

1 **Stable isotopes and gut contents indicate differential resource use by coexisting asp**
2 **(*Leuciscus aspius*) and pikeperch (*Sander lucioperca*)**

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16 **Running title:** Resource use by coexisting asp and pikeperch

17

18 **Abstract**

19 Differential use of habitat and prey resources is an important mechanism that may allow
20 coexistence of potentially competing species. Unlike interactions between smaller cyprinid
21 and percid fishes, the resource use by coexisting predatory asp (*Leuciscus aspius*) and
22 pikeperch (*Sander lucioperca*) has remained little studied. Here, we used gut content and
23 stable isotope analyses to study ontogenetic dietary shifts and interspecific trophic niche
24 overlap between asp and pikeperch coexisting in two reservoirs. We hypothesised that both
25 species show an ontogenetic dietary shift from small invertebrates to large fish prey, but at the
26 same time use different prey resources to reduce potential competitive interactions. Our
27 results demonstrated clear ontogenetic dietary shifts by and interspecific niche segregation
28 between asp and pikeperch. The isotopic niches of the two predators showed no or only a
29 moderate degree of overlap (0–65%). Interestingly, the degree of interspecific isotopic niche
30 overlap changed with body size differently in the two reservoirs, suggesting that trophic
31 segregation can be dynamic and variable among systems. Gut contents revealed that small
32 (<100 mm standard length) asp consumed mostly terrestrial invertebrates and emerged
33 aquatic insects, whereas small pikeperch foraged on zooplankton, larval and pupal stages of
34 aquatic insects and fish. Larger individuals (>100 mm) of both species were predominantly
35 piscivorous, with asp feeding more on cyprinid prey fish and pikeperch feeding more on
36 percid prey fish. Our study indicates that coexisting asp and pikeperch populations can use
37 different prey resources, thereby reducing the potential negative competitive interactions.

38

39 **Keywords:** dietary ontogeny, foraging strategy, interspecific competition, piscivory, stable
40 isotopes

41

42 **1 Introduction**

43 Differential resource use is perceived as an important mechanism allowing the coexistence of
44 species within ecological communities (Schoener, 1986; Chesson, 2000; Chase & Leibold,
45 2003). This view is based on the competitive exclusion principle (Hardin, 1960), which states
46 that species cannot stably coexist unless the utilisation of limiting resources is well
47 differentiated. The segregation of coexisting species can occur along various dimensions such
48 as the time of activity, the habitat used, or the type of prey eaten (Schoener, 1986). A
49 comprehensive review of resource use in fish communities by Ross (1986) suggested that
50 niche segregation among coexisting species is mainly driven by partitioning of available food
51 resources rather than habitat or time segregation. Species coexistence can, however, be
52 influenced also by other mechanisms. For example, stochastic events (e.g., unpredictable
53 environmental fluctuations) that affect demographic attributes of species may result in their
54 coexistence (Sale, 1978; Warner & Chesson, 1985; Grossman et al., 1998). Therefore, one
55 approach to improve our understanding of the ecological mechanisms that determine the
56 coexistence of species is to examine resource overlap among potential competitors. Most
57 studies of dietary segregation between coexisting fish species have focused on adult life
58 stages (e.g., Hodgson et al., 1997; Schulze et al., 2012; Walker et al., 2013; Zaia Alves et al.,
59 2017). Ontogenetic variations in the resource use among potential competitors have been
60 examined less frequently (Werner & Gilliam, 1984; Amundsen et al., 2003; Davis et al.,
61 2012) although this knowledge is important to fully understand the structure and functioning
62 of fish communities.

63 Asp (Cyprinidae, *Leuciscus aspius*) and pikeperch (Percidae, *Sander lucioperca*) are
64 important piscivorous fishes in freshwater communities of western Eurasia (Kottelat &
65 Freyhof, 2007; Vašek et al., 2013). In their native range, these two species often coexist in

66 large rivers, lakes and reservoirs. Adults reach similar sizes (usually up to 1000 mm in total
67 length) and prey on small fish (Baruš & Oliva, 1995; Mittelbach & Persson, 1998). Juveniles
68 of pikeperch forage on aquatic invertebrates whereas juveniles of asp may also feed on
69 terrestrial insects fallen on the water surface (Baruš & Oliva, 1995). The feeding ecology of
70 pikeperch has been explored extensively and thus it is well-known that this species usually
71 shifts to piscivory in the first summer of its life (Buijse & Houthuijzen, 1992; van Densen et
72 al., 1996; Mittelbach & Persson, 1998). Less is known, however, about the size and age at
73 which asp become piscivorous. Moreover, only limited attempts have been made to
74 quantitatively characterise the diets of coexisting asp and pikeperch populations (Specziár &
75 Rezsű, 2009). In general, similar feeding habits (i.e., invertivory followed by piscivory)
76 suggest that the two species may interact strongly. Sympatric populations of asp and
77 pikeperch thus provide a good opportunity to investigate whether and how the two predators
78 differ in resource use throughout their lives.

79 In this study, gut content (GCA) and stable isotope (SIA) analyses were used to
80 explore ontogenetic dietary shifts and niche segregation between asp and pikeperch co-
81 occurring in two artificial lakes. It was expected that both species undergo an ontogenetic
82 dietary shift from invertebrates to fish prey, but this shift occurs later (i.e., at a larger body
83 size) for asp due to its higher tendency to feed on invertebrates. It was also hypothesised that
84 coexisting asp and pikeperch use different prey resources, but the degree of trophic
85 segregation diminishes with increasing body size, i.e. when both species become piscivorous.

86

87

88 **2 Methods**

89 2.1 Study sites

90 The study was carried out in two reservoirs located in South Bohemia, Czech Republic. Lipno
91 Reservoir (hereafter Lipno; dam coordinates: 48°37'58"N, 14°14'13"E), situated on the upper
92 Vltava River, is a relatively shallow water body (Table 1). Due to its shallowness and
93 frequent wind action, most of the reservoir area does not thermally stratify through the
94 growing season. In contrast, Římov Reservoir (hereafter Římov; dam coordinates:
95 48°51'00"N, 14°29'28"E), situated on the Malše River, is a deep canyon-type lake (Table 1)
96 that is strongly thermally stratified during the growing season. Both reservoirs have similar
97 water transparency and a moderately eutrophic trophic status (Table 1).

