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1 **Title: Head shape disparity impacts pollutant accumulation in European eel**

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27 **ABSTRACT**

28 Several aspects of the life cycle of the critically endangered European eel (*Anguilla anguilla*)
29 remain poorly understood. One such aspect is the broad- versus narrow-head dimorphism,
30 and how this impacts their overall performance at different stages of their life cycle. At the
31 yellow eel stage, the phenotypes show a trophic divergence. We investigated whether
32 pollutant accumulation is affected by this disparity. We show that broad-headed eels
33 contained higher concentrations of mercury and several lipophilic organic pollutants,
34 compared to narrow-headed ones, irrespective of their fat content. The hereby confirmed link
35 between the phenotypic disparity, its associated feeding ecology and its impact on pollutant
36 accumulation thus raises further concerns about their migratory and reproductive
37 success. Considering that pollution is an important contributor to the European eel's decline,
38 our results demonstrate that broad-headed eels are more vulnerable to detrimental pollutant
39 accumulation. This compromises their successful contribution to their population's
40 reproduction and its restoration.

41 **Keywords** : Pollution – Biomagnification – Dimorphism – POPs – Trace metals

42 **CAPSULE**

43 Pollutant uptake rate in European eel can be governed by trophic segregation associated with
44 morphological specialization, with the rate depending on the contaminants lipophilicity.

45 **INTRODUCTION**

46 While European eels (*Anguilla anguilla*) have been studied extensively, many questions
47 regarding their unique life cycle still remain unanswered. Dependent on oceanic current
48 dynamics, this catadromous fish arrives as an un-pigmented glass eel at the continental waters
49 of Europe (1). They continue their journey up the rivers, where they feed and grow into yellow

50 eels. For several years, they accumulate a fat reserve of at least 12% of their body weight,
51 which is required for their migration back across the ocean and for gonadal development, to
52 become the mature silver eels that will eventually spawn in the Sargasso Sea (2, 3). Worryingly,
53 the analysis of the current glass eel recruitment indices at European monitoring stations have
54 revealed a decrease to only 1%, compared to the late 1970's, in the North Sea region and to
55 less than 5% in the rest of Europe (4). Consequently, the European eel is considered critically
56 endangered on the International Union for Conservation of Nature (IUCN) red list. Following
57 the EU Eel Regulation put in place in 2007 to restore the European eel stock (Council
58 Regulation (EC) No. 1100/2007), EU Member States have developed Eel Management Plans
59 (EMPs) with the objective to obtain a silver eel biomass escapement of 40%, compared to a
60 situation without anthropogenic disturbances. Alarmingly, more than 50% of the current EMP
61 progress reports show a failing to meet this target (5), indicating that much more effort is
62 needed to be put towards the European eel's recovery. A thorough understanding of its ecology
63 is thereby crucial to develop proper management measures. Among other factors, such as
64 habitat loss (6), migration barriers (7), parasites (8), overfishing (9), climate change and
65 changes in oceanic conditions (10, 11), the accumulation of pollutants could be one of the
66 possible synergistic causes that are contributing to the decline of the eel population (12).

67 Eels bio-accumulate many lipophilic pollutants in their fat tissue throughout their life. When
68 eels start migrating and turn into silver eels, they stop feeding and thrive on their fat reserves.
69 The continuous fat burning, in combination with endocrine-induced morphological and
70 physiological changes during migration, induces a remobilization of pollutants into the
71 bloodstream and leads to a subsequent increase in tissue concentrations of target organs,
72 such as the developing gonads (13-15). Bio-accumulation of toxic compounds can lead to
73 physiological disturbances, lowered resistance, disturbed reproduction and possibly even

74 death, before eels reach the spawning area (14, 16). Even when eels successfully complete the
75 migration, evidence has shown that 17-52% of the original fat reserves, together with its
76 pollutants, are incorporated in the oocytes. This maternal transfer of contaminants to eggs is
77 expected to increase the mortality of larvae, especially when the eels come from highly
78 polluted environments (17). Thus, pollution impacting physiological processes during the
79 European eel's life cycle is thought to be a crucial factor contributing to their decline.

80 While eels can absorb pollutants through their gills and skin, the major uptake of Persistent
81 Organic Pollutants (POPs) and trace metals, such as mercury, occurs through the ingestion of
82 contaminated food. European yellow eels demonstrate trophic divergence which is associated
83 with a head shape dimorphism: broad-headed eels consume more fish and crustaceans,
84 whereas narrow-headed eels mainly feed on benthic invertebrates (18-21). Also anatomically,
85 the dimorphism involves variation in the musculoskeletal components of the feeding
86 apparatus, which increases biting performance in broader heads (22). Here, we investigated
87 whether this disparity in trophic ecology could also affect the accumulation of pollutants,
88 which could alter reproductive ability. We hypothesize that broad-headed eels, which feed
89 higher in the food chain (23), are more vulnerable to pollutant accumulation and are thus
90 more susceptible to the detrimental pollutant effects, due to the consumption of more
91 contaminated prey. Gaining insight into the interaction of the eel's morphology with its
92 feeding ecology and pollutant accumulation would not only be a key element for improving
93 recovery efforts, but it would also shed some much needed light onto how feeding-associated
94 morphological variation can cause differential pollutant accumulation in eels.

95

96

97 MATERIAL AND METHODS

98 1. Experimental design

99 1.1 Sample collection

100 European eels (N = 377) were captured by fyke nets and electric fishing in Lake Weerde
101 (Belgium) in October 2015. Lake Weerde is a small (14 ha) lake in the Scheldt catchment,
102 which is located in a recreational and agricultural area, with no important industrial activity.
103 There is no open connection to a river system and all eels originated from glass eel restocking.
104 It was initially a lake with a depth of approximately 8 m, resulting from sand excavation
105 between 1968 and 1973. In order to create a more shallow lake with different shallow water
106 zones to increase biodiversity, 250 000 m³ of inert demolition material was used to partially
107 fill the lake during the 1990s. However, apparently, also toxic material had been dumped,
108 resulting in Lake Weerde being a highly contaminated lake for polychlorinated biphenyls
109 (PCBs) and other contaminants (considering the Flanders Eel Pollution Monitoring Network).
110 Because of these conditions, this lake was chosen for the present study. By annual monitoring,
111 it was shown that PCB body burden of eels were decreasing from 1998 to 2005 (24). However,
112 a steep increase of PCB contamination was observed in 2006, which was suspected to be due
113 to a local discharge or spilling of toxic waste containing PCBs (25).
114 All eels were anaesthetized with MS222 (Tricaine methanesulfonate). Subsequently, total
115 length (TotL) and total weight (W) were measured. Additionally, pictures of the head were
116 taken in dorsal view and head width (HW) was measured between the jaw hinges at the
117 nearest 0.1 mm, using a Mauser digital caliper. HW/TotL was calculated in the field. Eye size
118 was used to determine whether an eel was in the yellow or silver eel stage, as eye size
119 increases extensively during silvering (26). Out of the 377 eels, 75 yellow eels were selected
120 over a wide range of HW/TotL for further analyses. All the selected eels were larger than 46

121 cm (TotL_{min-max}: 46.5 – 67 cm), and thus were also all female (27), hence a sex effect could be
122 excluded. The other eels were released again into the wild once they had recovered from
123 anesthesia. The condition of the 75 eels was determined by calculating the Le Cren's condition
124 factor K (28) as follows: $K = W/W'$ where W is the observed weight and W' the calculated
125 weight based on the length-weight relationship ($W = a \times \text{TotL}^b$, where a is the intercept of the
126 slope and b the slope of the relationship). Subsequently, the 75 eels were anaesthetized by
127 MS222 and euthanized by an MS222 overdose, in accordance with the Belgian legislation. The
128 eels were decapitated and the heads were fixed in 10% formalin and preserved in 70%
129 ethanol. The body was skinned, intestines were removed and after decapitation, the body was
130 cut into four equal parts. The muscle tissue of the first three parts was used for further
131 analyses, while the final part acted as a reserve (figure S3). The muscle tissue of the first part
132 was used for stable isotope analysis, while the muscle tissue of the second part was used for
133 analysis of Persistent Organic Pollutants and the third part for the analysis of trace metals.
134 This protocol has been applied according to the methods developed by INBO during the Eel
135 Pollution Monitoring Network in Flanders (29) and described in ICES (30). It has been further
136 used during other international eel assessments (31, 32).

137 **1.2 Head shape determination**

138 We used a mixture analysis in PAST to visualize a bimodal distribution, based on the HW/TL of
139 the selected eels. We selected the point where the two unimodal distributions of the
140 frequency histogram overlapped as the separation value between broad- and narrow-heads.
141 The separation value was found at a HW/TL of 0.030. Based on this value, we considered eels
142 with a HW/TL lower than 0.0275 as narrow-heads, eels with a HW/TL higher than 0.0325 as
143 broad-heads and eels with a HW/TL between these values as intermediates. Using these cut-

144 off values, our dataset consisted of 26 narrow-heads (NH), 25 intermediates (INT) and 24
145 broad-heads (BH).

