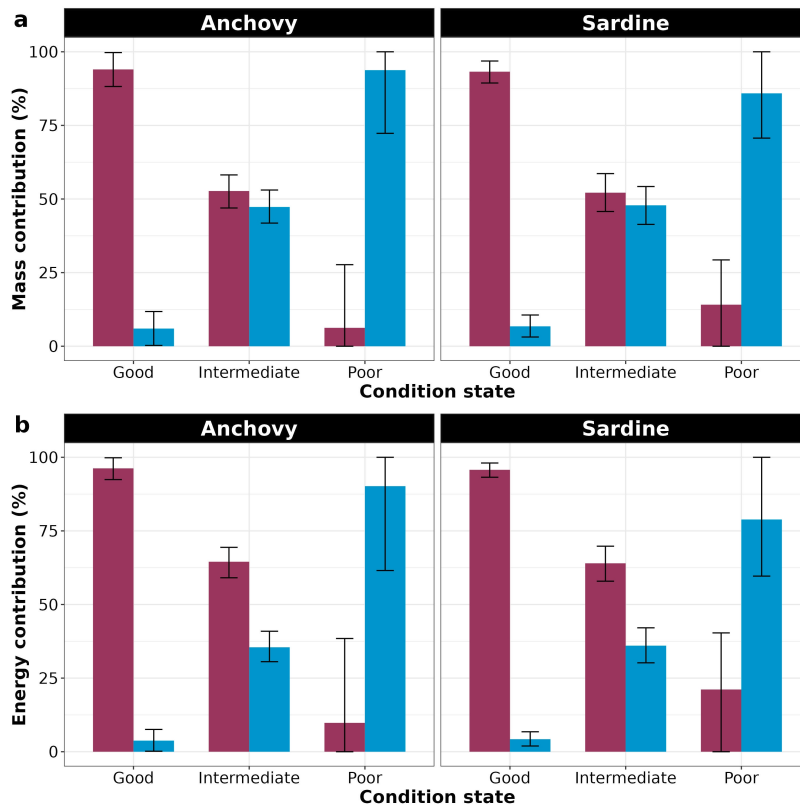
	Anchovy			Sardine		
	Proteins	Lipids	Ashes	Proteins	Lipids	Ashes
Water mass	< 2.10 ⁻¹⁶	< 2.10 ⁻¹⁶	< 2.10 ⁻¹⁶	< 2.10 ⁻¹⁶	1.10 ⁻¹³	< 2.10 ⁻¹⁶
State	< 2.10 ⁻¹⁶	< 2.10 ⁻¹⁶	0.207	< 2.10 ⁻¹⁶	< 2.10 ⁻¹⁶	0.357
Water mass * State	1.2.10 ⁻⁴	1.10 ⁻¹³	0.163	2.1.10 ⁻⁷	4.10 ⁻¹⁴	0.143

377

378 **3.5. Lipids and proteins contribution to energy mobilisation**

379 Based on the slope coefficients estimated within the three intervals, we determined the contributions
 380 and associated uncertainties of lipids and proteins to energy mobilization in terms of mass and
 381 energy (Figure 5, Table A5, Table A6). Within the good condition state, lipids emerged as the
 382 primary source of energy mobilisation in both anchovy and sardine, representing nearly all energy
 383 mobilized (94% and 96% in mass and energy, for anchovy and 93% and 96% in mass and energy,
 384 for sardine). In the intermediate condition state, lipids and proteins contributed roughly equally to
 385 energy utilization in terms of mass for both species (53% lipids and 47% proteins for anchovies;
 386 52% lipids and 48% proteins for sardines). However, given the higher energy density of lipids
 387 compared to proteins, lipids remained the dominant contributor to energy in terms of energy (65%
 388 lipids and 35% proteins for anchovies; 64% lipids and 36% proteins for sardines). In the poor
 389 condition state, proteins supplanted lipids as the primary source of energy for both species, due to
 390 lipid depletion (94% in mass and 90% in energy for anchovies; 86% in mass and 79% in energy for
 391 sardines).