98 Due to seasonal water level fluctuations, the littoral zone vegetation is poorly
99 developed and submerged macrophytes are practically missing in both reservoirs. The adult
100 fish community compositions are similar in Lipno and Římov, with a dominance of cyprinid
101 species (mostly roach *Rutilus rutilus*, bleak *Alburnus alburnus*, bream *Abramis brama* and
102 white bream *Blicca bjoerkna*) accompanied by perch *Perca fluviatilis* and ruffe
103 *Gymnocephalus cernua*. Asp and pikeperch naturally reproduce in both reservoirs (Jůza et al.,
104 2013; Blabolil et al., 2016). In Římov, however, populations of the two predators are also
105 regularly supported by stocking with pond-reared fingerlings in autumn (Vašek et al., 2013).

106

107 2.2 Sample collection

108 Asp, pikeperch and their fish prey were sampled from Lipno in August/September 2012 and
109 2013, and from Římov in August 2013 and 2014. Sampling was carried out with multi-mesh
110 survey gillnets set overnight in littoral, profundal and pelagic zones at four to five different
111 stations within each reservoir (for details of the gillnet sampling, see Vašek et al., 2016).
112 Additional samples of young-of-the-year (YOY) asp and pikeperch, as well as prey fish, were

113 collected from the littoral and pelagic zones of both reservoirs using a beach seine net and a
114 trawl, respectively (for details of these sampling methods, see Jůza et al., 2014).

115 Each fish was measured for standard length (mm) and a sample of dorsal muscle was
116 dissected and stored at -20°C until processed for stable isotope analysis. The analysed prey
117 fish included YOY perch, ruffe and roach, and one-year old bleak. The digestive tracts of asp
118 and pikeperch were dissected and preserved in a 10% formaldehyde solution for later diet
119 analysis. Moreover, scales or otoliths were taken and used for age determination. To evaluate
120 ontogenetic changes in the short-term (based on GCA) and long-term (based on SIA) diets of
121 asp and pikeperch, individuals of both species were grouped into <100 , $100\text{--}199$, $200\text{--}299$,
122 and ≥ 300 mm size classes that corresponded approximately to age categories 0+, 1+, 2+, and
123 $\geq 3+$, respectively.

124 At both reservoirs, invertebrate samples for SIA were collected from three to four
125 sampling stations and three to four times (June to August) during the same summers when the
126 fish were captured. Bulk zooplankton was collected from the pelagic zone by taking several
127 vertical hauls through the upper 5 m (Lipno) or 10 m (Římov) of the water column with a
128 $200\text{-}\mu\text{m}$ mesh plankton net. The live zooplankton was immediately sieved through a $350\text{-}\mu\text{m}$
129 mesh and stored frozen at -20°C . Before preparation for SIA, defrosted samples were visually
130 inspected using a stereomicroscope. Most of the samples were dominated by herbivorous
131 crustaceans, mainly cladocerans (*Daphnia*) and calanoid copepods. Bulk samples of
132 macroinvertebrates from the littoral zone (<2 m depth) were collected with a kick net (mesh
133 size 0.7 mm) and hand-picked from stones. Only non-predatory organisms (primary
134 consumers) were considered and they included mainly trichopteran, ephemeropteran and
135 chironomid larvae, and also waterlouse (*Asellus aquaticus*) and small snails (Lymnaeidae).
136 All the trichopteran larvae and snails were removed from their cases or shells. Bulk samples

137 of terrestrial insects were collected with a sweep net from the shoreline grasses and shrubs.
138 Adults of the aquatic insects (Trichoptera, Ephemeroptera, Odonata and Chironomidae) were
139 excluded when present, and thus the samples contained adult insects of purely terrestrial
140 origin, i.e. Hymenoptera, Hemiptera, Coleoptera, Lepidoptera, Brachycera and Orthoptera.
141 All littoral and terrestrial invertebrate samples were stored frozen at -20°C until further
142 processed for SIA.

143

144 2.3 Gut content analysis

145 In the laboratory, digestive tracts were opened and the contents were examined under a
146 stereomicroscope. Since asp lacks a true stomach, the contents of the entire gut from the
147 oesophagus to the anus were analysed, whereas only stomach contents were analysed from
148 pikeperch. The total gut or stomach fullness was first visually estimated on a percentage scale
149 ranging from empty (0%) to full (100%). The prey items were identified to the lowest feasible
150 taxonomic level, and their contribution to the total gut or stomach fullness was then
151 determined by the indirect volumetric method (Hyslop 1980). In addition, the number of prey
152 fish individuals discernible to species level was recorded for each digestive tract. When
153 possible, characteristic remains (e.g., scales, pharyngeal arches, opercula and other bones of
154 fish prey, and head capsules, thoraxes, tail spines and other exoskeleton parts of invertebrate
155 prey) were used for identification of masticated and partially digested prey items. The prey
156 items were subsequently grouped into six categories: (1) crustacean zooplankton, (2) larval
157 and pupal stages of aquatic insects, (3) emerged aquatic insects, (4) terrestrial insects, (5)
158 cyprinid fish, and (6) percid fish.

