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1	Narrower isotopic niche size in fish infected by the intestinal
2	parasite Pomphorhynchus sp. compared to uninfected ones
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#### 22 Abstract:

Examples of parasite-related effects on intermediate crustacean hosts are numerous but their 23 ecological consequences on their vertebrate hosts are scarce. Here, we address the role of 24 macroparasite infections on the trophic niche structure of definitive hosts and its potential 25 physiological consequences using wild fish populations infected with an acantochephalan 26 27 parasite Pomphorhynchus sp., a trophically-transmitted intestinal worm. Infected and 28 uninfected fish were sampled from six populations on the Marne River, France and the prevalence of intestinal parasites in the host populations ranged from 50% to 90%. Although 29 the isotopic ratios ( $\delta^{13}$ C and  $\delta^{15}$ N) did not differ between infected and uninfected fish, we 30 found a consistent pattern of isotopic niche size being considerably smaller in infected hosts 31 32 when compared with non-infected ones. This was not explained by interindividual differences in intrinsic factors such as length/age or body condition between infected and uninfected fish. 33 These results suggest a potential niche specialization of infected fish, which did not impair 34 35 their energetic status.

Keywords: Body condition, Carbon, European chub, Host-parasite system, Intestinal worms,
Nitrogen

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## 44 **1. Introduction**

Parasites are ubiquitous components of biological systems, found virtually in all freeliving organisms on earth (Strona, 2015). By definition, parasites live at the expense of their hosts and this interdependence of parasites is known to cause considerable changes in the physiology and metabolism of hosts (Shinagawa *et al.*, 2001; Hicks *et al.*, 2018), leading to potential reduced outcomes in host individuals (Johnson and Dick, 2001; Östlund-Nilsson *et al.*, 2005). Parasites are therefore recognized as important stress factors in all ecosystems.

In the last decade, stable isotopes have become popular in stress ecology studies (Wolf 51 et al., 2009), to assess stress-induced changes on the interactions between organisms and 52 53 between organisms and their environment. Changes in isotope signatures may, indeed, vary depending on various endogenous and environmental pressures via metabolic alterations 54 (Bowes et al., 2014; Yohannes et al., 2017; Karlson et al., 2018). Because stable isotopes 55 56 measures integrate dietary information over long timescales, they are useful tools to investigate the impact of parasite infections on individual hosts (Sánchez et al., 2013; 57 Knudsen et al., 2014), especially since some parasites are known to accumulate and survive in 58 their hosts over several weeks or months (Sandground, 1936). In addition, isotopic niche 59 being tightly correlated with trophic niche, stable isotopes are ecologically valuable for 60 quantifying changes in the trophic structure of communities and populations (Layman et al., 61 2007; Jackson et al., 2011). Consequently, stable isotopes could provide insight into parasite-62 related effects on the ecology of their host. Yet few studies have assessed the effects of 63 macroparasite infections on isotopic signature and niche structure of definitive hosts as well 64 as its functional significance (Britton and Andreou 2016; Gilbert et al., 2020; Riekenberg et 65 al., 2021). 66

67 Recent works suggest that the host's feeding behaviour, or diet, is likely to be a crucial
68 factor in shaping host–parasite interactions (Britton and Andreou 2016; Hite *et al.*, 2020),

especially for trophically transmitted parasites such as parasitic worms (i.e., helminths: 69 70 acanthocephalans, nematodes, cestodes and trematodes). The breadth of a host's trophic niche has long been associated with structuring helminth infections. For instance, greater diet 71 72 breadth increases potential encounter with parasite intermediate hosts, therefore leading to higher parasite abundance and taxon richness (Lagrue *et al.*, 2011; Paterson *et al.*, 2019). In a 73 recent metanalysis, Gutiérrez et al. (2019) found positive associations between trophic niche 74 75 breadth and trophically transmitted helminths in birds. On the other hand, parasites could affect the trophic niche of infected organisms trough impairment of host phenotypic traits or 76 behavioural alterations (Lefèvre et al., 2009; Weinersmith and Faulkes, 2014; Parker et al., 77 78 2015). The latter has been documented in manipulative parasites, which alter host habitat preferences, foraging activity, and antipredator behaviours (Loot et al., 2001; Poulin, 2007; 79 Friesen et al., 2018). Still, the relationship between trophic niche and trophically-transmitted 80 81 helminth infection is complex and not yet well understood, with contrasting results reported in the literature (Tierney, 2016; Pegg et al., 2017). The outcomes usually depend on several key 82 traits such as the life cycles, infection or feeding sites and modes of parasites. 83