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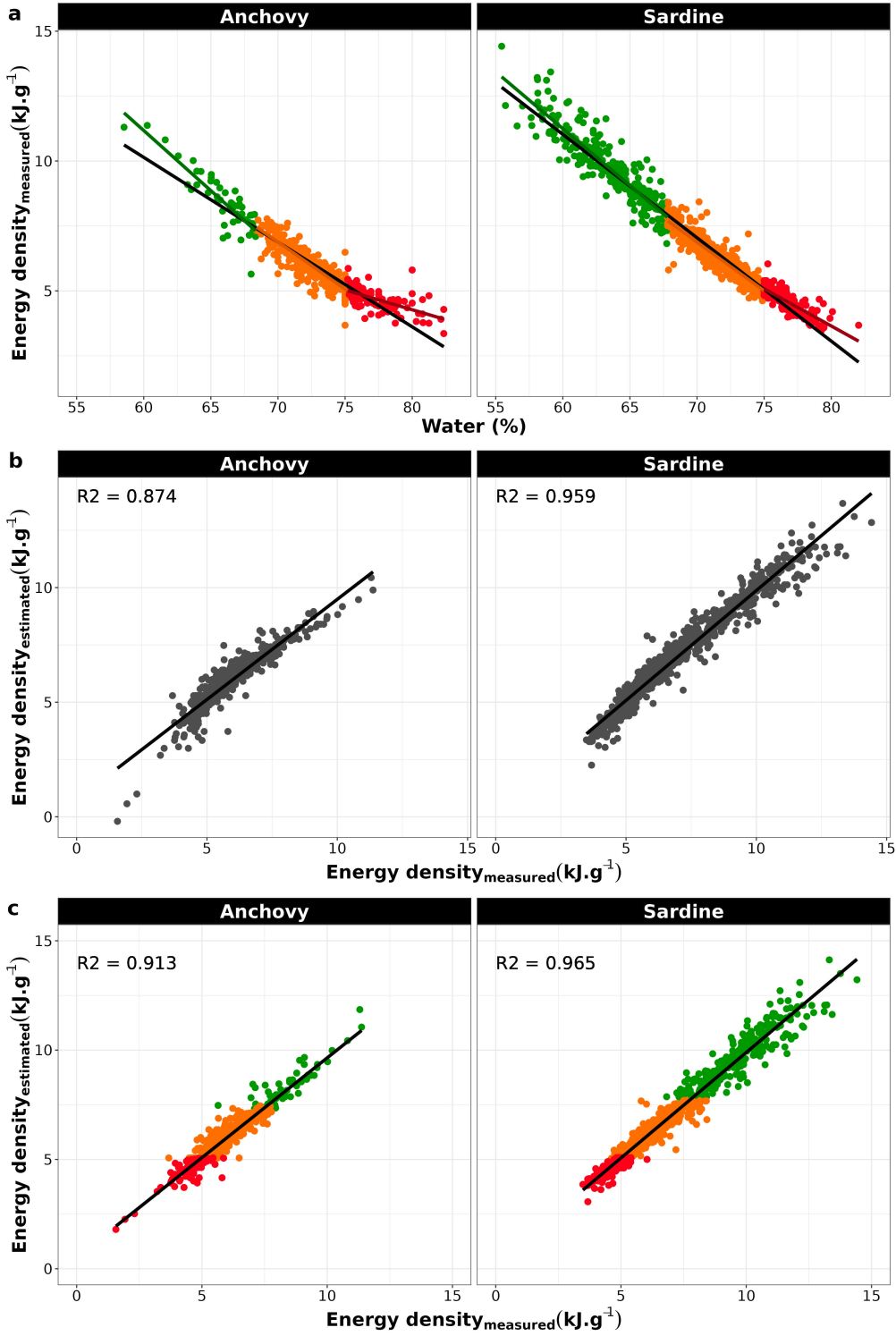
402 *Figure 5: Relative contribution to energy mobilisation of lipids (purple area) and proteins (blue*
403 *area) with associated uncertainty, expressed in terms of mass (a) and energy (b), for anchovy and*
404 *sardine according to the condition states (good, intermediate or poor).*

405

406 3.6 Relationship between fish water content and energy density

407 The linear models between energy density and water content, segmented or not by condition state,
408 are presented in Figure 6a. Significant differences in regression slopes (Table A7) were observed
409 between unsegmented and segmented linear regression, particularly for states of good and poor
410 conditions. The regression slopes for good condition (-46.3 ± 3.74 for anchovy and -43.5 ± 1.15 for
411 sardine) were higher, while the slopes for poor condition (-18.8 ± 1.14 for anchovy and -28.6 ± 1.29
412 for sardine) were lower than the regression slopes of unsegmented models (-31.2 ± 0.53 for anchovy
413 and -39.8 ± 0.26 for sardine). Using both segmented and unsegmented regression equations (for
414 good, intermediate, and poor states), we predicted energy density based on water content for both
415 anchovy and sardine. Energy density predictions from the unsegmented linear model were more
416 accurate for sardine ($R^2 = 0.959$) than for anchovy ($R^2 = 0.874$), primarily due to poor predictions of

417 extreme energy density values for anchovy (Figure 6b). Using a linear model segmented by each
418 condition state (Figure 6c) improved the accuracy of energy density predictions for fish in good and
419 poor condition, and also globally, particularly for anchovy ($R^2 = 0.913$).



421 *Figure 6: (a) Measured energy density as a function of water content (x-axis) and associated linear*
422 *models based on all individuals (black line), individuals in good (green dots and line), intermediate*
423 *(orange dots and line) and poor (red dots and line) condition states. Measured energy density (y-*
424 *axis) as a function of predicted energy density (x-axis) from a linear models fitted to all individuals*
425 *(b) and from segmented linear models fitted to individuals in good (green), intermediate (orange)*
426 *and poor (red) condition state (c).*

427 **4. Discussion**

428 Based on a unique dataset of energy density and proximate composition measured on almost 1500
429 anchovy and sardine sampled over a large region encompassing the Bay of Biscay and the English
430 Channel, we derived original patterns of variation and dependency between energy, water, lipid and
431 protein content. Working on wild individuals preclude any longitudinal studies in which individuals
432 and their condition are followed through time. Instead, we developed a cross-sectional approach
433 on a large number of individuals over a large range of size and from different seasons to uncover
434 those interesting patterns. We defined for the first time different condition states in wild fishes,
435 corresponding to different levels of lipid and protein contents, as well as to different contribution of
436 those metabolites when the lack of direct energy from food requires body reserves to be mobilised.
437 Following the identification of those condition states, we were able to i) assess that absolute water
438 mass, unlike relative water content, does not vary with fish condition, and ii) improve the modelling
439 of the non-linear relationship between energy density and water content.

440 *4.1. Three condition states based on relative proximate composition*

441 We identified three distinct condition states in wild anchovy and sardine, based on their lipid and
442 protein composition, relative to water content. Our finding is consistent with previous studies
443 describing three phases of starvation for mammals (Cherel *et al.*, 1988, 1991, 1992; Caloin, 2004),
444 birds (Le Maho *et al.*, 1981; Robin *et al.*, 1988; Groscolas and Robin, 2001) and fish (Bar, 2014).
445 We can not however ascertain that fish between good and intermediate states were starving, as we
446 did not follow individuals in a controlled longitudinal experiment. Instead, we may consider that the

447 three condition states identified in our study correspond to three different phases of energy
448 mobilisation or reserve constitution. The patterns of variation in the proximate composition were
449 strikingly similar between anchovy and sardine. Breck (2014) have also represented lipid and
450 protein variations relative to water content for five fish species (no small pelagic fish) but without
451 demonstrating different condition states. Investigating these variations in other fish species, in
452 particular small pelagics, would allow to investigate the genericity of these three condition states or
453 phases of energy mobilisation.