146 **2. Analyses**

147 ***2.1 Age determination***

148 The left and right sagittal otoliths were removed from the head, to determine the age of the
149 eels by the burning and cracking method (33, 34). This method is recommended for eel ageing,
150 especially for large eels (35). In short, the otoliths were first cut into two equal pieces and
151 were subsequently burned in a flame, revealing the annuli on the broken face. The otoliths
152 were then mounted cut face up onto a glass slide in silicone. Pictures of the otoliths were
153 taken using an SZX9 stereomicroscope, equipped with a ColorView 8 digital camera (Olympus,
154 Tokyo, Japan). These pictures were then used to determine the age of each eel. The age of
155 each eel was determined independently by two researchers in order to decrease possible
156 errors.

157 ***2.2 Body part 1: Stable isotope analyses***

158 One body part was used to obtain approximately 10 g of muscle tissue. These samples were
159 immediately frozen in liquid N₂ at -80°C and sent to ISOFYS (Isotope Bioscience lab, Ghent
160 University, Belgium), where they were stored in a freezer at -24°C. For analyses, the muscle
161 tissue was dried in an oven at 60°C for at least 48 h. The dried muscle tissues were then
162 completely homogenized and 100 µg per sample was used for stable isotope analyses. Total
163 N and $\delta^{15}\text{N}$ were measured using an elemental analyzer (ANCA-SL, SerCon, UK) coupled to an
164 isotope ratio mass spectrometer (20-22, SerCon, UK) (EA-IRMS). We used casein (%N = 13.32
165 and $\delta^{15}\text{N}$ = 5.94 per mil (Elemental Micro-analyses, UK)) as a laboratory standard that was
166 certified against NIST cysteine 143d for %N and IAEA-CH-6 for $\delta^{15}\text{N}$.

167 ***2.3 Body part 2: Persistent Organic Pollutants (POPs)***

168 The second body part of the eels was also stored in a freezer at -20°C and sent to the
169 Toxicological Centre of Antwerp University. Lipids and POPs were extracted and quantified
170 according to the method described by Malarvannan et al. (36). The measured analytes
171 included 36 PCB congeners (IUPAC numbers 18, 28, 31, 44, 47, 49, 52, 66, 70, 74, 87, 95, 99,
172 101, 105, 110, 118, 128, 138, 146, 149, 151, 153, 156, 167, 170, 171, 177, 180, 183, 187, 194,
173 196, 199, 206, 209), 7 PBDE congeners (IUPAC numbers: 28, 47, 99, 100, 153, 154, 183), three
174 DDTs (*p,p'*-DDT, *p,p'*-DDD and *p,p'*-DDE), hexachlorobenzene (HCB), three HCBs (α -, β - and
175 γ - isomers), cis- and trans-nonachlor (CN and TN, respectively), cis- and trans-chlordane (CC
176 and TC), oxychlordane (OxC) and anti- and syn-Dechlorane Plus (a-DP and s-DP). In short, a
177 homogenized sample of approximately 1 g eel muscle (ww) was weighed, then mixed with
178 anhydrous Na₂SO₄ and spiked with internal standards (BDE 77, BDE 128, ¹³C-BDE 209, CB 143
179 and ¹³C-HBCDs). Next, the samples were extracted for 2h by hot Soxhlet with 100 ml
180 hexane/acetone (3:1; v/v) and cleaned up on acidified silica. The lipid content was determined
181 gravimetrically on an aliquot of the extract (105 °C, 1 h). The remaining extract was cleaned
182 on 8 g of acidified silica (44%) and the analytes were then eluted with 20 mL hexane and 15
183 mL dichloromethane. The cleaned extract was concentrated to near dryness under a gentle
184 nitrogen stream, re-dissolved in 0.5 mL hexane and loaded on silica cartridges (500 mg/3 mL,
185 Agilent). The first fraction was eluted with 6 mL hexane and contained PCBs, DDTs, CHLs, HCB,
186 PBDEs and DPs, whereas the second fraction, containing the HBCDs, was eluted with 8 mL
187 DCM. Both fractions were evaporated to near dryness. The first fraction was re-solubilized in
188 100 μ L iso-octane, while the second was re-solubilized in 100 μ L methanol. PCBs, DDTs, CHLs,
189 HCB, PBDEs and DPs were analysed by GC-MS, whereas the HBCDs were quantified by LC-
190 MS/MS (25, 37). The detection limit was 0.05 ng/g ww for the PBDEs, 0.2 ng/g ww for the

191 lower chlorinated PCBs (CB18-CB66), 0.3 ng/g ww for the DDTs, and 0.1 ng/g for the remaining
192 analytes.

193 The POP analysis was evaluated by measuring the absolute recoveries of the internal
194 standards. Two criteria were used to quantify the peaks as target compounds. First, the
195 retention time of the peak had to match the retention time of the standard compound within
196 ± 0.1 min. Secondly, a signal-to-noise ratio (S/N) of 3:1 or higher was required. We determined
197 the *limit of quantification (LOQ)* by multiplying the standard deviation of the mean of the blank
198 measurements by three. Each batch contained seven samples and a procedural blank. The
199 latter was needed in order to detect possible interferences and/or contamination by solvents,
200 reagents or glassware. As consistent procedural blanks, with an RSD below 30%, were
201 obtained, we calculated the mean blank value of each compound. The values in the samples
202 were then subtracted by this mean value. The internal standard PCB 143 had a mean \pm SD
203 recovery of $86 \pm 6\%$, while the recovery was $93 \pm 10\%$ for BDE77. In order to validate the
204 analytical procedures, we analyzed the certified material SRM 1945 (organic contaminants in
205 whale blubber) which deviated less than 10% from certified values.

206 **2.4 Body part 3: Trace metals**

207 The third body part of the eels was stored in a freezer at -20 °C and sent to SPHERE (Systematic
208 Physiological & Ecotoxicological Research, University of Antwerp, Belgium). Of this body part,
209 0.5 g wet weight (ww) of homogenized fish muscle tissue was accurately weighed. Digestion
210 and analysis of the samples was done according to Mataba et al. (38). Dry weight (dw) was
211 determined after freeze drying. 1500 μ L of HCl (37%), 500 μ L of concentrated HNO₃ (69%),
212 200 μ L H₂O₂ and a magnetic stirrer were added to each of the samples. The samples were
213 subsequently digested using a SP-Discover microwave (CEM, USA) in two steps. The first step
214 ran at 120 °C, ramp and hold time of five minutes, with a maximum pressure of 34 bars at

215 300W and low stirring. Step 2 was identical to step 1 but ran at 160°C. To analyze Hg, the
216 samples were diluted upon 5-6% acid, whereas 1-2% acid was used for the other trace metals
217 (zinc, lead, nickel, copper, chromium, cobalt, cadmium and arsenic).

218 In order to control the quality, blanks and certified reference material (CRM) were added and
219 processed identical to the other samples. As CRM, both lyophilised Cod Muscle (BCR 422)
220 provided by the Institute for Reference Materials and Measurements (IRMM, Geel, Belgium)
221 and freeze dried muscle tissue (no. 2976) from NIST (National Institute of Standards and
222 Technology-USA) were used.

223 Subsequently, all eel samples were analyzed for total mercury in cold plasma mode by High
224 Resolution Inductively Coupled Mass Spectrometry (HR-ICP-MS; Thermo scientific Finnigan
225 element 3, Waltham, MA, USA), which has an instrumental detection limit of 0.01 µg/L. An
226 Inductively Coupled Plasma-Mass Spectrometer (ICP-MS, Varian UltraMass 700, Victoria,
227 Australia), with a detection limit of 0.1 µg/L, was used to measure the concentrations of the
228 remaining trace metals. The obtained concentrations are very similar to the certified
229 concentrations in the reference material, with a range from 90 to 110%, depending on the
230 metal.

231

232 **3. Data analysis**

233 Analytes that were below detection limit (BDL) in more than 50% of the specimens were not
234 used for further analysis (Table S1). Consequently, the BDEs 28, 99, 153, 154 and 180, the β-
235 and γ- isomers of HBCD and the trace metals Cd, Co and Ni were not used. Measurements that
236 were BDL in the remaining pollutants were estimated as detection limit*1/2. We need to
237 note, however, that the replacement of censored values by an arbitrary one, may have an
238 influence on the given means and standard deviation (Helsel, 2012). While little influence is

239 expected for the pollutants α -hexabromocyclododecane (α -HBCD; only 4% of the eels BDL),
240 HCB (4%) and BDE100 (1%), the replacement of the censored values could be more
241 pronounced for OxC (27%), Cr (23%), *p,p'*-DDT (20%) and TN (11%). Six PCBs (IUPAC numbers
242 28, 52, 101, 138, 153 and 180) and their sum were used for further statistical analysis, as these
243 are used to control for food safety according to European legislation. In addition, the sum of
244 *p,p'*-DDD, *p,p'*-DDT and *p,p'*-DDE (Σ DDT) was determined as a proxy of total DDTs.