Among parasites relying on trophic transmission to reach the definitive host, 84 acanthocephalans are ubiquitous obligatory intestinal worms encountered in a range of 85 taxonomically diverse host species, from aquatic to terrestrial habitats. These intestinal 86 parasites are commonly transmitted to vertebrate definitive hosts by aquatic crustacean preys 87 serving as intermediate hosts (Kennedy, 2006). In intermediate hosts, seemingly odd 88 89 behaviours of parasitized individuals have been reported, from reverse phototaxis (Franceschi et al., 2008) to changes in habitat and resource use (Fielding et al., 2003; Médoc et al., 2011). 90 These parasite-altered behaviours render hosts more susceptible to predation and increase the 91 92 likelihood of reaching the definitive host (Moore, 2002). Because parasite survival and transmission in intermediate host depends on the survival of the host itself, behavioural 93

responses to predation have been reported to switch from "predation suppression" during non-94 infective phase to "predation enhancement" when transmission is optimal, that is, when the 95 parasite reaches the stage infective to a definitive host (Dianne et al., 2011). In predatory final 96 97 hosts, parasites are likely to minimize pathogenicity to increase their own reproductive output and/or avoid multiple infections with other parasite species that may be harmful for the host. 98 One way to limit or avoid co-infections may occur by altering the trophic niche through 99 100 changes in the host feeding behaviour, which remains understudied for vertebrate definitive 101 hosts.

The objective of this study was to investigate interindividual variations in bivariate 102  $\delta^{13}$ C- $\delta^{15}$ N space between infected and uninfected fish cohorts and assess its potential 103 physiological consequence. To this purpose, we used a well-known and widespread host-104 parasite system, being composed of a freshwater fish the European chub, Squalius cephalus, 105 106 and its acanthocephalan parasite Pomphorhynchus sp., a thorny-headed worm living within the small intestines of vertebrates. Stable isotope data were used to calculate trophic niche 107 108 metrics and overlap using the Bayesian stable isotope approach. The Fulton's condition factor 109 provided relevant information on the energetic status and overall health of host individuals. We hypothesized that, under natural conditions, acanthocephalan parasites would prevent 110 111 multiple infection of their host with other parasite species through a reduction of the host trophic niche, hence closing the encounter filter for parasites acquired by ingestion, leading to 112 different host body condition according to their infection status. While controlling for 113 114 potential confounding factors such as age and body length of host individuals, we expected *i*) altered isotopic signatures between infected and uninfected fish, *ii*) decreased health status in 115 infected host, reflected by a poorer body condition, and *iii*) smaller isotopic niche in infected 116 117 cohorts as an evolutionary strategy in parasitism.

#### 119 2. Materials and Methods

#### 120 2.1 Ethic statement

Fish sampling was conducted in accordance with the relevant national and European
guidelines (L436-9, EN14011). The authorization for the scientific fish capture was delivered
by local administration authorities (Departmental Direction of Territories of Seine-et-Marne,
Val de Marne, Aisne, and Marne).