159

160 2.4 Stable isotope analysis

161 Fish muscle and invertebrate samples were dried at 60°C for 48 h and ground to a fine powder
162 using either a porcelain mortar or a mixer mill MM 200 (Retsch GmbH, Haan, Germany).
163 Stable carbon and nitrogen isotopes and the element (C, N) composition of all samples were
164 measured using a Europa Scientific elemental analyser interfaced with a Europa Scientific 20-
165 20 isotope ratio mass spectrometer (Sercon Ltd, Crewe, UK) at the Iso-Analytical Ltd, Crewe,
166 UK. Vienna Pee Dee Belemnite and atmospheric N₂ were used as the international standards
167 for carbon and nitrogen, respectively, while NBS-1577B (powdered bovine liver, $\delta^{13}\text{C}_{\text{V-PDB}} =$
168 -21.60‰ , $\delta^{15}\text{N}_{\text{Air}} = 7.65\text{‰}$) was used as a working standard. NBS-1557B was calibrated in-
169 house as a secondary reference material and is directly traceable to IAEA-CH-6 (sucrose,
170 $\delta^{13}\text{C}_{\text{V-PDB}} = -10.43\text{‰}$) and IAEA-N-1 (ammonium sulphate, $\delta^{15}\text{N}_{\text{Air}} = 0.40\text{‰}$). Isotope ratios
171 in each sample were expressed in conventional delta notation ($\delta^{13}\text{C}$, $\delta^{15}\text{N}$) as parts per
172 thousand (‰) differences from the international standard. The analytical error (standard
173 deviation), estimated from replicated runs of the reference material, was less than 0.1‰ for
174 both $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$. Every fifth sample was run in duplicate and the mean difference \pm
175 standard deviation (SD) between replicates was $0.03 \pm 0.04\text{‰}$ for $\delta^{13}\text{C}$ and $0.06 \pm 0.06\text{‰}$ for
176 $\delta^{15}\text{N}$. The fish muscle $\delta^{13}\text{C}$ values were not corrected for lipids due to the generally low C:N
177 ratios (<3.5) indicating negligible lipid content in the samples (Hoffman et al., 2015).

178 The relative contributions of different diet sources assimilated by each size class of
179 asp and pikeperch were modelled using the SIAR package in R (Stable Isotope Analysis in R;
180 Parnell et al., 2010). Inputs to the model were the $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ values of the individual
181 consumers (asp and pikeperch) and the reservoir-specific mean \pm SD $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ values of
182 the potential prey resources (see Appendix Figure). In both reservoirs, pelagic zooplankton
183 and littoral macroinvertebrates did not differ in isotope values (t -tests, $P > 0.05$ for both $\delta^{13}\text{C}$
184 and $\delta^{15}\text{N}$) and hence they were grouped as “aquatic invertebrates” for SIAR. Furthermore, the

185 isotope data collected in the subsequent years were pooled because isotope values for major
186 trophic level groups (i.e. predatory fish, prey fish, aquatic invertebrates and terrestrial insects)
187 did not substantially differ between years. Therefore, diet composition for both asp and
188 pikeperch was estimated from three possible diet sources: aquatic invertebrates, terrestrial
189 insects, and fish. Fractionation factors (mean \pm SD) between resources and the consumers
190 were assumed to be $0.91 \pm 1.04\text{‰}$ for $\delta^{13}\text{C}$ and $3.23 \pm 0.41\text{‰}$ for $\delta^{15}\text{N}$ (Vander Zanden &
191 Rasmussen, 2001). Element concentrations (proportions of C and N) directly measured in the
192 prey resources were included into the model (Phillips & Koch, 2002).

193 Trophic position (TP) of individual asp and pikeperch was estimated from stable
194 isotope data, using the equation described by Cabana and Rasmussen (1996):

$$195 \text{TP}_{\text{consumer}} = (\delta^{15}\text{N}_{\text{consumer}} - \delta^{15}\text{N}_{\text{baseline}}) / 3.23 + 2$$

196 where $\delta^{15}\text{N}_{\text{consumer}}$ is the $\delta^{15}\text{N}$ value of asp or pikeperch, $\delta^{15}\text{N}_{\text{baseline}}$ is the $\delta^{15}\text{N}$ value of the
197 baseline organisms (calculated as the average value from aquatic invertebrates), 3.23 is the
198 assumed diet-tissue enrichment in $\delta^{15}\text{N}$ per trophic level (Vander Zanden & Rasmussen,
199 2001), and the constant 2 refers to the trophic position of the baseline organisms.

200 Finally, the isotopic niche widths of each size class of asp and pikeperch were
201 calculated as sample size-corrected standard ellipse areas (SEA_{C}) using the SIBER package in
202 R (Stable Isotope Bayesian Ellipses in R; Jackson et al., 2011). SEA_{C} was also used to
203 determine the degree of isotopic niche overlap between the two species, using the equation of
204 Stasko et al. (2015):

$$205 \% \text{ Overlap} = [\{ (\text{area of overlap between } \text{SEA}_{\text{C}1} \text{ and } \text{SEA}_{\text{C}2}) \times 2 \} / (\text{SEA}_{\text{C}1} + \text{SEA}_{\text{C}2})] \times 100$$

206 where $\text{SEA}_{\text{C}1}$ and $\text{SEA}_{\text{C}2}$ are the ellipse areas calculated from asp and pikeperch samples,
207 respectively.

208

209 2.5 Statistical analysis

210 Non-parametric one-way analysis of similarities (ANOSIM) was run in PAST ver. 3.19
211 (Hammer et al., 2001) to compare volumetric proportions of different prey categories in the
212 digestive tracts of different size classes of asp and pikeperch in the Lipno and Římov
213 reservoirs. ANOSIM was based on Bray-Curtis similarity index and the one-tailed
214 significance was computed by permutation of group membership with 9,999 replicates. The
215 size at piscivorous shift was compared between the species by using binomial data of prey
216 fish presence in gut contents (0 = no fish remains in gut, 1 = fish remains in gut) as the
217 response variable and fish length and species as the predictor variables in logit-regression
218 models. Furthermore, the ontogenetic (i.e., size-related) changes in asp and pikeperch TP
219 were analysed by fitting asymptotic regression models using the SSasypm function in R (Ritz
220 et al., 2015). The differences in TP between asp and pikeperch of each size class in each
221 reservoir were also compared using *t*-test. Finally, the likelihood test in the SIBER (Jackson et
222 al., 2011) was used to test for between-species differences in isotopic niche widths of asp and
223 pikeperch size-classes. All statistical analyses except ANOSIM were performed in the R
224 computing programme ver. 3.4.1 (R Core Team, 2017).