245 A Shapiro-Wilk test was performed to investigate whether age, total length, weight, trophic
246 position, condition and fat content followed a normal distribution in the different phenotypes
247 (BHs, INTs and NHs). In case the assumptions were met, an ANOVA was performed to test
248 whether these variables differed significantly between the three phenotypes. If the data were
249 not normally distributed, a non-parametric permutation test (10,000 replicates) was used. In
250 addition, the measurements of these variables were log-transformed and were then used for
251 a Between-group Principal Component analysis (PCA) to find potential differences between
252 the phenotypes

253 Next, the lipid weight concentrations of the different POPs and the wet weight concentrations
254 of the trace metals were log-transformed. Subsequently, a Shapiro-Wilk test was performed
255 to test whether these concentrations in each phenotype followed a normal distribution.
256 Potential outliers were identified by performing a mixture analysis. In case one specimen
257 differed strongly from all the others, it was removed for further analysis. In case multiple
258 outliers were present, we performed non-parametric tests. Subsequently, we calculated the
259 Pearson (if data were normally distributed) or Spearman's ranked (if data were not normally
260 distributed) correlation between POPs (lipid weight basis), metals (wet weight basis), age and
261 TotL.

262 Furthermore, in case a normal distribution was not supported, we used a non-parametric
263 permutation test (10,000 replicates) to detect significant differences in the log-transformed
264 pollutant concentrations between the different phenotypes. When the assumptions were
265 met, an ANCOVA was performed to find significant differences in the log-transformed
266 pollutant concentrations between the different phenotypes. The ANCOVAs were performed
267 both with the log-transformed total length (TotL) and age as co-variate.

268 In addition, a between-group PCA were performed on the log-transformed lipid weight
269 concentrations of the POPs.

270 Finally, we determined the mean lipid weight POPs concentrations in the three phenotypes
271 and then calculated the mean(INT)/mean(NH) and mean(BH)/mean(INT) ratios. These were
272 then plotted against the log(Kow)-values of these POPs (39-42). This allowed for a simplified
273 data model, and comprehensible graphical representations, in order to clearly visualize POP
274 behaviour over the different phenotype groups. Kow represents the octanol/water partition
275 coefficient and the higher this value, the more hydrophobic (and thus lipophilic) a chemical is.
276 For the POPs that included eels with concentrations BDL (OxC, TN, HCB, α -HBCD, *p,p'*-DDT and
277 BDE100), we performed a sensitivity analysis by replacing the censored value by the minimum-
278 value (0), by detection limit*1/2 and by the detection limit and then calculating the means of
279 each phenotype. In case different patterns were observed, depending on the replacement
280 value, the POP was not used for this analysis.

281 **RESULTS**

282 **Head shape and trophic position**

283 Based on the head width/total length ratio (HW/TotL), 26 narrow-headed (NH), 25
284 intermediate (INT) and 24 broad-headed (BH) female yellow (size class: 46.5 – 67cm) eels were
285 selected. These groups did not differ significantly in total length, condition factor and body

286 weight. Intermediates were, however, significantly younger (Table 1). The fat content of NHs
287 was significantly higher compared to INTs and BHs (Table 1). Accordingly, a significant,
288 negative correlation was found between HW/TotL and fat content ($r=-0.36$; $P < 0.01$; Figure
289 1A). Trophic position (measured as $\delta^{15}\text{N}$), on the contrary, was positively correlated with
290 HW/TotL ($r=0.66$; $P < 0.01$; Figure 1B), but showed no correlation with TotL ($r=-0.07$; $P = 0.57$).
291 The $\delta^{15}\text{N}$ concentrations were significantly higher in BHs compared to INTs, while the
292 concentrations of the latter were significantly higher compared to NHs (Table 1). The
293 difference in average $\delta^{15}\text{N}$ between BHs and NHs equalled only 2.82‰, indicating that the
294 different phenotypes were less than one trophic position apart (43), which corresponds to
295 results obtained previously (23). Rather, the increase of $\delta^{15}\text{N}$ with HW/TotL indicates that eels
296 with a broader head consume relatively more prey items that are higher in trophic position.
297 This observation is supported by the literature, where a shift in $\delta^{15}\text{N}$ from 3.0 to 3.6 in omul
298 (*Coregonus migratorius*), for example, suggested that this fish did not only feed on benthic
299 invertebrates, but also on carnivores and omnivores (44). Our observations are also consistent
300 with the results obtained from stomach analyses, which indicated that broad-headed eels
301 consume proportionally more, but not exclusively larger prey items, such as fish and
302 crustaceans (2). The above described results are consistent with those of the between-group
303 PCA (Figure S1)

304 **Head shape and pollution**

305 Because of the lipophilic nature of POPs (45), we used the lipid-normalized concentrations
306 (expressed as ng/g fat) for our analyses. This also prevented a confounding effect of
307 differences in fat content. Several pollutants were found to be below the detection limit in
308 more than 50% of the eels and were, therefore, not included for further statistical analyses
309 (Table S1). Furthermore, of all measured PCBs, only the six ICES indicator PCBs and their sum

310 were included for further analyses (CB 28, 52, 101, 138, 153 and 180). The median \pm SD of the
311 pollutant concentrations is found in Table S2. None of the pollutants correlated with size or
312 age (Table S3). The results of the ANCOVA with TotL and age were comparable (Table 2). For
313 *p,p'*-DDE and -DDT, BDE47 and 100, OxC and TN, a non-parametric permutation test was used
314 since these data did not fall within a normal distribution (Table S4). HCB concentrations were
315 found to be significantly lower in eels with a broader head. The concentrations of the higher
316 chlorinated PCBs (CB 138-180), *p,p'*-DDE, *p,p'*-DDT and both Σ PCB and Σ DDT were found to be
317 similar between INTs and BHs, with only the concentrations of the higher chlorinated PCBs
318 and *p,p'*-DDT being highest in BHs, although for the latter, the median is used (Table 2-3).
319 These concentrations were significantly higher compared to NHs. The *p,p'*-DDD, BDE100 and
320 CB101 concentrations, on the other hand, were comparable between NHs and BHs, with
321 higher values measured in INTs. Finally, BDE47 and α -HBCD, OxC and TN differed strongly
322 between the three groups, with the concentrations being significantly higher in INTs (Table 2-
323 3). Interestingly, the mean concentration of α -HBCD and BDE47 and the median concentration
324 of OxC and TN were generally the highest in INTs.

325 However, the between-group PCA on the log-transformed pollutant concentrations indicated
326 that PC1 scores showed a lot of variation in BHs (Figure 2). The lower mean values of BHs,
327 compared to those of INTs, resulted from some BHs having low pollutant concentrations.
328 Furthermore, the between-group PCA indicated that BHs were generally characterized by a
329 combination of lower PC1-scores and higher PC2-scores, while the opposite was true for NHs.
330 INTs take up a rather intermediate morphospace. As such, BHs are mainly associated with
331 higher concentrations of higher chlorinated CBs (CB 138, 153, 180), *p,p'*-DDE and *p,p'*-DDT,
332 whereas NHs have a higher amount of CB28, α -HBCD and especially HCB. While INTs also have

333 high pollutant concentrations, these concentrations are not as high as in BHs or NHs (as PCA
334 shows).

335 Interestingly, the compounds with the highest concentrations in BHs are also the most
336 lipophilic ones. Therefore, the mean POP concentrations (not log-transformed) of NHs, INTs
337 and BHs were determined and used to calculate the mean(INT)/mean(NH),
338 mean(BH)/mean(INT) and mean(BH)/mean(NH) ratios. The sensitivity analysis showed only
339 different patterns for TN depending on the value by which a concentration BDL was replaced
340 and was therefore not included in this analysis (Table S5). All three ratios were significantly
341 positively correlated with lipophilicity (K_{ow} ; $r=0.84$, $r=0.83$ and $r=0.88$ for INT/NH, BH/INT and
342 BH/H respectively; $P < 0.01$ in all cases; figures 3 and S2), indicating that the more lipophilic a
343 pollutant is, the higher its concentration will be in broad-headed eels. Lipophilicity has also
344 been related to bio-magnification potential, i.e. the more lipophilic a chemical is, the more
345 likely it is to accumulate in food chains (45, 46). As such, the positive correlation between the
346 POP ratios and lipophilicity can be explained by differences in prey consumption, since the
347 studied eels did not differ in age, length or sex, except for the significantly younger age of INTs
348 compared to BHs and NHs. This indicates that the proportionally higher consumption of such
349 prey items leads to a cumulative uptake of highly lipophilic pollutants, which is consistent with
350 previous studies suggesting that the uptake of such pollutants occurs through food (47, 48).