454 Our results confirmed that fish in good condition have high lipid (> 5%) and protein (> 20%) and
455 low water (< 68 %) contents. While some studies indicated that fish may use protein as primary
456 energy source (Marqueze *et al.*, 2017) or a combination of lipid and protein (Van Dijk *et al.*, 2005;
457 Gunasekera *et al.*, 2001.), we showed that anchovy and sardine primarily used lipids as source of
458 energy which was also described for other fish species (Savitz, 1971; Hung *et al.*, 1997). In the
459 intermediate condition state, fish exhibited lower lipid levels (< 5%), high yet decreasing protein
460 levels (> 20%), and average water content ranging from 68 to 75%. Within this state, fish continued
461 to utilise lipids as a source of energy but started to consume proteins, which is in agreement with
462 observations from other studies (Bar, 2014). Within the poor condition state, fish had nearly
463 depleted all lipids and only relied on protein mobilisation for energy production (Black and Love,
464 1986). We considered that the remaining ~1% lipids were structural (Bayse *et al.*, 2018; Biro *et al.*,
465 2021) and could not be mobilised. De Leonardis and Macciola (2004) showed that the difference of
466 total lipid between lean and fat sardine were exclusively due to increase of reserve lipid and not
467 affecting structural lipids. Fish in poor condition exhibited high water content in our study (>75%),
468 consistent with Breck's (2008) model results. Because protein metabolism is energetically
469 demanding (Schmidt-Nielsen., 1997; Anthony *et al.*, 2000), poor condition state impacts metabolic
470 rates (Queiros *et al.*, 2021) and is likely brief and critical. However, Queiros *et al.* (2021) showed
471 that some sardine were able to survive 57 days without food *in-vivo*. Survival time when starving
472 should be even shorter *in-situ*, as poor condition can impact swimming capacity (Martínez *et al.*,

473 2003; Faria *et al.*, 2011), and reduce predator escapement capacity (Biro *et al.*, 2021). Reversibility
474 from poor to intermediate condition state has been observed in controlled environments (Marqueze
475 *et al.*, 2017), but has never been documented in the wild so far.

476 4.2. What determines the fish condition state?

477 We identified three condition states along the gradient of water content corresponding to different
478 utilisation strategies (when water content increases), or creation (when water content decreases), of
479 lipid reserves or proteins. Since we were not in a starvation experiment framework, we explored the
480 internal and external drivers that could explain the position of the fish in a given state.

481 We found that season strongly influenced anchovy and sardine condition state. Our results are in
482 agreement with other studies studying anchovy and sardine condition in the Bay of Biscay through
483 energy density (Gatti *et al.*, 2018; Rosa *et al.*, 2010) or Le Cren body condition index (Véron *et al.*,
484 2020). They found minimum energy density (or Le Cren index) in winter and spring when food is
485 limiting and a peak in summer and autumn after energy storage during the feeding season. Same
486 patterns have been shown for the Northwest Atlantic forage species (Wuenschel *et al.*, 2024).
487 Seasonality in energy density in small pelagic fish has also been demonstrated in Mediterranean
488 sea, although the peak of energy density was in spring for anchovy and in autumn for sardine (Albo-
489 Puigserver *et al.*, 2017; Campanini *et al.*, 2021; Schismenou *et al.*, 2024). The seasonal effect under
490 environmental variability could also influence the lipid and fatty acid composition (Bandarra *et al.*,
491 1997 ; Biton-Porsmoguer *et al.*, 2020), affecting the fish physiology as some fatty acid are important
492 components for growth, survival and egg quality (Almansa *et al.*; 1999; Bruce *et al.*; 1999; Garrido *et*
493 *al.*, 2007; Tocher, 2003). However, variations in energy density are mainly due to changes in total
494 lipid content varying between 36.2 to 39.5 kJ.g⁻¹ (Schloesser and Fabrizio, 2015). Despite this
495 strong effect of season on the condition state, we found some fish in very poor condition in autumn,
496 i.e. two large sardines in reproduction. Conversely, no fish in good condition were found in winter
497 when food was limited, nor in spring when reserves did not have time to replenish.

498 Region has also an important impact on condition state with anchovy and sardine in good condition
499 mostly sampled in the English Channel, in agreement with other studies (Gatti *et al.*, 2018; Menu *et*
500 *al.*, 2023) which is most likely explained by more abundant food (Menu *et al.*, 2023), and can be
501 generalised along a latitudinal gradient in temperate systems (Huret *et al.*, 2019; Ljungström *et al.*,
502 2023).

503 For both species, length significantly influenced the condition state, with larger individuals of both
504 anchovy and sardine more likely to be in good condition, yet also more prone to being in poor
505 condition (though not significant for anchovy). Larger individuals seemed capable of accumulating
506 substantial reserves (Albo-Puigserver *et al.*, 2017) but also mobilizing proteins when lipid reserves
507 are depleted, in agreement with Gatti *et al.* (2018), who observed a decrease in energy density in
508 larger size classes of sardine. Conversely, the prevalence of smaller individuals in the intermediate
509 state suggested limited reserves and their absence in the poor condition state indicate an inability to
510 utilize protein for energy mobilization. These may use protein to prioritize rapid growth, but if
511 unsuccessful, they may succumb quickly when facing adverse conditions such as during winter
512 (Dubreuil and Petitgas, 2009).