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#### 126 2.2 Sample collection

In September 2016, sampling was conducted on six stations of the Marne River and its 127 tributaries, France, along a 290-km-long stretch of the river (Table 1 and Table S1). Fish were 128 captured using a DEKA 3000 portable electrofishing unit ("Martin-pêcheur") in an upstream 129 direction, within a period of twelve days to limit temporal variations. Given the relatively 130 131 short-range movements of chubs (Penczak, 2006), fish from different sampling sites were considered distinct populations. A total of 118 acanthocephalan-infected and uninfected 132 European chub, Squalius cephalus, from 8.8 to 41.5 cm in length were captured (Table 1) and 133 fish were euthanized in 1 g/L MS222 solution. Total length (L<sub>T;</sub> nearest mm) and whole-fish 134 wet weight ( $W_{T_1} \pm 0.1$  g) were recorded. Prior tissue sampling, the Fulton's condition factor 135 (K, n = 118) was calculated according to the following equation:  $K = (W_T / L_F^3) \times 100$ . Note 136 that parasite mass was considered negligeable to fish mass when estimating its body condition 137 as Pomphorhynchus sp. represented less than 1% of fish mass even for heavily parasitized 138 139 ones (unpublished data). Scales were collected from each specimen above the lateral line and age determination was independently made by two different operators to limit interpretation 140 bias (see Molbert et al., 2020 for additional information). Chub intestine and body cavity were 141 examined for the presence of parasitic worms. For ectoparasites, external examination of fish 142 surface (skin and gills) was carried out. In our study, chub harbored only one parasite species, 143

which was initially identified *as Pomphorhynchus laevis* based on morphological
characteristics. However, without molecular identification to distinguish with other closely
related acanthocephalan species (e.g., *Pomphorhynchus tereticollis*), parasites were reported
as *Pomphorhynchus* sp. All parasites were found within the host intestine. A sample of dorsal
muscle was collected on each individual for isotopic measurements. These samples were then
lyophilized (Alpha1- 4LD plus; Martin Christ) for 48h, ground to obtain a fine powder and
weighed into aluminium capsules.

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Isotopic measurements. Isotopic measurements were performed on 0.20-0.25 mg of freeze-152 dried and powdered muscle samples (n = 113, some samples were lost because of combustion 153 issues) and analyzed by means of a Flash HT elemental analyzer (Thermofisher) coupled to a 154 Delta V mass spectrometer (Thermofisher) at the IRD laboratory in Bondy (France). Stable 155 156 isotope abundances are expressed as the deviation from international standards in parts per thousand ( $\delta$ ,  $\infty$ ) as follow:  $\delta X \approx [(Rsample / Rstandard) -1] * 1000, with X corresponding$ 157 to <sup>13</sup>C or <sup>15</sup>N and R representing <sup>13</sup>C/<sup>12</sup>C or <sup>15</sup>N/<sup>14</sup>N ratios. Atmospheric N2 and Pee Dee 158 Belemite were used as standards for nitrogen and carbon, respectively. The analytical 159 precision was found to be 0.1‰ and 0.7‰ for  $\delta^{13}$ C and  $\delta^{15}$ N, respectively. There was no lipid 160 161 correction applied to the data.

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#### 163 *2.3 Data analyses*

Infection levels of *Pomphorhynchus* sp. in *S. cephalus* were described as their prevalence (*i.e.*, number of infected fish/total number of fish x 100). Stable isotope data ( $\delta^{13}$ C and  $\delta^{15}$ N) from infected and uninfected fish cohorts were used to characterize isotopic signature and trophic structure from wild fish host populations in the Marne River. Two metrics of standard ellipse areas (SEAs) were calculated with the R package SIBER (Jackson *et al.*, 2011) to

assess the distribution of individuals in isotopic space, which enclose approximately 40% of 169 170 the data within groups (Jackson et al., 2011). SEA<sub>C</sub>, which stand for sample size corrected standard ellipse, gives a single estimate of ellipse area whereas SEA<sub>B</sub>, a Bayesian-estimated 171 standard ellipse area, gives a distribution of estimates that reflects variability in the data. Both 172 metrics, SEAc and SEA<sub>B</sub>, were calculated for infected and uninfected fish hosts and the 173 isotopic niche width was compared between groups (infected vs uninfected) by calculating the 174 175 probability that the distributions of SEA<sub>B</sub> differed from one another. The overlap in the isotopic niches of infected and uninfected host cohorts was calculated through the area of 176 overlap between two fitted ellipses based on the maximum likelihood and then as the 177 proportion overlap (%). Note that we did not obtain sufficient samples of uninfected hosts to 178 explore isotopic space across all locations. 179