351 Finally, to analyze the accumulation of trace metals, wet weight muscle concentrations (ng/g
352 eel) were used. Of all trace metals, only the measurements of mercury (Hg) and zinc (Zn)
353 followed a normal distribution (Table S4). According to the ANCOVA with Totl and age as co-
354 variate, the mercury concentrations were found to be significantly higher in BHs, compared to
355 INTs and NHs (Table 2), whereas no significant differences were found for Zn. Using a
356 permutation test, we also found that copper (Cu) concentrations were significantly higher in

357 BHs, compared to NHs (Table 3). Chromium(Cr) and arsenic (As), on the other hand, were
358 found to be significantly lower in BHs, compared to NHs. Arsenic mainly accumulates in the
359 exoskeleton of small invertebrates (49). Consequently, the higher consumption of these
360 invertebrates by NHs could explain this observation. Lead (Pb) concentrations were found to
361 be significantly lower in NHs compared to INTs, but did not differ significantly from BHs. The
362 results of the trace metal analysis were thus consistent with results obtained previously,
363 showing that Hg demonstrates a tendency to bio-magnify in food chains (50), whereas this is
364 generally not the case for the other metals (51). Moreover, most metals are hydrophilic and
365 will be taken up via the water rather than via the food. Additionally, most metals accumulate
366 in the organs rather than in the muscle tissue (38). Together with Hg, the only exception in
367 our study is Cu, which also increased with head width and thus trophic position.

368 Our novel findings show that there is a link between head shape, trophic position and
369 pollutant accumulation. Eels with broader heads feed more on prey items which are higher up
370 in the food chain, leading to an increased uptake of more lipophilic POPs and Hg, compared
371 to eels with a narrower head.

372 **DISCUSSION**

373 A driver for phenotypic-trophic disparity in European yellow eels could be to alleviate the
374 pressure on food competition, thus increasing their survival rates (52). Additionally, a recent
375 study has indicated that eels can adapt to differences in diet; young eels that were given hard
376 prey, such as crustaceans, developed broader heads than eels that were given soft prey, such
377 as chironomid larvae (53). The broader heads were associated with larger jaw muscles that
378 facilitate consumption of hard prey (22). The capability to show such an adaptive, phenotypic
379 response to differences in food accessibility and the decrease in intra-specific competition
380 through this dimorphism are likely to increase the eel's survival. As such, the dimorphism

381 could be sustained in the population through natural selection, and thus be considered as
382 adaptive. However, our results suggest that the causality between diet and head shape also
383 controls the pollutant accumulation. Anthropogenic influences may thus turn an adaptive trait
384 into a maladaptive one, as the advantage of adapting towards consuming larger and harder
385 preys in the broad-headed yellow eel may ultimately lead to a lower reproduction potential
386 as silver eel, due to increased pollutant accumulation (17).

387 While measurements of fat content in a particular subsample of muscle tissue may not be
388 representative for the complete fish (54), we observed that the fat content of broad-head eels
389 was significantly lower than that of NHs. This phenomenon could be explained by the trophic
390 difference, with broad-head eels feeding proportionally more on larger and harder prey items,
391 such as fish and crustaceans. The consumption of such larger prey is energetically more
392 profitable than the consumption of small, benthic invertebrates (55). However, obtaining such
393 prey items requires biting or spinning behaviour, in contrast to NHs sucking in small, benthic
394 invertebrates. It has been demonstrated that the net energy gain when using biting and
395 spinning behaviour is lower than when relying on suction feeding (56). It is thus possible that
396 the observed differences in fat content are a consequence of the different feeding methods.
397 Secondly, it is possible that prey items, such as benthic invertebrates, are more abundant and
398 thus easier to capture than the more evasive fish and crustaceans and that NHs hence require
399 less energy for prey capture.

400 Alternatively, several pollutants could have a negative impact on the eel's fat content, for
401 example due to a higher energy demand induced by chemical stress (14, 16). The lower lipid
402 energy reserves of BHs may thus also be a consequence of significantly higher concentrations
403 of the more lipophilic pollutants (57). The measured lower fat content of BHs could, in turn,
404 imply that BHs will also require a longer time to accumulate enough fat to start migration due

405 to pollution. Indeed, while similar in age, 54% of the NHs had accumulated enough fat (12%
406 of body weight (3)) to start migrating, while this was only 17% for the BHs. For INTs, this was
407 28%, but they were significantly younger. Pollution therefore may affect the BHs at four levels.
408 Firstly, during the fat accumulation period as yellow eels, where BHs need considerably more
409 time to accumulate enough lipid energy to migrate successfully. A delayed migration
410 inevitably increases the risk of being killed by predators or other factors. Secondly, if BHs
411 would still start their migration as silver eels, but with low fat concentrations, energy storage
412 would potentially be insufficient to fuel the transatlantic migration and production of
413 gametes, and thus leads to lowered reproduction potential (57). Thirdly, BHs have a higher
414 risk of being exposed to the detrimental effects of higher concentrations of pollutants, when
415 fat is being metabolised during migration. These effects may include disturbances of the
416 immune, reproductive, nervous and endocrine systems (14). Finally, being more loaded with
417 the most toxic pollutants, such as PCBs, could have a negative impact on the mean weight of
418 eggs by interfering with the development of the ovaries (58), possibly resulting in less viable
419 eggs. Our results therefore strongly indicate that BHs are expected to suffer more from POP
420 and mercury pollution than NHs at several (potentially cumulative) concentrations, due to the
421 trophic ecology related to head shape disparity. Additionally, the eel's head morphology has
422 also been related to parasite infection, showing that the higher piscivorous behavior of BHs
423 leads to a higher exposure and thus infection with the parasite *Anguillicoloides crassus* (59).
424 This parasite damages the swim bladder and drains the eel's energy by sucking blood, thus
425 also impairing the eel's migration success (8, 60). So, while natural selection would favour
426 dimorphism because of decreased intra-specific competition, anthropogenic effects may well
427 disrupt this adaptive process in polluted areas as the reproduction potential of BHs is expected
428 to be lower than that of narrow-heads. This is especially worrisome since current eel

429 restoration measures in the EU member states focus on producing maximal quantities of silver
430 eels leaving the river catchments (at least a 40% silver eel biomass escapement), and do not
431 take into consideration their overall body quality. It has been argued, however, that also the
432 health status of silver eels should be improved at the level of pollution load, infection levels
433 by pathological agents and fitness indicators, such as lipid levels and condition. These are all
434 factors that are crucial for maximizing the eel's reproductive potential, and hence for the
435 successful restoration of the European eel stock. Our results indicate that there could be
436 differences in reproductive potential between broad- and narrow-headed eels from polluted
437 environments. We would advise that the current stock assessment, which is based on
438 quantitative indicators, should be jointly merged with indicators characterizing the quality of
439 the eels leaving our catchments. This could potentially lead to an overall, much needed,
440 successful restoration of European eel stocks.

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596 **FIGURE CAPTIONS**

597 **Figure 1 (1 column width): Relation between fat content and trophic position with relative**
598 **eel head width.** A: Relation of fat content with eel head width relative to total length
599 (HW/TotL). B: Relation of trophic position ($\delta^{15}\text{N}$) with eel head width relative to total length
600 (HW/TotL). Light blue dots represents narrow-heads, blue dots intermediates and dark blue
601 dots broad-headed eels. A representative of each phenotype is given above. Black dots
602 represent the means of each phenotype; the error bars represent standard deviation.

603 **Figure 2 (1 or 1.5 column width): Between-group PCA on the log-transformed pollutant**
604 **concentrations and the corresponding biplot.** The pollutant concentrations were expressed
605 as lipid weight. Light blue dots represents narrow-heads, blue dots intermediates and dark
606 blue dots broad-headed eels. The amount of variation explained by each Principal Component
607 (PC) is shown in parentheses.

608 **Figure 3 (1 column width): Relation of the mean(BH)/mean(NH) ratio of the different**
609 **pollutants plotted against the log of their octanol/water partition coefficient (K_{ow}).** The
610 pollutants are expressed as lipid weight. NH: Narrow-heads, BH: Broad-heads. Blue diamonds
611 represent pentachlorobiphenyls (PCBs), green diamonds Dichlorodiphenyltrichloroethanes
612 (DDTs) and red the remaining measured pollutants.