225

226

227 **3 Results**

228 Both GCA and SIA data demonstrated clear ontogenetic dietary shifts and differential use of
229 the prey resources by coexisting asp and pikeperch. The GCA results indicated significant
230 between-species differences in the prey compositions (ANOSIM: $R = 0.457$, $P < 0.001$), but
231 the diets of asp and pikeperch became more similar with increasing size (Table 2). Small

232 (<100 mm) asp fed on terrestrial and emerged aquatic insects, whereas larger asp consumed
233 mostly fish (Fig. 1). Correspondingly, small (<100 mm) pikeperch foraged on zooplankton,
234 larval and pupal stages of aquatic insects and fish, whereas larger pikeperch were mainly
235 piscivores (Fig. 1). Contrary to asp, no terrestrial insects or emerged aquatic insects were
236 found in pikeperch stomachs. The two species showed contrasting prey fish compositions,
237 with asp feeding more often on cyprinid prey fish and pikeperch feeding mostly on percid fish
238 (Fig. 1). When only prey fish discernible to species level were considered, the most abundant
239 species found in asp guts were ruffe in Lipno and bleak in Římov, whereas the most abundant
240 species observed in pikeperch stomachs were perch and conspecifics in Lipno and ruffe in
241 Římov (Table 3). The logit-regression models (Fig. 2) demonstrated that pikeperch shifted to
242 piscivory at a smaller size than asp, both in Lipno (parameter estimate \pm SE for species effect:
243 2.4 ± 0.9 ; $Z = 2.6$, $P = 0.009$) and Římov (2.3 ± 0.6 ; $Z = 3.7$, $P < 0.001$).

244 The SIA results confirmed the ontogenetic dietary shifts of asp and pikeperch to
245 piscivory, as illustrated by the positive non-linear relationship between size and TP (Fig. 3,
246 Table 4) and by the SIAR estimates showing a shift from invertebrate to fish prey with
247 increasing predator size (Fig. 4). In Lipno, small (<100 mm) asp had a significantly lower TP
248 than similar-sized pikeperch, whereas no between-species differences were observed among
249 larger size classes (Table 5). An opposite pattern was observed in Římov, where no between-
250 species differences in TP were observed for small asp and pikeperch, whereas larger asp had
251 consistently lower TP as compared to similar-sized pikeperch (Table 5). The results from
252 SIAR isotopic mixing model suggested that terrestrial insects contributed only little to the
253 long-term diet of all size classes of asp and pikeperch in Římov but had a relatively high
254 contribution to the long-term diet of both small (<100 mm) asp (54%) and pikeperch (32%) in
255 Lipno (Fig. 4).

256 The isotopic niche widths generally did not differ between the coexisting asp and
257 pikeperch populations (Table 5). In Lipno, there was no overlap between isotopic niches
258 (SEAC) of small (<100 mm) asp and pikeperch, whereas the isotopic niches of larger asp and
259 pikeperch overlapped partially (i.e., 49–65%; Table 5). An opposite pattern was observed in
260 Římov, where the interspecific SEAC overlap was highest (65%) in the <100 mm size class,
261 intermediate (42%) in the 100–199 mm size class, and none in the 200–299 and \geq 300 mm size
262 classes (Table 5).

263

264

265 **4 Discussion**

266 Our results demonstrated clear ontogenetic dietary shifts and interspecific niche segregation
267 between asp and pikeperch. So far, only limited information on resource use has been
268 available for co-occurring asp and pikeperch populations (Specziár & Rezsú, 2009), and the
269 current study is also the first that applied SIA approach (together with conventional GCA
270 method) to characterise dietary niches of the two sympatric predators. Consequently, findings
271 of this study provide important insights into the trophic ecology of coexisting asp and
272 pikeperch populations and their roles in freshwater food webs.

273

274 4.1 Ontogenetic dietary shifts in asp and pikeperch

275 Gut content and stable isotope data both showed that asp and pikeperch consumed more fish
276 prey with increasing body size. According to the GCA and SIAR results, fish prey
277 overwhelmingly dominated in the short- and long-term diets of large- and medium-sized
278 predators, whereas they contributed only around 50% or less to the diets of small-sized (<100
279 mm) asp and pikeperch. Our results demonstrate that asp and pikeperch can begin feeding on

280 fish as early as their first summer, although shifting to piscivory was completed in their
281 second summer of life (i.e., at length >100 mm). Both the logit-regression models (based on
282 absence/presence of prey fish in predators' digestive tracts) and SIA-based TP estimates
283 consistently indicated that pikeperch shifted to piscivory at a smaller size than asp. These
284 results confirmed our expectation that juvenile asp have a higher tendency to feed on
285 invertebrates and shift to piscivory somewhat later (i.e., at a larger size) than pikeperch.
286 However, although piscivory occurred later for asp, the TP estimates suggest that both species
287 accomplished shifting to predominantly piscivorous feeding in their second summer of life
288 since individuals of the 100–199 mm size class attained mean TP values of ≥ 3.5 , indicating
289 piscivory. Consequently, both species can be characterised as “specialist piscivores” (sensu
290 Keast, 1985) because they shift to piscivory relatively early in life.

291 GCA indicated that small-sized (<100 mm) asp consumed mainly terrestrial and
292 emerged aquatic insects and to a lesser extent fish in both reservoirs. SIAR results suggested
293 that terrestrial insects dominated (54%) the long-term diet of small-sized (<100 mm) asp in
294 Lipno, whereas small asp in Římov showed a greater reliance upon prey fish. Hence, our
295 SIAR results indicate that small-sized Římov asp probably consumed more fish than
296 suggested by GCA which reflects only recently ingested prey items (e.g., Paradis et al., 2008).
297 According to GCA, small-sized (<100 mm) pikeperch fed on zooplankton, aquatic insects and
298 fish, but not on terrestrial and emerged aquatic insects. Although terrestrial insects were
299 absent in pikeperch digestive tracts, we always included terrestrial invertebrates as a third
300 prey source (besides aquatic invertebrates and fish) to make the SIAR analysis consistent
301 between asp and pikeperch. Consequently, the SIAR results confirmed that terrestrial insects
302 generally represented an unimportant prey source for pikeperch of all size classes. Small-sized
303 (<100 mm) Lipno pikeperch were, however, an exception because the SIAR suggested that

304 terrestrial insects might be a substantial diet source (32%) for these fish. This apparent bias
305 (i.e., overestimation of the terrestrial prey contribution to pikeperch diet) can be explained by
306 the fact that stable isotope values of terrestrial and aquatic invertebrates partially overlapped
307 (particularly in terms of $\delta^{13}\text{C}$) in Lipno, hindering assessment of the relative contributions of
308 these prey sources to higher trophic levels. Nevertheless, the SIAR results appropriately
309 revealed the ontogenetic niche shift from feeding on invertebrates to piscivory in both species.