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#### 181 *2.4 Statistical analyses*

All statistical models were performed in R v. 3.3.2 software (R Core Team, 2016). All data 182 were tested for normality and homogeneity before further statistical analyses and a 183 significance level of  $\alpha < 0.05$  was used for all tests. Since age and body length were highly 184 correlated ( $F_{(1,109)}$ = 484.5 , p < 0.001), we only used body length in our models to avoid 185 collinearity. Linear Mixed Models (LMMs) with site as random effects, to account for 186 different fish populations and environmental conditions, were used to assess i) the effect of 187 infection and body length on isotopic ratios ( $\delta^{15}N$  and  $\delta^{13}C$ ) and *ii*) the effect of parasite 188 infection and  $\delta^{15}N$  (as a proxy of trophic position) on condition factor. Analyses were 189 performed using the *lmer* statistical procedure available in the *lme4* and *lmerTest* packages for 190 linear mixed models (Pinheiro and Bates, 2000; Kuznetsova et al., 2017). Variance 191 components were estimated using a restricted maximum likelihood (REML) function and all 192 models had normal error structures. Data are presented as mean ( $\beta$ ) ± standard error (SE). 193

# 195 **3. Results**

Data and descriptive statistics are summarized in Table 1. The prevalence of intestinal worms 196 *Pomphorhynchus* sp, ranging from 50% to 90%, did not significantly differ among sampling 197 sites (Fisher-exact, p = 0.072). The probability of infection by acanthocephalan parasites was 198 199 not significantly correlated with body length (binomial GLMM: Z = 4.476, p=0.064) or age (Z = 4.645, p=0.067), and similarly, the body weight of infected and uninfected chub did not 200 significantly differ when the effect of body length was controlled (Gaussian GLMM: 201  $F_{(1,112)}=1.490$ , p>0.05). Importantly, the body condition was not significantly altered by 202 intestinal parasites *Pomphorhynchus* sp. (*K*; LMM:  $F_{(1.98.6)}$ =0.152, *p*>0.05), but was positively 203 correlated to  $\delta^{15}$ N (LMM:  $F_{(1.53.9)}$ =4.471, p=0.039,  $\beta \pm$  SE: 0.018  $\pm$  0.009). 204

The isotopic signature of chub varied among the different populations studied (LMM;  $\delta^{15}$ N: 205  $F_{(5,105)}=8.688$ , p<0.001 and  $\delta^{13}$ C:  $F_{(5,105)}=9.152$ , p<0.001) and ranged from 5.92 to 12.9 % for 206 and -30.2 to -22.8 % for  $\delta^{15}$ N and  $\delta^{13}$ C respectively (Table 1). Interestingly, there were no 207 208 significant effects of parasite infection by Pomphorhynchus sp. on stable isotopes ratios (LMM;  $\delta^{15}$ N:  $F_{(1,106.1)}=0.411$ , p=0.522 and  $\delta^{13}$ C:  $F_{(1,106.4)}=0.963$ , p=0.329). There was 209 however a consistent pattern of isotopic niche size (SEA<sub>B</sub> and its sample-corrected version, 210 SEA<sub>C</sub>) being considerably smaller in infected cohorts when compared with their non-infected 211 conspecifics (Table 2 and Figure 1), with niche width of uninfected groups being up to 3 212 times larger than that of infected ones. Overall, there were various degrees of isotopic overlap 213 214 between infected and uninfected cohorts among chub populations. Specifically, the isotopic niche of infected hosts has some overlap with uninfected conspecifics, ranging from 60% to 215 216 97% (i.e., infected niche within that of uninfected conspecifics; Table 2 and Figure 1).