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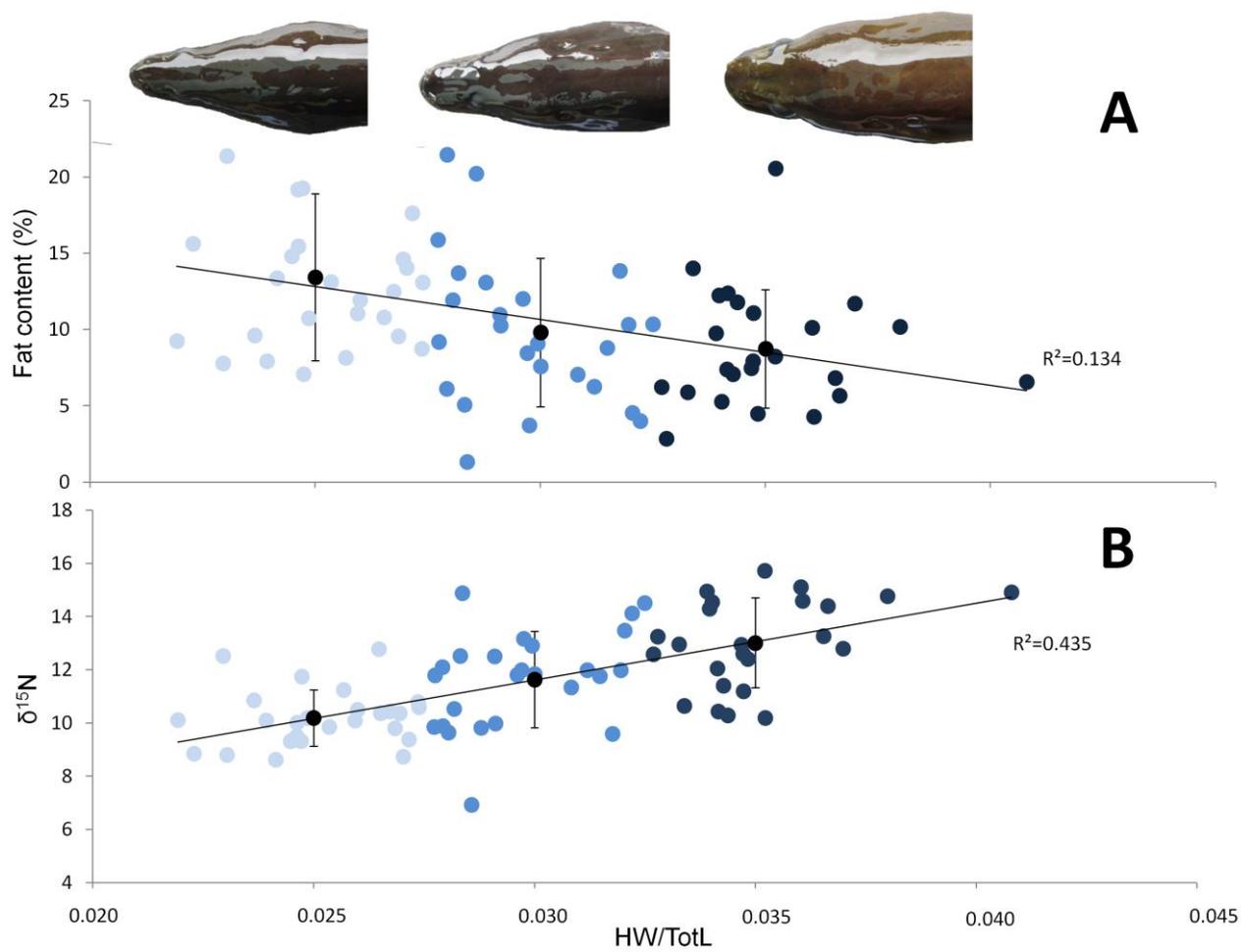
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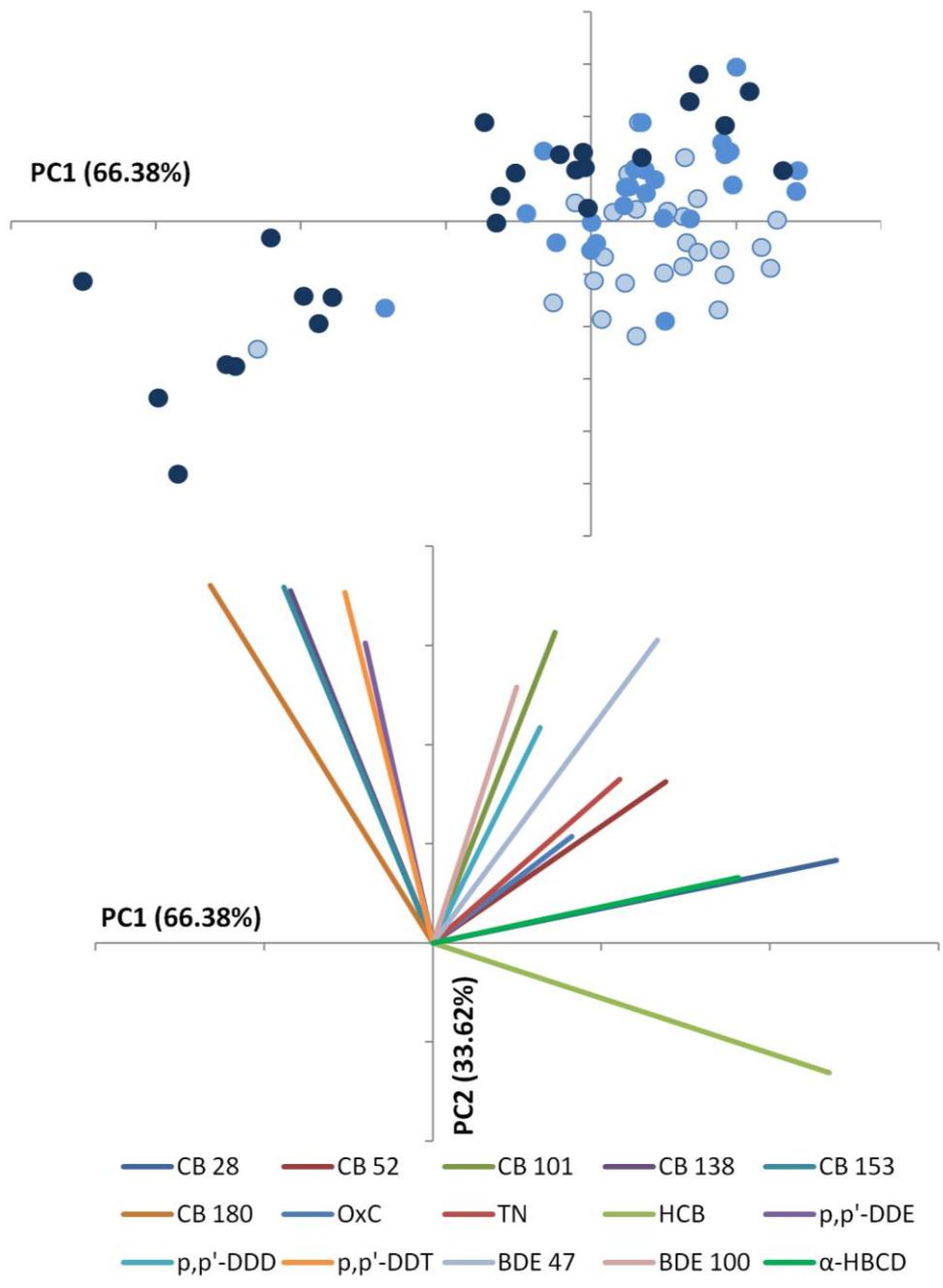
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619 Figure 1

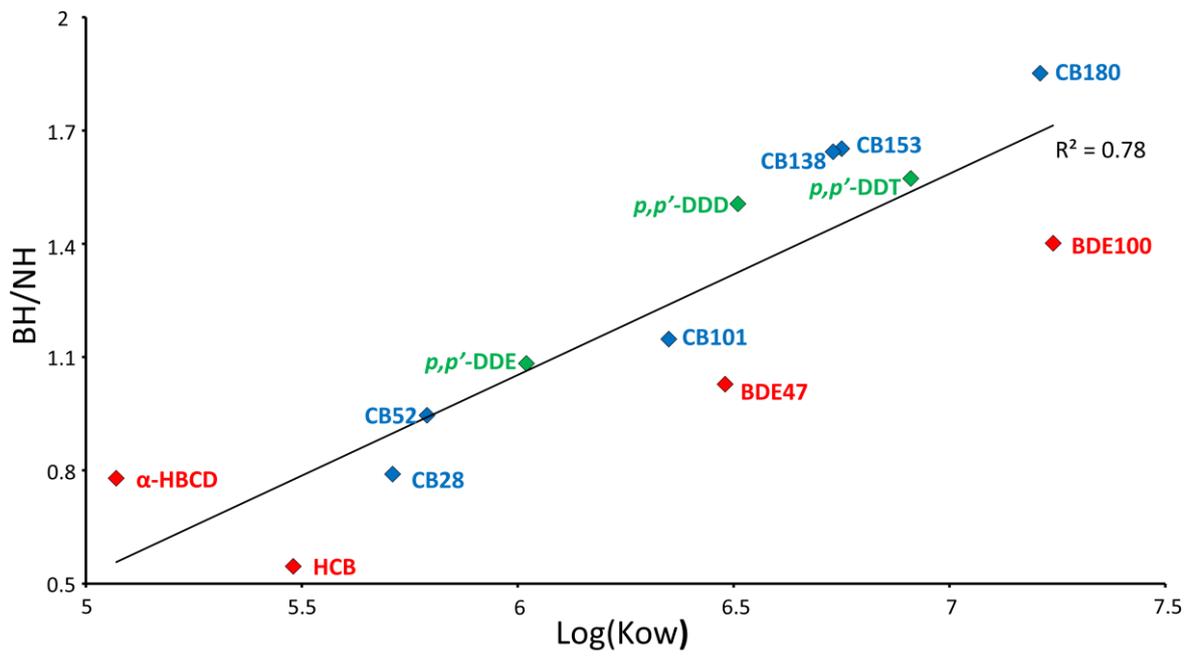


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624 **Figure 3**



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628 **TABLES**629 **Table 1: Basic characteristics in narrow-head, intermediate and broad-head eels**