310 Previous studies have shown that, under favourable growth conditions (i.e., high
311 temperature and food availability), pikeperch become piscivorous during their first summer
312 and reach sizes well above 100 mm (Buijse & Houthuijzen, 1992; van Densen et al., 1996). In
313 contrast, under less suitable conditions, YOY pikeperch either remain invertivorous and reach
314 generally small sizes (Specziár, 2005; Vinni et al., 2009, Ginter et al., 2011) or develop a
315 bimodal size distribution with a minor group becoming piscivorous and a majority staying
316 invertivorous (van Densen, 1985; Frankiewicz et al., 1996). Information on ontogenetic
317 dietary shifts in asp is limited. Yet, the data available from Lake Balaton (Specziár & Rezsú,
318 2009) correspond well with our results: the <40 mm asp were invertivorous, the 41–120 mm
319 asp had a mixed diet containing both invertebrates and fish, and the 121–500 mm fish were
320 entirely piscivorous. The fact that transition to piscivory in our study systems was not
321 completed during the first summer implies growth-limiting conditions for juvenile stages of
322 both species. Persson & Brönmark (2002) highlighted the importance for YOY predators to
323 be synchronised with fluctuations in resource availability. Hence, we can speculate that
324 discontinuous availability of suitable food resources might reduce growth and delay shifting
325 to piscivory in our study systems. However, growth rates of YOY predators might have also
326 been restricted by water temperatures. Římov is a deep reservoir situated in a canyon and
327 hence it warms slowly in spring which may delay the spawning period and shorten the first-

328 year growth season (cf., Wysujack et al., 2002; Jůza et al., 2013). In contrast, Lipno is a
329 shallow reservoir, but because of its location at a higher altitude, thermal conditions may
330 delay spawning and reduce the growth of juvenile stages similarly to that in Římov.

331

332 4.2 Interspecific niche segregation between asp and pikeperch

333 The isotopic niche widths generally did not differ between the coexisting asp and pikeperch
334 populations, indicating a similar extent of trophic specialisation in both species. Moreover,
335 the isotopic niches of the two predators showed no or only a moderate degree of overlap (i.e.,
336 0–65%). Interestingly, overlap between the isotopic niches of asp and pikeperch increased
337 with increasing predator size in Lipno but decreased in Římov. These findings do not support
338 our hypothesis that the degree of trophic segregation between asp and pikeperch should
339 diminish with increasing body size (i.e., with a shift to piscivory). Instead, the results suggest
340 that size-related trophic segregation between asp and pikeperch may be dynamic and variable
341 among systems, probably reflecting varying availability of prey sources.

342 The results also demonstrate that coexisting asp and pikeperch used rather different
343 prey resources both at small and large sizes. The GCA and SIA data both indicated that the
344 interspecific trophic segregation in the smallest (<100 mm) predator size class was likely due
345 to the exclusive utilisation of terrestrial invertebrates and emerged aquatic insects by asp,
346 whereas pikeperch used zooplankton, larval and pupal stages of aquatic insects, and small
347 fish. Similarly, Specziár & Rezsú (2009) observed that small (16–40 mm) asp foraged mostly
348 on adult Chironomidae, whereas co-occurring similar-sized pikeperch relied on zooplankton.
349 Moreover, our GCA results showed that piscivorous stages of the two predators consumed the
350 same fish species, but in different proportions. The fact that asp used relatively more cyprinid
351 prey fish while pikeperch consumed more percoid fish might be another reason for the

352 observed segregation of the isotopic niches of the two predators, particularly those of
353 medium- and large-sized classes. Because pikeperch ingest prey intact, with no mastication, it
354 was usually possible to identify (at least to family level) most of prey fish. In contrast,
355 ingested prey fish in asp were often strongly masticated by pharyngeal teeth and digested
356 beyond recognition. Hence, we suppose that the relative contribution of cyprinid prey fish in
357 the diet of asp might even be higher than suggested by the GCA, because small and soft
358 cyprinid species such as bleak were probably under-represented due to their rapid digestion.

359 Empirical data on piscivorous diets of coexisting asp and pikeperch populations have
360 been missing so far. Single-species studies indicate that the piscivorous diets of both asp and
361 pikeperch are dominated by cyprinid (Wysujack et al., 2002; Krpo-Ćetković et al., 2010;
362 Specziár, 2011) and by percid prey fish (Vostradovský & Váša, 1981; Frankiewicz et al.,
363 1999, Keskinen & Marjomäki, 2004). Hence, both predators can behave rather
364 opportunistically and consume the most abundant fish species. However, in sympatry, asp and
365 pikeperch can differentiate prey fish resources as illustrated by our study. In summary, by
366 using a combination of GCA and SIA, our study indicates that coexisting asp and pikeperch
367 populations can use different prey resources at both juvenile and adult life stages, thereby
368 reducing the potential negative competitive interactions (Vanni et al., 2009).

369

370 4.3 Conclusion and recommendation for future studies

371 The present study demonstrates that coexisting asp and pikeperch forage at the top of the food
372 webs and thereby play similar functional roles in lake ecosystems. Notably, our study showed
373 that the trophic niches of the two predators were relatively well separated, both at juvenile and
374 older life stages. The observed niche segregation may help to reduce potential interspecific
375 resource competition between coexisting asp and pikeperch populations.

376 Asp and pikeperch can induce top-down cascading impacts on lower trophic levels
377 (Benndorf, 1990; Brabrand & Faafeng, 1993; Donabaum et al., 1999). They are also popular
378 game fishes for anglers and therefore often stocked into various systems (e.g., Ruuhijärvi et
379 al., 1996; Wysujack et al., 2002; Vašek et al., 2013). Our results are relevant to fisheries
380 management, because they indicate that different use of the prey resources may potentially
381 mitigate interspecific competition between co-occurring asp and pikeperch populations. In
382 future studies, comparison of trophic niches of the two species under conditions of allopatry
383 and sympatry may help to determine whether relatively low overlap in resource use is the
384 consequence of interspecific competition or different foraging strategies that evolved in the
385 past.

