1	Parallel trophic diversifications in polyploid cyprinid fish from East
2	Africa: from preadaptive polymorphism to trophic specialization
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Abstract

28 Trophic diversification is one of the main mechanisms driving the adaptive radiation. The 29 polyploid lineage of the cyprinid genus Labeobarbus represent an excellent model for 30 studying the trophically-based adaptive radiation in either lacustrine or riverine 31 environments. Recently discovered four diversifications in rivers of the Ethiopian Highlands 32 (East Africa) demonstrate independently evolved repeated mouth polymorphisms each 33 represented by four core mouth phenotypes: (i) generalized, (ii) thick-lipped, (iii) scraping, 34 and iv) large-mouthed. Mouth phenotypes in some radiations can be further divided to 35 subtypes representing from four to eight sympatric ecomorphs. Using the stable isotope and 36 gut content analyses we tested hypothesis on trophic resource partitioning within each 37 radiation, revealed disparity in degree of diversification between radiations and tried to 38 reconstruct the process of trophic diversification. Three of four radiations demonstrated 39 partitioning of trophic resources within five trophic niches: i) detritophagy, ii) 40 macrophytophagy, iii) invertivorous benthophagy, iv) periphyton feeding, and v) piscivory. 41 The studied riverine radiations were likely at the different stages of the diversification. One 42 radiation having a similar set of mouth phenotypes was not trophically divergent displaying a 43 remarkable decouple of form and function. A unique case of ecologically non-functional 44 mouth polymorphism at an incipient stage of trophic diversification supports a concept of the 45 plasticity-first evolution. This phenomenon stems from the pre-existing genomic templates of 46 mouth polymorphism ancestrally inherited upon the allopolyploid origin of the *Labeobarbus* 47 lineage. The predetermined and preadaptive mouth polymorphism can be considered a key 48 innovation of the *Labeobarbus* that promoted to resource-based diversification via adaptive 49 radiation.

51 Running Title: Preadaptive polymorphism and trophic specialization

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53 Key words: adaptive radiation, polymorphism, trophic resource partitioning, ecological
54 speciation, vertebrates

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56 1. Introduction

57 Trophic divergence is one of the main drivers of ecological speciation and adaptive radiation 58 (Schluter, 2000; Seehausen & Wagner, 2014; Martin & Richards, 2019). The fishes being the 59 most diverse group among vertebrates display numerous examples of trophic diversification 60 during adaptive radiation - not only among textbook examples from cichlids, whitefish, 61 Arctic charrs (Seehausen & Wagner, 2014) - but also in other taxa. Despite tremendous 62 amount of studies focused on the mechanisms underpinning the rapid diversification during 63 adaptive radiation (and achieved progress – e.g., Brawand et al., 2014; McGee et al., 2020; 64 Ronco *et al.*, 2021), the phenomenon is still poorly understood (Martin & Richards, 2019; 65 Gillespie et al., 2020). One of the greatest findings of the last decade – a contribution of the 66 heterogenous genome (usually evolved via past hybridization) into accelerated rates of 67 ecological diversification and speciation (Nosil et al., 2009; Genner & Turner, 2012; Meier et 68 al., 2017; Irisarri et al., 2018; Marques et al., 2019; Svardal et al., 2020). One of the 69 mechanisms to increase genetic heterogeneity is a polyploidization. After a round of fish-70 specific genome duplication (Meyer & Van de Peer, 2005) occurred ca. 350 mya in the ray-71 finned fish (Actinopterygii) lineage, the numerous further polyploidization events happened 72 within and resulted in >1000 species mostly within Cypriniformes, Salmoniformes and 73 Acipenseriformes (Yang et al., 2022). Contrary to plants, the animal polyploidization is