	Mean \pm SD			ANOVA			
	NH	INT	BH	<i>P</i>	<i>P</i> _{NH-Int}	<i>P</i> _{Int-BH}	<i>P</i> _{NH-BH}
Head Width/TotL (-)	0.025 \pm 0.002	0.030 \pm 0.002	0.035 \pm 0.002	< 0.01	< 0.01	< 0.01	< 0.01
Total Length (cm)	52.1 \pm 3.4	53.2 \pm 4.9	54.5 \pm 6.0	0.22	0.71	0.61	0.20
Total Weight (g)	224 \pm 56	253 \pm 86	273 \pm 112	0.10	0.46	0.58	0.08
Age (y)	15 \pm 3	13 \pm 2	15 \pm 3	<0.01	0.01	0.01	0.99
Trophic Position ($\delta^{15}\text{N}$, ‰)	10.2 \pm 1.1	11.6 \pm 1.8	13.0 \pm 1.7	< 0.01	<0.01	< 0.01	0.01
Le Cren Condition Factor (-)	0.996 \pm 0.022	1.00 \pm 0.02	1.00 \pm 0.02	0.38	0.40	0.97	0.54
Muscle Fat Content (%)	13.4 \pm 5.5	9.80 \pm 4.88	8.77 \pm 4.88	<0.01	0.03	0.72	<0.01

630

631 SD : Standard Deviation

632 NH: Narrow-head

633 INT: Intermediate

634 BH: Broad-head

635

636 **Table 2: Results obtained from the ANCOVA of the different pollutant concentrations**
637 **between narrow-heads, intermediates and broad-heads. The adjusted means for each**
638 **group are given, as well as the P-values for slope equality ($P > 0.05$ for ANCOVA) and the P-**
639 **values of the ANCOVA (P_{anc}). Significant values are shown in bold.**

		TotL (log-transformed)								
		Adj. Means			Slope			$P_{s/eq}$	P_{anc}	
		NH	INT	BH	NH	INT	BH			
	Hg	-1.02	-0.97	-0.81	-1.028	-0.451	0.471	0.62	< 0.01	
	Zn	1.00	1.01	1.04	-1.133	1.247	0.400	0.06	0.52	
	CB 101	2.62	2.77	2.59	1.730	0.775	0.710	0.91	0.07	
	CB 153	3.12	3.29	3.30	0.179	0.978	0.719	0.93	0.01	
	CB 138	3.05	3.23	3.24	-0.139	0.176	0.213	0.98	< 0.01	
	CB 180	2.62	2.80	2.86	-0.576	0.527	0.582	0.81	< 0.01	
	HCB	1.08	1.00	0.85	3.030	-0.092	0.731	0.26	< 0.01	
	p,p'-DDE	2.45	2.60	2.58	0.380	-0.007	0.300	0.97	0.03	
	α-HBCD	0.99	1.03	0.81	-3.444	-1.017	-1.394	0.73	0.08	
	∑PCB	3.58	3.74	3.72	0.383	0.518	0.639	0.99	0.02	
	∑BDE	1.13	1.27	1.09	1.442	0.581	0.619	0.92	0.03	
Wet Weight/ Lipid Weight (log-transformed)	∑DDT	2.53	2.68	2.64	0.597	-0.062	0.285	0.94	0.05	
		Age (not log-transformed)								
	Hg	-1.02	-0.96	-0.82	0.058	-0.015	-0.001	< 0.01	< 0.01	
	Zn	1	1.03	1.05	-0.001	-0.005	-0.002	0.97	0.29	
	CB 101	2.62	2.79	2.64	-0.044	0.009	0.021	0.10	0.18	
	CB 153	3.11	3.31	3.35	-0.044	0.011	0.007	0.04	< 0.01	
	CB 138	3.04	3.25	3.27	-0.039	0.015	0.011	0.04	< 0.01	
	CB 180	2.6	2.82	2.89	-0.028	0.009	0.007	0.24	< 0.01	
	HCB	1.07	1.01	0.85	0.003	0.018	0.023	0.77	0.01	
	p,p'-DDE	2.44	2.61	2.61	-0.031	0.008	0.015	0.11	0.01	
	α-HBCD	0.98	1.05	0.88	0.010	0.034	-0.001	0.70	0.33	
	∑PCB	3.57	3.75	3.76	-0.039	0.013	0.011	0.05	< 0.01	
	∑BDE	1.1	1.28	1.13	-0.021	0.016	0.000	0.47	0.07	
	∑DDT	2.52	2.69	2.67	-0.033	0.008	0.016	0.08	0.02	

640 TotL: Total Length

641 NH: Narrow-head

642 INT: Intermediate

643 BH: Broad-head

644

645 **Table 3. Results of the non-parametric permutation test (10,000 replicates) performed on**
646 **the log-transformed pollutant concentrations. Wet weight concentrations were used for**
647 **metals, lipid weight concentrations for Persistent Organic Pollutants. Significant differences**
648 **are shown in bold. * indicate pollutants for which the median is presented (as the measured**
649 **concentrations were not above detection limit in all eels).**

		Mean/Median \pm SD			P-values		
		NH	INT	BH	NH-INT	INT-BH	BH-NH
	As	-1.36 \pm 0.22	-1.49 \pm 0.42	-1.55 \pm 0.15	0.12	0.75	< 0.01
	Cu	0.92 \pm 0.11	-0.88 \pm 0.14	-0.77 \pm 0.25	0.24	0.07	< 0.01
	Pb	-1.81 \pm 0.40	-1.33 \pm 0.66	-1.55 \pm 0.67	< 0.01	0.26	0.10
	Se	-0.84 \pm 0.31	-1.00 \pm 0.62	-1.06 \pm 0.63	0.26	0.72	0.10
	Cr*	-1.56 \pm 0.49	-1.57 \pm 0.96	-2.23 \pm 0.92	0.24	0.12	< 0.01
Wet Weight/	OxC*	0.40 \pm 0.20	0.46 \pm 0.21	0.35 \pm 0.26	0.37	0.07	0.24
Lipid Weight	TN*	0.53 \pm 0.13	0.63 \pm 0.18	0.54 \pm 0.33	0.07	0.02	0.17
	<i>p,p'</i> -DDD	1.70 \pm 0.19	1.81 \pm 0.16	1.67 \pm 0.33	< 0.01	0.70	0.04
	<i>p,p'</i> -DDT*	0.77 \pm 0.20	0.88 \pm 0.21	0.91 \pm 0.30	< 0.01	0.69	0.04
	CB 28	1.81 \pm 0.22	1.84 \pm 0.19	1.53 \pm 0.46	0.55	< 0.01	< 0.01
	CB 52	2.60 \pm 0.20	2.68 \pm 0.17	2.47 \pm 0.40	0.14	0.02	0.15
	BDE47	0.94 \pm 0.21	1.09 \pm 0.20	0.85 \pm 0.40	0.01	< 0.01	0.31
	BDE100*	0.45 \pm 0.18	0.55 \pm 0.21	0.58 \pm 0.54	0.02	0.32	0.99