386

387

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397

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- 549

550 **Tables**

551 **Table 1.** Basic environmental characteristics of the two reservoirs studied. Mean values for
552 the growing season (May–September) are shown for Secchi depth, total phosphorus and
553 chlorophyll-*a*.

Characteristic	Lipno	Římov
Year of filling	1960	1978
Surface altitude (m a.s.l.)	725	471
Surface area (km ²)	48.7	2.1
Mean depth (m)	6	16
Maximum depth (m)	22	43
Hydraulic retention time (days)	244	85
Secchi depth (m)	1.9	2.6
Total phosphorus (µg L ⁻¹)	25	27
Chlorophyll- <i>a</i> (µg L ⁻¹)	14	19

554

555 **Table 2.** Sample sizes in GCA (n) and results from pairwise one-way analysis of similarities
 556 (ANOSIM) comparisons of volumetric prey proportions in digestive tracts of asp and
 557 pikeperch in the Lipno and Římov reservoirs. Statistically significant differences ($P < 0.05$)
 558 are shown in bold.

Reservoir	Size class (mm)	n		ANOSIM
		Asp	Pikeperch	P
Lipno	<100	16	17	<0.001
	100-199	4	7	0.024
	200-299	6	11	0.262
	≥ 300	7	6	0.217
Římov	<100	12	20	<0.001
	100-199	14	10	0.043
	200-299	15	11	<0.001
	≥ 300	7	5	0.176

559

560

561 **Table 3.** List of fish species preyed upon and their total numbers found in all digestive tracts
 562 of asp and pikeperch collected from the Lipno and Římov reservoirs.

Prey fish species (family)	Asp		Pikeperch	
	Lipno	Římov	Lipno	Římov
Perch (Percidae)	3	2	14	8
Pikeperch (Percidae)	1	-	11	2
Ruffe (Percidae)	7	-	6	13
Bleak (Cyprinidae)	3	9	1	-
Bream (Cyprinidae)	-	-	-	1
Roach (Cyprinidae)	1	1	-	6
Total number of prey fish discernible to species level	15	12	32	30

563

564 **Table 4.** Parameter estimates and corresponding *t*- and *P*-values for the non-linear
565 (asymptotic) regression models with trophic position (TP) as a response variable and standard
566 length (mm) as a predictor variable, given for asp and pikeperch in Lipno and Římov
567 reservoirs. The models are fitted using SSasympt function in R (R Core Team 2017),
568 producing estimates for the horizontal asymptote on the right side (Asym), the response value
569 (i.e., TP) when length is zero (R0), and the natural logarithm of the rate constant (lrc). See
570 Fig. 3 for sample sizes and predicted regression curves.

Reservoir	Species	Parameter	Estimate	SE	<i>t</i>	<i>P</i>
Lipno	Asp	Asym	3.85	0.08	45.79	<0.001
		R0	0.22	0.88	0.25	0.806
		lrc	-4.29	0.27	-16.00	<0.001
Lipno	Pikeperch	Asym	3.77	0.05	81.09	<0.001
		R0	2.03	0.26	7.93	<0.001
		lrc	-4.16	0.20	-21.28	<0.001
Římov	Asp	Asym	3.91	0.07	54.77	<0.001
		R0	2.50	0.39	6.35	<0.001
		lrc	-4.45	0.35	-12.90	<0.001
Římov	Pikeperch	Asym	4.31	0.08	51.30	<0.001
		R0	2.86	0.18	16.07	<0.001
		lrc	-4.66	0.24	-19.16	<0.001

571

572 **Table 5.** Mean trophic position (TP; standard deviation in parentheses), isotopic niche width
573 (SEAC) and niche overlap for asp and pikeperch of four size classes collected from the Lipno
574 and Římov reservoirs. P^\dagger and P^\ddagger refer to statistical significances for TP comparisons and for
575 niche size comparisons, respectively, between asp and pikeperch in each size class.
576 Statistically significant differences ($P < 0.05$) are highlighted in bold. For number of
577 predators analysed for stable isotopes in each size class see Fig. 4.

Reservoir	Size class (mm)	TP (SD)		P^\dagger	SEAC		P^\ddagger	Overlap (%)
		Asp	Pikeperch		Asp	Pikeperch		
Lipno	<100	2.5 (0.3)	3.1 (0.2)	<0.001	2.6	3.9	0.88	0.0
	100-199	3.5 (0.2)	3.6 (0.3)	0.64	3.0	2.9	0.38	48.8
	200-299	3.7 (0.3)	3.7 (0.2)	0.94	3.2	1.3	0.03	54.9
	≥ 300	3.8 (0.2)	3.8 (0.2)	0.64	2.1	1.1	0.08	64.7
Římov	<100	3.3 (0.3)	3.5 (0.3)	0.07	2.7	2.7	0.51	65.2
	100-199	3.7 (0.2)	3.9 (0.1)	<0.001	1.0	1.2	0.74	42.3
	200-299	3.8 (0.1)	4.2 (0.2)	<0.001	1.2	1.1	0.28	0.0
	≥ 300	3.9 (0.2)	4.2 (0.2)	<0.001	1.4	0.9	0.18	0.0

578

579 **Figure Legends**

580 **Figure 1.** Mean volumetric proportion of different prey categories in the digestive tracts of
581 different size classes of asp and pikeperch in the Lipno and Římov reservoirs. Predators that
582 contained unidentified prey fish in their digestive tracts are not shown to better illustrate
583 interspecific differences in piscivorous foraging on percid and cyprinid fish. The number of
584 examined digestive tracts with discernible prey items is indicated above the bars.

585

586 **Figure 2.** Logistic regressions showing the probability of finding fish remains in gut contents
587 as a function of asp ($n = 43$ & 65) and pikeperch ($n = 56$ & 69) standard length in the Lipno
588 and Římov reservoirs.

589

590 **Figure 3.** Trophic positions of asp ($n = 43$ & 77) and pikeperch ($n = 62$ & 88) in the Lipno
591 and Římov reservoirs as a function of standard length. Lines indicate predicted values for the
592 non-linear (asymptotic) regression models (see parameter estimates in Table 4).