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#### 218 **4. Discussion**

Parasitism is among the most common stress factors affecting free-living animals and how 219 parasites affect their host varies greatly, from causing minor alteration to death. One 220 understudied topic, albeit of greater ecological significance, is to evaluate potential shifts that 221 222 occur in trophic ecology of hosts following parasite infections, especially by trophically transmitted parasites. In this study, the use of a widespread chub-acanthocephalan system 223 revealed different patterns in the isotopic niche of infected and non-infected chub cohorts, as 224 measured by the standard ellipse area (*i.e.*, the isotopic niche) in the bivariate  $\delta^{13}$ C- $\delta^{15}$ N 225 226 space. Apart from two fish populations, infected subgroups showed a more restricted isotopic breadth compared to uninfected ones, although the extent of divergence between infected and 227 uninfected niches differed between populations. Additionally, the isotopic niches of infected 228 chub sat within that of uninfected ones for half of the populations under study. This illustrates 229 a potential trophic specialization, with infected fish consuming specific food items that are 230 231 also within the dietary range of uninfected chub, and this non-lethal consequence of parasite infection may have strong implications for the feeding ecology of the host. 232

There are empirical examples of an association between parasite infection and 233 234 shifts in host trophic niche, but they remain somewhat restricted for macroparasites (Bernardo and Singer, 2017). Parasite-mediated shifts in trophic niche can be associated to the cost 235 exerted by parasites or modifications of the host phenotype, which can influence diet 236 preferences or prey selectivity (Britton and Andreou, 2016; Vivas Muñoz et al., 2021). These 237 changes, which can limit hosts' niche usage, are either adaptative for the host or the parasite 238 as they redress physiological deficits imposed by parasite infections (Britton, Pegg and 239 Williams, 2011; Sánchez et al., 2013; Pegg et al., 2015), provide nutritional requirements for 240 parasites growth, reproduction and survival (Esch and Fernández, 1993) or limit co-infection 241 which might be detrimental for the established parasite (direct competition for space and 242 resources, or indirect through the release of toxic products; Ferrari et al., 2009). For instance, 243

the tapeworm Schistocephalus solidus causes infected sticklebacks, Gasterosteus aculeatus, to 244 245 eat smaller prey (Milinski, 1984), or selectively preyed on larger items than uninfected individuals, which compensated for the energy cost of infection (Ranta, 1995). Similarly, 246 247 Pegg et al. (2017) reported a consistent niche specialization in two cyprinid fish species infected by a copepod gill parasite. They speculated that infected fishes increased their 248 predation on prey requiring relatively low-energy expenditure to capture (*i.e.*, abundant and/ 249 250 or relatively slow moving) as a consequence of parasite-associated energetic costs. In wild 251 common carp Cyprinus carpio, individuals infected with cestodes shifted to lower trophic status prey items as a result of pathological damage to the gastrointestinal tract (Britton, Pegg 252 253 and Williams, 2011). Despite the variation in isotopic niche between chub cohorts, the absence of differences in relative body condition (as measured by the Fulton's condition 254 factor) between infected and uninfected fish may suggest that hosts maintained their energetic 255 256 requirements. While adult acanthocephalans are usually less virulent in the final host, thornyheaded parasites are known to perforate the gut wall with their proboscis and, under heavily 257 258 infection, obstruct the digestive tube and cause severe local inflammation (Taraschewski, 259 2000). Cases of serious illness, enhanced mortality or altered reproduction in acanthocephalan-infected fish are seldom, although fitness cost for the vertebrate host has 260 already been observed in fish in the face of infection by acanthocephalan parasites (Salal et 261 al., 2001; Rajeshkumar et al., 2013; Sakthivel, 2014; Atalabi et al., 2018). Here, the low 262 abundance of *Pomphorhynchus* sp. might explain the absence of pathological effects on the 263 264 host.