513 The influence of reproductive status on fish condition has been extensively discussed (Pethybridge
514 *et al.*, 2014; Brosset *et al.*, 2015a; Albo-Puigserver *et al.*, 2017; Campanini *et al.*, 2021). In our
515 study, reproductive status was only retained in the anchovy's model, indicating that anchovies were
516 less likely to be in good condition and more likely to be in poor condition during reproduction.

517 Working on two species with two different spawning strategies highlighted that the main pattern of
518 variability in energy or proximate composition in our temperate study area was driven by external
519 factors (i.e. season and region). The significant effect of internal factors was mostly driven by their
520 interaction with external factors. Indeed, spawning sardines could be in poor or good condition,
521 mostly depending on their reproduction timing in spring or autumn, but not because of their
522 reproductive status. However some studies have demonstrated that reproduction has an effect on
523 lipid content and composition in the Mediterranean Sea (Vila-Belmonte *et al.*, 2024). We were not

524 able to highlight this effect for sardine in the Bay of Biscay. We mention that our dataset, although
525 extensive, is not fully representative of all variable combinations (season, region, length, and
526 reproductive status). This limitation was mainly due to species phenology as well as migration
527 patterns, e.g., anchovy being distributed in the North Sea and not in the English Channel during
528 spring and summer (Huret et al., 2020).

529 *4.3. Water mass does not vary with body proximate composition*

530 We showed that length had a significant effect on water mass, unlike the proportions of lipid or
531 protein. Therefore, water mass primarily relies on fish ontogeny rather than its lipid and protein
532 content. Although a significant negative correlation exists between lipids and water, it is only valid
533 in relative content. As lipid content decreases, water content increases but only in relative terms, not
534 in mass. These results did not agree with previous fish studies, which stated that fish replaced the
535 loss of lipids or protein with water to maintain osmotic pressure, cell size, and body shape (Love,
536 1970; Pangle and Sutton, 2005; McCue, 2010; Rosa *et al.*, 2010; Bar, 2014; Nemova *et al.*, 2021) or
537 on the opposite found that water did not replace lipid (Savitz, 1971; Breck, 2008) but rather
538 decreased together with lipids explaining the largest part of weight loss (Niimi, 1972; Breck, 2008
539 but see Collins and Anderson, 1995). Focusing on small pelagic fish, Dubreuil and Petitgas, (2009)
540 stated that anchovy replaced water with organic matter during feeding season and replaced organic
541 matter with water in winter. Breck (2008) argued that proteins are strongly associated with water,
542 leading to a loss of water when proteins are consumed. Groves (1970), suggested that proteins are
543 lost more rapidly than water and argued that a fish in starvation will have a lower protein:water
544 ratio than a fish in good condition, for the same length. We also showed that fish in poor condition
545 state exhibited a lower protein:water ratio, or lipid:water ratio, although we showed, thanks to an
546 analysis based on absolute mass rather than relative content, that this discrepancy is only
547 attributable to protein and lipid loss and not variation in water mass.

548 *4.4. Differential use of lipids and proteins among condition states explain the non*
549 *linear relationship between water content and energy density*

550 The strength of the relationship between water content (or dry content) and energy density has been
551 extensively documented (Hartman and Brandt, 1995; Tirelli *et al.*, 2006; Dubreuil and Petitgas,
552 2009; Gatti *et al.*, 2018). Non-linearity has been reported in this relationship (Hartman and Brandt,
553 1995; Gatti *et al.*, 2018), often attributed to ontogenetic and size-related effects (Wuenschel *et al.*,
554 2006) but without clear consensus on the mechanisms at stake. In our study, we showed that the
555 well-established relationship between water content and energy density varies for both anchovy and
556 sardine in response to fish condition state and body composition. Non-linearity in the energy to
557 water relationship can be modelled by applying a piece-wise regression, over each condition state
558 derived from proximate composition vs. water content models. The regression slopes for good
559 condition were steeper while the slopes for poor condition were flatter as compared to the
560 regression slopes of unsegmented models. A pronounced negative slope means a more pronounced
561 reduction in energy as water content increases, consistent with the initial loss of lipids with higher
562 energy density than proteins, followed by the loss of proteins within the poor state. Our findings
563 increase the predictability of energy density based on a water content proxy. The absence of non-
564 linearity in this relationship reported in some studies (Tirelli *et al.*, 2006; Dubreuil and Petitgas,
565 2009) may have stemmed from two main reasons. First, the relationship may not be nonlinear if
566 most fish are in the same condition state, e.g. in the case of reduced sample size and/or space-time
567 range. Second, differences in the contributions of lipids and proteins may be difficult to detect in the
568 case of lean fish with low lipid content (Anthony *et al.*, 2000). However, the genericity of this non-
569 linearity across species and beyond small pelagic species only (Hartman and Brandt, 1995) calls for
570 consideration of segmented regressions in future studies before using it to predict energy density.