593

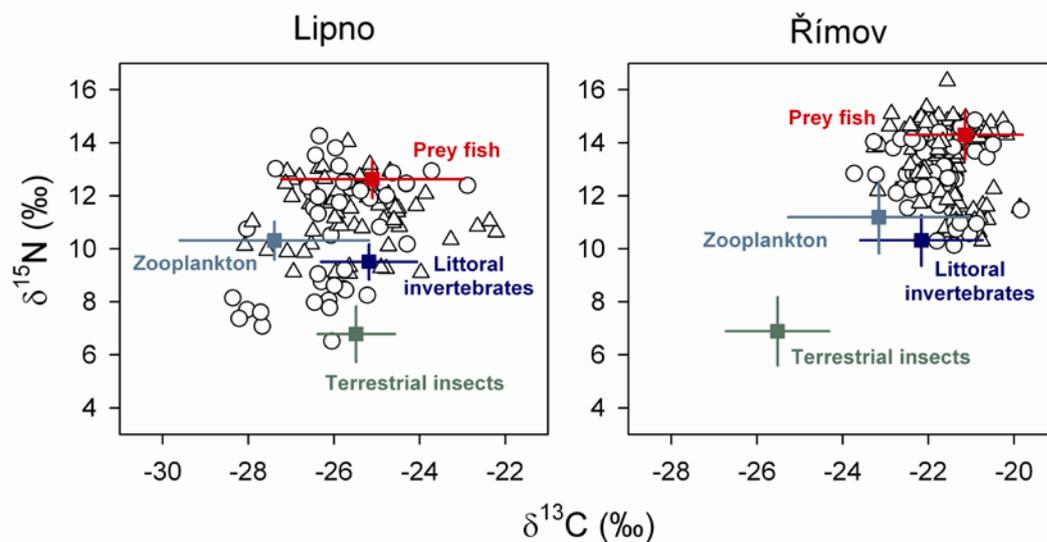
594 **Figure 4.** Proportional contribution (mean \pm 95% credibility intervals) of different prey types
595 (aquatic invertebrates, terrestrial insects and fish) to the diet of different size classes of asp
596 and pikeperch in the Lipno and Římov reservoirs, estimated using SIAR mixing model. The
597 number of predators analysed for stable isotopes in each size class is indicated across the top
598 of the graphs.

599

600 **Appendix**

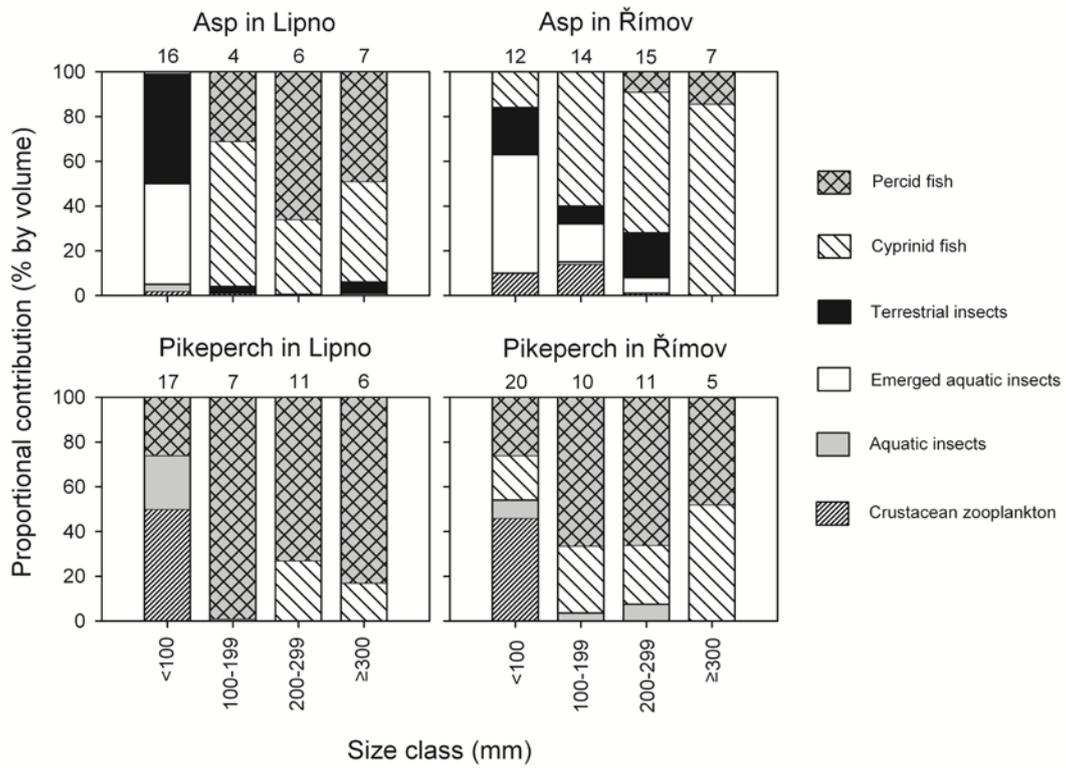
601 **Appendix figure**

602 Bi-plots of $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ values for individual asp (circles) and pikeperch (triangles), and
603 their principal diet sources in the Lipno and Římov reservoirs. Filled squares represent mean
604 \pm standard deviation for pelagic zooplankton, littoral macroinvertebrates, terrestrial insects
605 and prey fish. All diet sources were corrected for trophic fractionation using values ($\delta^{13}\text{C} =$
606 0.91 , $\delta^{15}\text{N} = 3.23$) from Vander Zanden & Rasmussen (2001). Because pelagic zooplankton
607 and littoral macroinvertebrates did not differ in their isotope values, they were merged as
608 “aquatic invertebrates” for the purpose of SIAR estimates (see Fig. 4).