Interestingly, trophic niche use at the study sites showed different trends for infected and uninfected cohorts among chub populations. This result suggests that parasite infection on niche specialization is context-dependent (Thorburn *et al.*, 2022). There is mounting evidence that stress factors, other than parasite infection, can alter physiological

performance and affect interindividual variability of the isotopic signature (Karlson et al., 269 270 2018). Chub populations were, among other things, exposed to various contaminants from industrial, agricultural, and urban activities along the Marne River (Molbert et al., 2020). A 271 272 previous study, based on the same host-parasite system, evidenced enhance oxidative stress in non-infected chubs exposed to high contaminant levels, which could drive fitness costs for the 273 274 host (Molbert et al., 2021). Although the body condition did not differ with the infection status, alteration in metabolic pathways through differential exposure to contaminants could 275 276 explain interpopulation variations in trophic niches between infected and non-infected cohorts. In fact, the lack of evidence at higher biological scale could be explained by the 277 "selection hypothesis", stating that less fit animals are sampled less frequently as a result of 278 more efficient predation on sick and weak animals. 279

280 Since our study only reveals association, we cannot ascertain causation between 281 niche specialization and parasite infection. Alternatively, dietary specialization could be a driver of Pomphorhynchus sp. infection (Britton and Andreou, 2016, Gutiérrez et al., 2017), 282 as specialized feeding behavior can lead to large variations in exposure to parasites among 283 individual hosts. Infection in chub populations at the study sites, was solely explained by a 284 single genus, which suggests intense feeding on gammarid intermediate hosts or infected 285 286 paratenic fish hosts. Smaller SEAc in infected chub cohorts compared to uninfected ones also support that infected fish had more selective diet, while uninfected conspecifics appeared to 287 be more generalist feeders. In line with those findings, Costa- Pereira et al. (2019) 288 demonstrated that thin-toed grogs, Leptodactylus spp, with lower helminth parasite load 289 adopted a more flexible feeding strstrategy and suggested that diet consistency may lead to 290 increase infection rates. Although parasite infection may lead to smaller niches formed by 291 292 groups of trophically specialised individuals (Bolnick et al., 2003; Quevedo et al., 2009; Pegg et al., 2017), Tierney (2016) showed an opposite trend, with infected cohorts of brown trout, 293

Salmo trutta, having greater SEAc than non-infected ones. Where brown trout were infected by seven species of helminth parasites, including the acanthocephalan *Pomphorhynchus tereticollis*, we only found one acanthocephalan genus among 6 populations along a 290-kmlong stretch of the Marne River. These results support the hypothesis that higher parasite species richness is associated with greater niche breadth, and thus higher probability to encounter parasite intermediate hosts.

300 Last but not least, isotopic turnover rates are important for understanding how parasite infection affects stable isotope fractionation. While carbon and nitrogen half-lives in 301 fish muscle are ca. 50 and 100 days, respectively (Church et al., 2009; Buchheister and 302 303 Latour, 2010; Busst and Brittons, 2018), the lifespan of most fish acanthocephalans in the intestine of fish host does usually not exceed 7-8 months (Nachev and Sures, 2016). In this 304 305 study, however, the dynamics of current and past infection by *Pomphorhynchus* sp., or other 306 parasitic worms, is unknown. As a result, parasite infection could be confounded given the slow turnover rate of stable isotope in muscle. With naturally infected fish, parasite effect on 307 308 the isotopic niche or composition should therefore be treated with caution. Overall, future challenges lie in the determination of the influence of trophic niche on helminth infection (or 309 vice versa) and its functional significance, especially for host-parasite systems composed of 310 311 macroparasites with complex life cycle and vertebrate hosts. To this purpose, emphasis should be placed on studies combining field studies and experimental approaches, and parasite effect 312 on host trophic niche should be considered within a comprehensive framework rather than in 313 314 isolation.

315

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319

# 320 **Conflicts of interest**

321 The authors declare no conflict of interest.

322

## 323 Contributions

A.G. acquired the funding, conceived the idea, designed the methodology and captured the fish. N.M. carried out stable isotope analyses, ran statistical analyses and drafted the manuscript. Both authors contributed critically to the manuscript draft and gave final accordance for publication.

328

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