571 *4.5. Conclusion*

572 This study highlights the importance of monitoring and predicting the intraspecific variations in
573 water content, proximate composition and energy content in small pelagic fish. We revealed
574 striking similarities between anchovy and sardine, suggesting that similar studies should be
575 conducted on other small pelagic fish species to verify the larger genericity of the observed patterns.
576 However, before doing so, these relationships should be determined on a case by case basis
577 (McPherson) to validate the use of proxy such as water content or fatmeter measurements, despite
578 some limitations of this device (Brosset et al, 2015, Schismenou et al, 2024).In particular, lipid
579 storage organs may vary from one species to another (e.g. muscle, liver, gonads), which should
580 influence the generalisation of this relationship. Integrating water content measurements during
581 surveys could provide new insight on individual and population conditions, especially in the context
582 of the decline condition in the European Atlantic (Doray *et al.*, 2018a; Véron *et al.*, 2020; Taboada
583 et al., 2024) and Mediterranean areas (Brosset *et al.*, 2017). Sardine spawners in better condition
584 have higher reproduction outcomes, i.e. more batches and more viable eggs (Rosa *et al.*, 2010;
585 Brosset *et al.*, 2016), potentially impacting recruitment. Monitoring changes in the proportion of
586 fish in good and poor condition states could offer insights into the evolution of natural mortality
587 rates, which may be increasing in the Bay of Biscay (Véron *et al.*, 2020) but considered as invariant
588 in stock assessment models (ICES, 2023). Finally, our observation of individuals in very poor
589 condition in the wild raises the following questions: Is this state reversible? If so, what proportion
590 of individuals survives annually? Is it a common state in the fish life cycle? A better
591 characterisation of the overwintering dynamics, especially the survival of individuals, and of the
592 interplay with reproductive strategies would certainly bring important knowledge for a better
593 understanding of the recruitment process. Furthermore, our research improved our understanding of
594 the dynamics of reserves (lipids and proteins) that should be taken into account for improved
595 realism of bioenergetic models (Breck, 2008; Gatti *et al.*, 2017).

596

597 ***Author contributions***

598 A.F. and M.H. designed the methodology. A.F. ran the analysis. S.L.M acquired the data. M.H.
599 raised fundings. A.F. wrote the original manuscript.. A.F., M.H., J.S., and M.D. contributed to
600 revising and editing the manuscript.

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609 ***Supporting information***

610 Additional supporting information can be found online in the Supporting Information section at the
611 end of this article.

612

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614 **Appendix A. Calculation of lipid and protein contributions to the variation in**
 615 **fish mass**

616 The contribution of lipid and protein, in both mass and energy, to energy mobilisation has been
 617 estimated as the mass variation of each of the two components in relation to the variation in total
 618 body mass. As we did not follow individuals through time like in experimental studies, we had to
 619 assume that the patterns of variation between the body components explored in our study based on
 620 our field sampling is representative of the patterns of variation of these components for an
 621 individual through time.

622 The starting point is thus the segmented regressions between the lipid or protein content and water
 623 content and its associated slope (equation 1A), which we had to transform in order to obtain a
 624 contribution in absolute mass of lipid or protein to the total body mass variation (equation 8A).

$$625 \quad \frac{d\left(\frac{W_C}{W_T}\right)}{d\left(\frac{W_W}{W_T}\right)} = a_C \quad (1A)$$

626 With W_c the component (lipid or protein) mass, W_w the water mass, W_T the fish body mass and a the
 627 component slope of the segmented regression.

628 Using $\left(\frac{u}{v}\right)' = \frac{u'v - uv'}{v^2}$ we can transform equation (1A) into equation (2A)

$$629 \quad \frac{\frac{dW_C * W_T - W_C * dW_T}{W_T^2}}{\frac{dW_W * W_T - W_W * dW_T}{W_T^2}} = a_C \quad (2A)$$

630 This can be simplified as:

$$631 \quad \frac{dW_C * W_T - W_C * dW_T}{dW_W * W_T - W_W * dW_T} = a_C \quad (3A)$$

632 Then, as we demonstrated (see our result section) that water mass does not vary, $dW_w = 0$:

$$633 \quad dW_C * W_T - W_C * dW_T = -a_C * W_W * dW_T \quad (4A)$$

634

635 Dividing both terms by dW_T we obtain:

$$636 \quad \frac{dW_C * W_T}{dW_T} - W_C = -a_C * W_W \quad (5A)$$

637 Or:

638
$$\frac{dW_C * W_T}{dW_T} = -a_C * W_W + W_C \quad (6A)$$

639 Dividing both terms by W_T we obtain:

640
$$\frac{dW_C}{dW_T} = \frac{-a_C * W_W + W_C}{W_T} \quad (7A)$$

641 Finally, simplifying W_W/W_T and W_C/W_T we obtain:

642
$$Contrib_C = \frac{dW_C}{dW_T} = -a_C * \%W + \%C \quad (8A)$$

643 With $Contrib_C$ the mass contribution of the component C to energy mobilisation, $\%W$ the water
644 content and $\%C$ the component (lipid or protein) content determined from the mean of all individual
645 data on considered interval. Since the contributions are based on independent regressions, we do not
646 obtain contributions somming to 1. Thus we normalised protein and lipid contribution.

647 Finally, to consider the energy contribution of each component, we had to weight the mass
648 contributions by the energy density of both components (equation 10A).

649

650
$$Contrib_{LEnergy} = \frac{Contrib_{Lnorm} * 38.5}{Contrib_{Lnorm} * 38.5 + Contrib_{Pnorm} * 23.6} \quad (9A)$$

651 With $Contrib_{LEnergy}$ the lipid energy contribution to energy mobilisation, $Contrib_{Lnorm}$ and $Contrib_{Pnorm}$
652 the normalized lipid and protein, respectively, masses contribution to energy mobilisation. Protein
653 energy contribution $Contrib_{PEnergy} = 1 - Contrib_{LEnergy}$.

654 To calculate the uncertainties associated to both mass and energy contribution to energy
655 mobilisation, we calculated mass and energy contribution (equations 8A to 10A) by taking the
656 minimum and maximum value of slope $a_{Cmin} = a_C - standard\ error$ and $a_{Cmax} = a_C + standard\ error$
657 for each component.