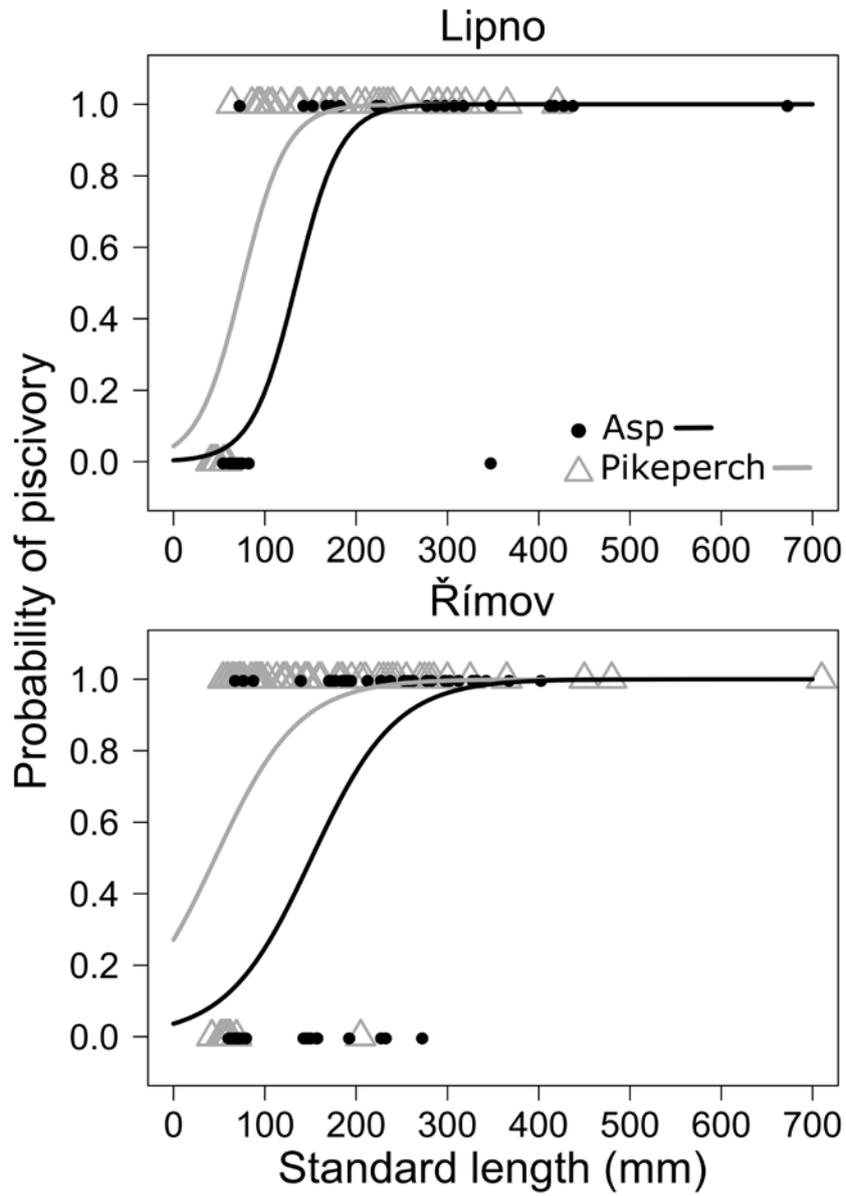


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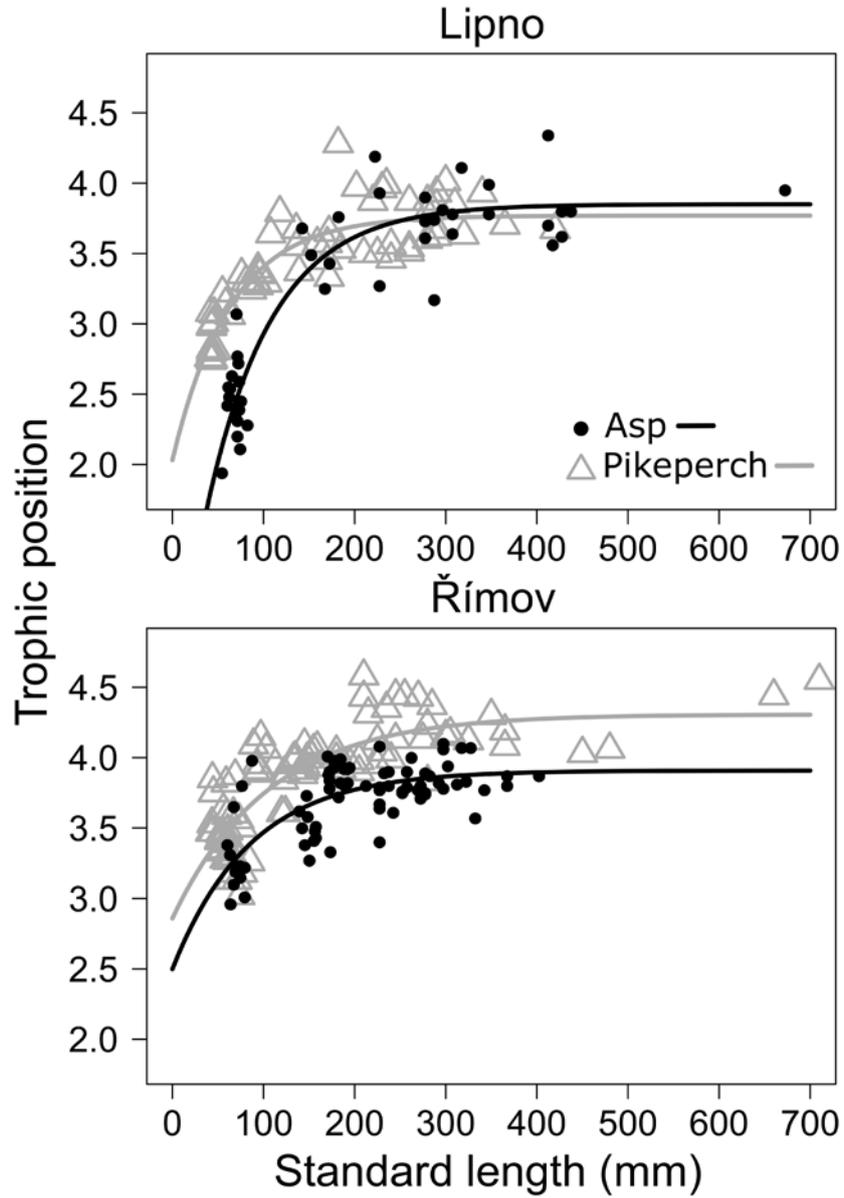
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