650

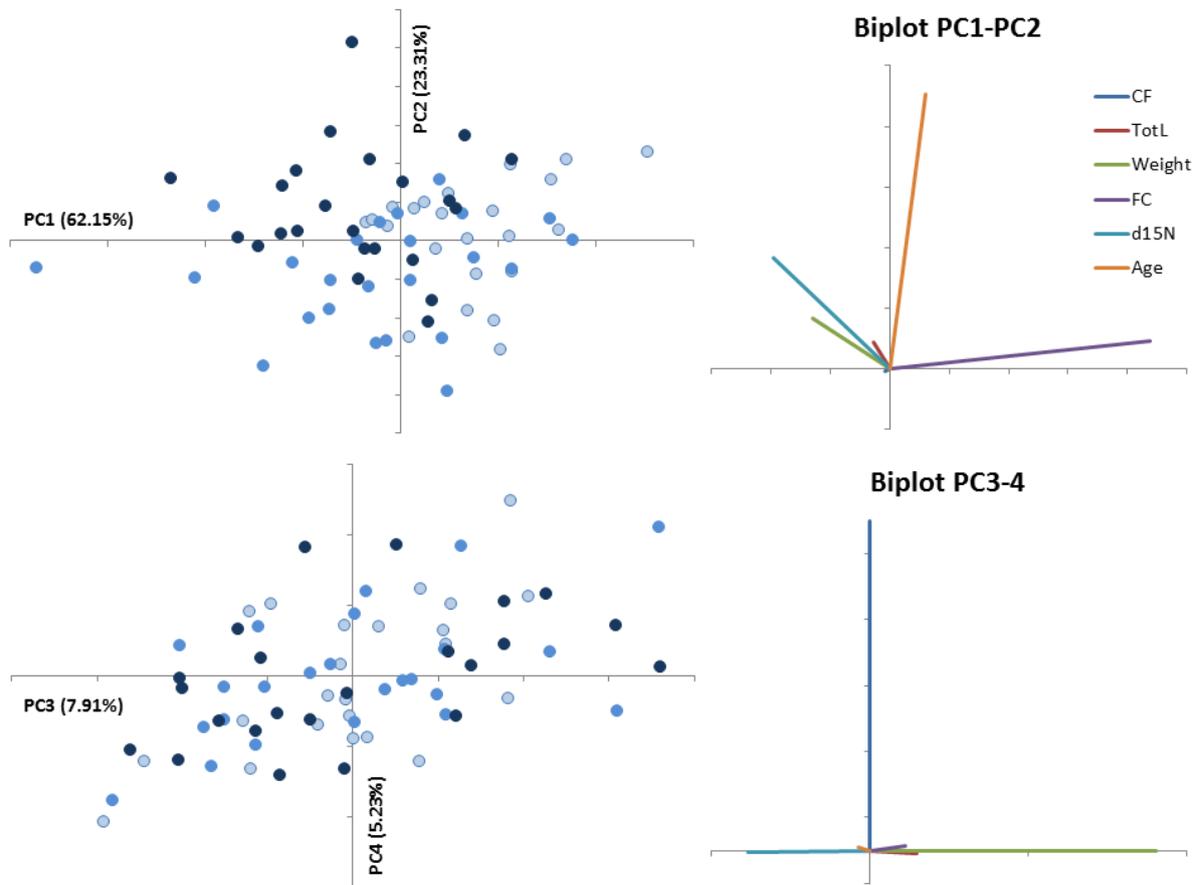
651 SD: Standard Deviation

652 NH: Narrow-head

653 INT: Intermediate

654 BH: Broad-head

655



657

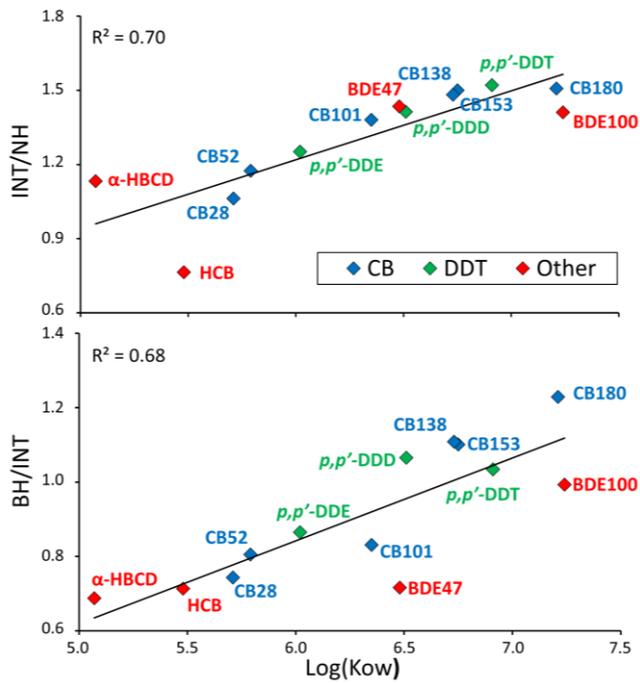
658 **Fig. S1: Between-group PCA on the different eel measurements and the associated biplot.**

659 The pollutant concentrations were expressed as lipid weight. Light blue dots represents

660 narrow-heads, blue dots intermediates and dark blue dots broad-headed eels. The amount of

661 variation explained by each Principal Component (PC) is shown in parentheses.

662



663

664 **Fig. S2: The mean(INT)/mean(NH) and mean(BH)/mean(INT) of the different pollutants**
 665 **plotted against the log of their octanol/water partition coefficient (Kow).** The pollutants are
 666 expressed as lipid weight. NH: Narrow-heads, INT: Intermediates, BH: Broad-heads. Blue
 667 diamonds represent pentachlorobiphenyls (PCBs), green diamonds
 668 Dichlorodiphenyltrichloroethanes (DDTs) and red the remaining measured pollutants.

669

Head: Age &
head shape

Body part 1:
Stable isotopes

Body part 2:
POPs

Body part 3:
Trace metals

Body part 4:
Reserve



670

671 **Fig. S3: The different eel body parts used for the different analyses.**

672

673 **SUPPLEMENTARY TABLES**

674 **Table S1: The absolute number of specimens for which the pollutant measurement was**
 675 **below the detection limit.** The relative number can be found in parentheses (expressed in
 676 %). Pollutants that were below the detection limit for more than 50% of the eels were not
 677 used for further analyses. NH: Narrow-heads, INT: Intermediates, BH: Broad-heads.

	NH	INT	BH	TOTAL
CB 28	0 (0)	0 (0)	0 (0)	0 (0)
CB 52	0 (0)	0 (0)	0 (0)	0 (0)
CB 101	0 (0)	0 (0)	0 (0)	0 (0)
CB 138	0 (0)	0 (0)	0 (0)	0 (0)
CB 153	0 (0)	0 (0)	0 (0)	0 (0)
CB 180	0 (0)	0 (0)	0 (0)	0 (0)
OxC	5 (19)	5 (20)	10 (42)	20 (27)
TN	0 (0)	1 (4)	7 (29)	8 (11)
CN	26 (100)	25 (100)	24 (100)	75 (100)
TC	26 (100)	25 (100)	24 (100)	75 (100)
CC	24 (92)	23 (92)	23 (96)	70 (93)
HCB	0 (0)	0 (0)	3 (13)	3 (4)
<i>p,p'</i>-DDE	0 (0)	0 (0)	0 (0)	0 (0)
<i>p,p'</i>-DDD	0 (0)	0 (0)	0 (0)	0 (0)
<i>p,p'</i>-DDT	7 (27)	2 (8)	6 (25)	15 (20)
BDE 28	24 (92)	20 (80)	21 (88)	65 (87)
BDE 47	0 (0)	0 (0)	0 (0)	0 (0)
BDE 100	0 (0)	0 (0)	1 (4)	1 (1)
BDE 99	23 (88)	25 (100)	23 (96)	71 (95)

BDE 154	18 (69)	8 (32)	15 (63)	41 (55)
BDE 153	19 (73)	12 (48)	17 (71)	48 (64)
BDE 183	26 (100)	25 (100)	24 (100)	75 (100)
s-DP	26 (100)	25 (100)	24 (100)	75 (100)
a-DP	26 (100)	25 (100)	22 (92)	73 (97)
α-HBCD	0 (0)	0 (0)	3 (13)	3 (4)
β-HBCD	25 (96)	24 (96)	24 (100)	73 (97)
γ-HBCD	22 (85)	24 (96)	24 (100)	70 (93)
As	0 (0)	0 (0)	0 (0)	0 (0)
Cd	25 (96)	25 (100)	24 (100)	74 (99)
Co	15 (58)	13 (52)	18 (75)	46 (61)
Cr	1 (4)	7 (28)	9 (38)	17 (23)
Cu	0 (0)	0 (0)	0 (0)	0 (0)
Hg	0 (0)	0 (0)	0 (0)	0 (0)
Pb	0 (0)	0 (0)	0 (0)	0 (0)
Ni	24 (92)	20 (80)	18 (75)	62 (83)
Se	0 (0)	0 (0)	0 (0)	0 (0)
Zn	0 (0)	0 (0)	0 (0)	0 (0)

678

679

680 **Table S2: Measurements of the different pollutants, expressed as lipid and wet weight.**

681 The median \pm Standard deviation (SD) is given for each phenotype; NH: Narrow-heads, INT:

682 Intermediates, BH: Broad-heads. Also the mean(INT)/mean(NH), mean(BH)/mean(INT) and

683 mean(BH)/mean(NH) ratio of each pollutant is given.