659 **Appendix B. Segmented regression equations**

660 *Table A1: Slope coefficients and associated standard error (SE) and confidence interval at 95% (IC*
 661 *95%) of observed proportions of proteins and lipids in sardine and anchovy body composition, as a*
 662 *function of water content for each condition state determined by the segmented regression analysis.*

	Anchovy				Sardine		
	State	Coefficient	SE	IC 95 %	Coefficient	SE	IC 95 %
Water ~ Lipids	Good	-1.38	±0.11	[-1.59 ; -1.18]	-1.43	±0.12	[-1.66 ; -1.20]
	Intermediate	-0.64	±0.09	[-0.82 ; -0.46]	-0.78	±0.10	[-0.97 ; -0.59]
	Poor	-0.09	±0.31	[-0.71 ; 0.53]	-0.16	±0.18	[-0.53 ; 0.20]
Water ~ Proteins	Good	0.23	±0.09	[0.05 ; 0.42]	0.21	±0.06	[0.09 ; 0.33]
	Intermediate	-0.32	±0.06	[-0.43 ; -0.21]	-0.47	±0.11	[-0.69 ; -0.26]
	Poor	-1.36	±0.51	[-2.36 ; -0.34]	-0.87	±0.21	[-1.29 ; -0.45]

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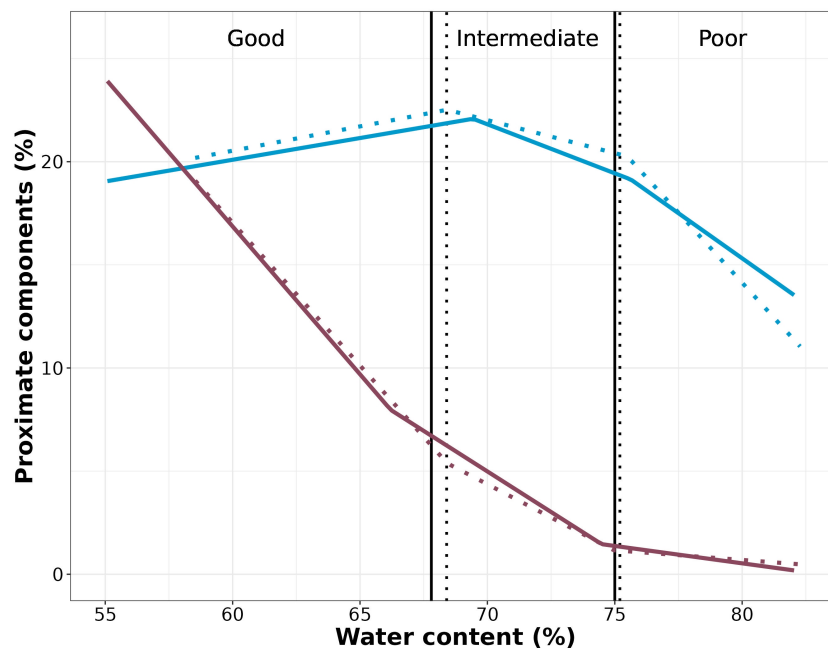
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675 *Figure A1: Comparison of segmented regressions of lipids (purple lines) and proteins (blue lines)*
 676 *versus water content for sardine (solid lines) and anchovy (dotted lines), across different condition*
 677 *states. The vertical lines represent the mean values of the breakpoints identified from proteins and*
 678 *lipids segmented regressions for sardine (solid lines) and anchovy (dotted lines).*

679 **Appendix C. Length, season, reproductive status and region versus**
 680 **condition state**

681 *Table A2. Multinomial logistic regression table showing the odd ratio and 95% confidence interval*
 682 *of explanatory variables associated with condition states for sardine (reference group = intermedi-*
 683 *ate state).*

SARDINE	Good state	Poor state
Season		
<i>Winter</i>	1	1
<i>Spring</i>	5.09.10⁻³ [5.09.10⁻³ ; 5.10.10⁻³]***	1.02.10⁻¹ [1.47.10⁻² ; 7.10.10⁻²]*
<i>Summer</i>	4.09.10⁵ [1.02.10⁵ ; 1 62.10⁶]***	0.45 [0.05 ; 3.70]
<i>Autumn</i>	4.55.10⁵ [1.45.10⁵ ; 1 42.10⁶]***	0.26 [0.04 ; 1.86]
Region		
<i>Bay of Biscay</i>	1	1
<i>English Channel</i>	3.13 [0.95 ; 10.4]	0.62 [0.13 ; 2.93]
Length	1.37 [1.15 ; 1.63]***	1.57 [1.25 ; 1.97]***
P-value: * <0.05, ** <0.01, *** =<0.001		

684 *Table A3. Multinomial logistic regression table showing the odd ratio and 95% confidence interval*
 685 *of explanatory variables associated with condition states for anchovy (reference group = intermedi-*
 686 *ate state).*

ANCHOVY	Good state	Poor state
Season		
<i>Winter</i>	1	1
<i>Spring</i>	6.68.10⁻⁴ [6.68.10⁻⁴ ; 7.00.10⁻⁴]***	3.58.10⁻¹⁴ [5.08.10⁻¹⁵ ; 2.54.10⁻¹³]***
<i>Summer</i>	6.91.10¹ [2.27 ; 2.10.10³]*	4.33.10⁻¹³ [8.39.10⁻¹⁴ ; 1.88.10⁻¹⁶]***
<i>Autumn</i>	9.75.10¹ [4.76 ; 2.00.10³]**	1.36.10⁻⁶ [2.18.10⁻⁸ ; 1.00.10⁻⁴]***
Region		
<i>Bay of Biscay</i>	1	1
<i>English Channel</i>	1.06.10 ¹ [0.97 ; 1.15.10 ²]	6.33.10⁻⁶ [6.32.10⁻⁶ ; 6.34.10⁻⁶]***
Length	1.72 [1.03 ; 2.87]*	1.70 [0.89 ; 3.22]
Reproductive status		
<i>No</i>	1	1
<i>Yes</i>	1.98.10⁻⁸ [1.98.10⁻⁸ ; 1.98.10⁻⁸]***	8.16.10⁷ [8.56.10⁶ ; 7.77.10⁸]***

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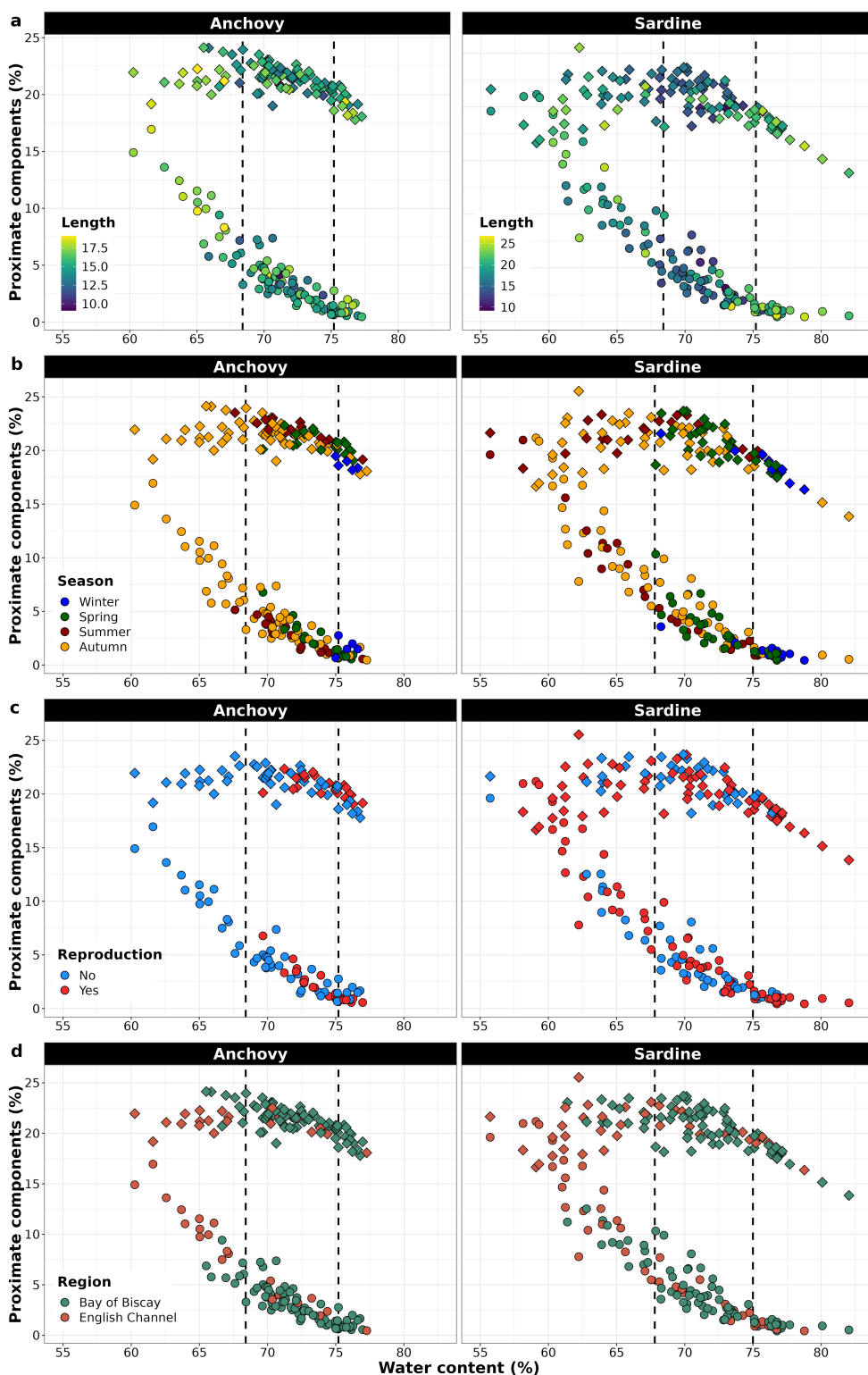
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714 *Figure A2: Observed content of proteins (diamond) and lipids (circle) in sardine and anchovy body*
715 *composition, as a function of water content with colors representing (a) the length, (b) the season,*
716 *(c) the reproductive status and (d) the region. The dotted vertical lines represent the mean values of*
717 *the breakpoints identified from proteins and lipids segmented regressions, allowing to separate the*
718 *three condition states.*

719 **Appendix D. Regression equation characteristics of proximate components**
 720 **versus water for each condition state**

721 *Table A4: Regression equation and associated standard error (\pm SE) and determination coefficient*
 722 *(R^2) of observed mass of proteins, lipids and ash in sardine and anchovy body composition, as a*
 723 *function of water mass for all individuals (all) and for each condition state (good, intermediate,*
 724 *poor)*