		Median \pm SD			Ratio		
		NH	INT	BH	INT/NH	BH/INT	BH/NH
Lipid weight concentration (ng/ g fat)	CB 28	68.6 \pm 27.8	72.0 \pm 25.4	54.0 \pm 64.3	1.06	0.743	0.790
	CB 52	413 \pm 180	480 \pm 195	415 \pm 297	1.17	0.805	0.946
	CB 101	401 \pm 285	620 \pm 300	507 \pm 405	1.38	0.831	1.15
	CB 138	967 \pm 605	1,808 \pm 782	1,926 \pm 1,071	1.50	1.10	1.65
	CB 153	1,150 \pm 807	2,139 \pm 966	2,336 \pm 1,251	1.48	1.11	1.64
	CB 180	381 \pm 234	721 \pm 301	771 \pm 432	1.51	1.23	1.85
	Sum PCB	3,367 \pm 2,013	5,626 \pm 2,353	6,763 \pm 3,287	1.44	1.06	1.52
	HCB	11.0 \pm 8.6	10.4 \pm 3.3	7.48 \pm 4.69	0.764	0.714	0.546
	α-HBCD	10.3 \pm 9.2	10.4 \pm 16.6	8.3 \pm 9.9	1.13	0.688	0.779
	TN	3.38 \pm 1.17	4.20 \pm 1.46	3.43 \pm 2.64	1.23	0.803	0.987
	OxC	2.50 \pm 0.97	2.89 \pm 1.13	2.25 \pm 1.50	1.14	0.808	0.919
	BDE 47	8.99 \pm 4.22	11.3 \pm 7.5	11.3 \pm 6.9	1.43	0.716	1.03
	BDE 100	2.83 \pm 1.35	3.56 \pm 2.74	3.80 \pm 3.63	1.41	0.993	1.4
	<i>p,p'</i>-DDE	262 \pm 122	421 \pm 152	461 \pm 244	1.41	1.07	1.51
	<i>p,p'</i>-DDD	49.0 \pm 24.4	67.4 \pm 27.1	61.3 \pm 34.1	1.25	0.865	1.08
<i>p,p'</i>-DDT	5.93 \pm 2.27	7.49 \pm 3.88	8.07 \pm 5.19	1.52	1.03	1.57	
Sum DDT	312 \pm 145	485 \pm 178	529 \pm 280	1.39	1.04	1.44	
Wet weight concentration (ng/g)	As	0.039 \pm 0.039	0.037 \pm 0.038	0.028 \pm 0.012	0.741	0.794	0.588
	Cr	0.028 \pm 0.025	0.027 \pm 0.065	0.006 \pm 0.064	1.48	0.633	0.924
	Cu	0.131 \pm 0.029	0.131 \pm 0.051	0.158 \pm 0.316	1.13	1.58	1.79
	Hg	0.084 \pm 0.064	0.095 \pm 0.049	0.154 \pm 0.044	1.06	1.39	1.48

Pb	0.013 ± 0.074	0.023 ± 0.178	0.015 ± 0.156	4.30	0.743	3.20
Se	0.123 ± 0.411	0.103 ± 0.622	0.098 ± 0.280	1.08	0.702	0.759
Zn	9.57 ± 2.6	10.4 ± 3.5	10.2 ± 2.7	1.04	1.06	1.10
CB 28	22.8 ± 13.8	20.7 ± 14.4	9.46 ± 19.7	-	-	-
CB 52	49.1 ± 26.4	41.9 ± 35.9	32.7 ± 38.3	-	-	-
CB 101	47.2 ± 34.1	57.9 ± 45.6	45.6 ± 46.8	-	-	-
CB 138	138 ± 89	140 ± 113	166 ± 102	-	-	-
CB 153	156 ± 118	168 ± 146	189 ± 130	-	-	-
CB 180	53.8 ± 40.1	51.0 ± 41.5	68.8 ± 36.5	-	-	-
Sum PCB	445 ± 301	497 ± 381	493 ± 352	-	-	-
HCB	1.33 ± 0.80	0.86 ± 0.57	0.47 ± 0.50	-	-	-
α-HBCD	1.23 ± 1.16	0.82 ± 2.00	0.51 ± 0.543	-	-	-
TN	0.454 ± 0.170	0.347 ± 0.215	0.273 ± 0.218	-	-	-
OxC	0.285 ± 0.149	0.248 ± 0.141	0.139 ± 0.116	-	-	-
BDE 47	1.14 ± 0.47	0.96 ± 1.65	0.58 ± 0.81	-	-	-
BDE 100	0.367 ± 0.172	0.284 ± 0.523	0.270 ± 0.300	-	-	-
<i>p,p'</i>-DDE	33.3 ± 20.9	32.0 ± 23.8	37.5 ± 24.3	-	-	-
<i>p,p'</i>-DDD	5.80 ± 4.53	5.09 ± 4.57	4.28 ± 3.84	-	-	-
<i>p,p'</i>-DDT	0.60 ± 0.36	0.59 ± 0.49	0.69 ± 0.42	-	-	-
Sum DDT	39.6 ± 25.4	37.0 ± 28.7	42.4 ± 28.3	-	-	-

684

685

686 **Table S3: The correlation of the log-transformed wet weight (metals) and lipid weight**
 687 **(POPs) concentrations of all specimens (n =75) with total length (TotL) and age.**

		TotL	Age
	As	-0.038	0.039
	Cr	-0.146	-0.161
	Cu	0.062	0.203
	Hg	-0.026	0.213
	Pb	0.129	-0.23
	Se	-0.1	-0.039
	Zn	0.118	-0.068
	CB 28	0.149	-0.055
	CB 52	0.144	-0.054
Wet Weight/	CB 101	0.133	-0.12
Lipid weight	CB 153	0.139	-0.15
(log-	CB 138	0.064	-0.11
transformed)	CB 180	0.096	-0.08
	OxC	-0.237	0.002
	TN	-0.141	-0.079
	HCB	0.064	0.12
	<i>p,p'</i> -DDE	0.065	-0.117
	<i>p,p'</i> -DDD	0.117	-0.119
	<i>p,p'</i> -DDT	-0.117	-0.046
	BDE47	0.145	-0.069
	BDE100	0.047	-0.121
	α -HBCD	-0.189	0.04

Σ PCB	0.109	-0.12
Σ BDE	0.111	-0.13
Σ DDT	0.062	-0.09

688

689

690 **Table S4: The P-values of the Shapiro-Wilk test.** Values lower than 0.05 indicate that the data
 691 did not fall within a normal distribution. * indicates *P*-values after removal of 1 outlier. Wet
 692 weight concentrations were used for the metals, lipid weight concentrations for the Persistent
 693 Organic Pollutants. NH: Narrow-heads, INT: Intermediates, BH: Broad-heads.

		NH	INT	BH	ALL
Original	TotL	0.51	0.09	0.04	-
	Weight	0.75	0.008	0.01	-
	Age	0.09	0.2	0.41	-
	Muscle Fat				
	content	< 0.01	0.47	0.07	-
	CF	0.64	0.45	0.28	-
	$\delta^{15}\text{N}$	0.13	0.35	0.15	-
Log- transformed	Totl	0.56	0.18	0.07	< 0.01
	Weight	0.96	0.06	0.30	0.08
	Age	0.04	0.07	0.19	0.15
	Muscle Fat Content	0.97	0.79*	0.98	0.03
Wet Weight/ Lipid weight (log- transformed)	As	< 0.01	< 0.01	0.47	< 0.01
	Cr	< 0.01	< 0.01	< 0.01	< 0.01
	Cu	0.02	0.03	< 0.01	< 0.01
	Hg	0.26*	0.75	0.28	< 0.01
	Pb	< 0.01	< 0.01	< 0.01	< 0.01
	Se	< 0.01	< 0.01	< 0.01	< 0.01
	Zn	0.51	0.18	0.52	< 0.01
	CB 28	0.85	0.82	0.03	< 0.01
	CB 52	0.16	0.93	0.03	< 0.01

CB 101	0.77	0.60	0.24	0.10
CB 153	0.39	0.21	0.33	0.62
CB 138	0.22	0.29	0.30	0.65
CB 180	0.65	0.36	0.10	0.42
OxC	< 0.01	< 0.01	< 0.01	< 0.01
TN	0.80	< 0.01	0.01	< 0.01
HCB	0.06	0.17	0.57*	< 0.01
<i>p,p'</i> -DDE	0.30	0.76	0.07	0.43
<i>p,p'</i> -DDD	0.53	0.35	0.03	< 0.01
<i>p,p'</i> -DDT	< 0.01	0.17	0.02	< 0.01
BDE47	0.35	0.59	0.02	< 0.01
BDE100	0.80	0.37	< 0.01	< 0.01
α -HBCD	0.73	0.10	0.19	0.23
Sum PCB	0.22	0.24	0.09	0.48
Sum BDE	0.79	0.40	0.18	0.27
Sum DDT	0.51	0.61	0.06	0.27

694

695

696 **Table S5: Results of the sensitivity analysis. The table shows the mean(INT)/mean(NH),**
 697 **mean(INT)/mean(BH) and mean(BH)/mean(NH) ratios for pollutants where concentrations**
 698 **below detection limit were measured and where the non-detects were replaced by the**
 699 **detection limit/2, the minimum concentration (0) and the minimum detection limit.**

		DL/2	Minimum	DL
BDE100	INT/NH	1.41	1.41	1.41
	BH/INT	0.99	0.99	0.99
	BH/NH	1.4	1.4	1.41
HBCD	INT/NH	1.13	1.13	1.13
	BH/INT	0.69	0.68	0.7
	BH/NH	0.78	0.77	0.79
HCB	INT/NH	0.76	0.76	0.76
	BH/INT	0.71	0.7	0.72
	BH/NH	0.55	0.54	0.55
DDT	INT/NH	1.52	1.6	1.39
	BH/INT	1.03	1.01	1.09
	BH/NH	1.57	1.61	1.51
TN	INT/NH	1.23	1.22	1.24
	BH/INT	0.8	0.75	0.86
	BH/NH	0.99	0.92	1.07

700