		Anchovy	Sardine
log(Proteins) ~ log(Water)	All	-1.26 (\pm 0.07) + 1.01(\pm 0.03) x, $R^2 = \mathbf{0.93}$	-1.01 (\pm 0.07) + 0.93(\pm 0.02) x, $R^2 = \mathbf{0.95}$
	Good	-1.05 (\pm 0.09) + 0.98(\pm 0.03) x, $R^2 = \mathbf{0.98}$	-0.90 (\pm 0.12) + 0.94(\pm 0.03) x, $R^2 = \mathbf{0.96}$
	Intermediate	-1.24 (\pm 0.06) + 1.01(\pm 0.02) x, $R^2 = \mathbf{0.97}$	-1.10 (\pm 0.05) + 0.96(\pm 0.02) x, $R^2 = \mathbf{0.98}$
	Poor	-1.14 (\pm 0.15) + 0.92(\pm 0.05) x, $R^2 = \mathbf{0.95}$	-1.11 (\pm 0.17) + 0.91(\pm 0.04) x, $R^2 = \mathbf{0.95}$
log(Lipids) ~ log(Water)	All	-4.58 (\pm 0.64) + 1.51(\pm 0.23) x, $R^2 = \mathbf{0.29}$	-2.39 (\pm 0.53) + 0.85(\pm 0.15) x, $R^2 = \mathbf{0.21}$
	Good	-3.83 (\pm 0.44) + 1.62(\pm 0.15) x, $R^2 = \mathbf{0.85}$	-3.00 (\pm 0.54) + 1.35(\pm 0.15) x, $R^2 = \mathbf{0.71}$
	Intermediate	-3.25 (\pm 0.56) + 0.98(\pm 0.21) x, $R^2 = \mathbf{0.23}$	-2.56 (\pm 0.31) + 0.86(\pm 0.10) x, $R^2 = \mathbf{0.59}$
	Poor	-6.93 (\pm 1.32) + 1.92(\pm 0.46) x, $R^2 = \mathbf{0.53}$	-3.67 (\pm 0.59) + 0.81(\pm 0.15) x, $R^2 = \mathbf{0.53}$
log(Ash) ~ log(Water)	All	-2.80 (\pm 0.09) + 0.92(\pm 0.03) x, $R^2 = \mathbf{0.89}$	-3.08 (\pm 0.07) + 1.04(\pm 0.02) x, $R^2 = \mathbf{0.96}$
	Good	-2.44 (\pm 0.15) + 0.81(\pm 0.05) x, $R^2 = \mathbf{0.93}$	-2.83 (\pm 0.16) + 0.98(\pm 0.04) x, $R^2 = \mathbf{0.94}$
	Intermediate	-2.84 (\pm 0.12) + 0.93(\pm 0.04) x, $R^2 = \mathbf{0.87}$	-3.11 (\pm 0.09) + 1.05(\pm 0.03) x, $R^2 = \mathbf{0.96}$
	Poor	-3.42 (\pm 0.30) + 1.12(\pm 0.10) x, $R^2 = \mathbf{0.89}$	-2.98 (\pm 0.20) + 1.01(\pm 0.05) x, $R^2 = \mathbf{0.94}$

726 **Appendix E. Lipid and protein contributions to the variation in fish mass**

727 Table A5: Slope coefficients in segmented regression of protein and lipids and their relative
 728 contribution as energy source (calculated using equations 2 and 3) in anchovy across the condition
 729 states. a_L : lipid slope and associated standard error, a_P : protein slope and associated standard
 730 error C_L mass: lipid mass contribution, C_P mass: protein mass contribution, C_L energy: lipid energy
 731 contribution, C_P energy: protein energy contribution.

	Anchovy					
	Lipids			Proteins		
	a_L	C_L mass	C_L energy	a_P	C_P mass	C_L energy
Good	1.38 ± 0.11	94%	96%	0.23 ± 0.09	6%	4%
Intermediate	-0.64 ± 0.09	53%	65%	-0.32 ± 0.06	47%	35%
Poor	-0.09 ± 0.31	6%	10%	-1.36 ± 0.51	94%	90%

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734 Table A6: Slope coefficients in segmented regression of protein and lipids and their relative
 735 contribution as energy source (calculated using equations 2 and 3) in sardine across the condition
 736 states. a_L : lipid slope and associated standard error, a_P : protein slope and associated standard
 737 error C_L mass: lipid mass contribution, C_P mass: protein mass contribution, C_L energy: lipid energy
 738 contribution, C_P energy: protein energy contribution.

	Sardine					
	Lipids			Proteins		
	a_L	C_L mass	C_L energy	a_P	C_P mass	C_L energy
Good	-1.43 ± 0.12	93%	96%	0.21 ± 0.06	7%	4%
Intermediate	-0.78 ± 0.10	52%	64%	-0.47 ± 0.11	48%	36%
Poor	-0.16 ± 0.18	14%	21%	-0.87 ± 0.21	86%	79%

739 **Appendix F. Difference between simple and segmented regression for**
 740 **water~energy density regression**

741 *Table A7: Regression equation and associated standard error (\pm SE) and determination coefficient*
 742 *(R^2) of observed energy density as a function of percent water for all individuals (all) and for each*
 743 *condition state (good, intermediate, poor)*

		Anchovy	Sardine
Energy density ~ water	Simple	28.7 (\pm 0.39) – 31.2 (\pm 0.53) x, $R^2 = \mathbf{0.87}$	34.9 (\pm 0.19) – 39.8 (\pm 0.26) x, $R^2 = \mathbf{0.96}$
	Strong	38.9 (\pm 2.46) – 46.3 (\pm 3.74) x, $R^2 = \mathbf{0.75}$	37.3 (\pm 0.73) – 43.5 (\pm 1.15) x, $R^2 = \mathbf{0.83}$
	Intermediate	32.2 (\pm 0.78) – 36.2 (\pm 1.07) x, $R^2 = \mathbf{0.77}$	33.1 (\pm 0.55) – 37.5 (\pm 0.77) x, $R^2 = \mathbf{0.83}$
	Poor	19.2 (\pm 0.88) – 18.8 (\pm 1.14) x, $R^2 = \mathbf{0.71}$	26.5 (\pm 0.99) – 28.6 (\pm 1.29) x, $R^2 = \mathbf{0.71}$

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