74 realized predominantly via allopolyploidization (Gregory & Mable, 2005) that means the 75 most of fish polyploid lineages contains the different subgenomes. Therefore, the polyploid 76 lineages look as perfect candidates to test a hypothesis on role of complex genome in further 77 diversification during adaptive radiation. Herein a polyploidy apart prospective gains also has 78 many restrictions on fish biology that can dismiss the expected benefits (Gregory & Mable, 79 2005; Van de Peer et al., 2017). Nevertheless, the most researchers agreed that polyploid taxa 80 are characterized by increased ecological and phenotypic plasticity that (i) facilitate to 81 overcome the stressful and harsh environment and (ii) can promote to further diversification 82 and speciation that is within 'flexible stem' hypothesis (West-Eberhard, 2003; Wund et al., 83 2008; Li & Guo, 2020; Van de Peer et al., 2021). Some polyploid lineages, for instance – 84 Salmonidae and Coregonidae – represent nice examples of phenotypic and ecological 85 plasticity along with possibility to evolve small-scale adaptive radiation especially within 86 genera Salvelinus, Salmo, and Coregonus (Reshetnikov, 1980; Bernatchez et al., 1999; 87 Alekseyev et al., 2002; Markevich et al., 2018; Segherloo et al., 2021; Levin et al., 2022).

88 The cyprinid fishes (fam. Cyprinidae sensu Tan & Armbruster, 2018) being one of the 89 most diverse families of teleostean fishes (> 1780 species - Fricke et al., 2023) are the 90 champions in the number of the polyplod species (ca. 600 species – Yang et al., 2022). Some 91 of cyprinid polyploid lineages display bright examples of adaptive radiation based on trophic 92 polymorphism (Nagelkerke et al., 1994; Roberts & Khaironizam, 2008; Levin et al., 2019; 93 2021; Qiao et al., 2020; Komarova et al., 2021). Among these, the most distinguished one in 94 terms of the morpho-ecological diversification is the cyprinid genus Labeobarbus 95 representing species-rich African hexaploid lineage (2n=150 – Oellerman & Skelton, 1990; 96 Golubtsov & Krysanov, 1993) that comprises >130 species (Vreven et al., 2016; Fricke et al., 97 2023) and displays numerous trophically-based adaptive radiations (Nagelkerke *et al.*, 1994; 98 2015; Mina et al., 1996; Sibbing & Nagelkerke, 2000; Dimmick et al., 2001; Golubtsov,

2010; Shkil *et al.*, 2015; Levin *et al.*, 2020). The *Labeobarbus* is widespread in Africa and
presents in each of the ten African ichthyofaunal provinces (Vreven *et al.*, 2016). Based on
the recent data on phenotypic, ecological, and genetic data, the taxonomic diversity of the *Labeobarbus* might be greatly underestimated (Levin *et al.*, 2019, 2020; Decru *et al.*, 2022).

103 The Ethiopian Highlands is remarkably distinguished as a hotspot for multiple adaptive 104 radiations of the *Labeobarbus* discovered during last 30 years (see Golubtsov et al., 2021). 105 The most famous example of rapid adaptive radiation among non-cichlid fishes belongs to 106 the lacustrine radiation of the Labeobarbus in Lake Tana, Ethiopia, where up to 15 107 species/ecomorphs were described that partitioned trophic resources (Nagelkerke *et al.*, 1994; 108 Mina et al., 1996; Zworykin et al., 2006; Shkil et al., 2015). Apart from the lacustrine 109 radiation strongly predominating among fishes (e.g. Seehausen & Wagner, 2014), there are 110 striking examples of the riverine adaptive radiations in non-cyprinid (Turner *et al.*, 1985; 111 Piálek et al., 2012; Esin et al., 2021; Říčan et al., 2021; Burress et al., 2022) as well as 112 cyprinid fishes - Labeobarbus, in particular (Levin et al., 2019; 2020).

113 The most conspicuous phenotypic difference between sympatric ecomorphs of the 114 Labeobarbus refers to mouth type variation. The core of mouth polymorphism is composed 115 of four main phenotypes (Figure 1) that were recorded throughout Africa (Worthington, 116 1929; 1933; Matthes, 1963; Banister, 1973; Skelton et al., 1991; Nagelkerke et al., 1994): i) 117 generalized (typical for barbs), ii) thick-lipped ('rubberlip'), iii) scraping ('chiselmouth'), and 118 iv) large-mouthed. Mouth phenotypes are subjected to further sub-diversification and can 119 include subtypes or variants that are recruited for occupying the new ecological niches (Levin 120 et al., 2021b). Moreover, there are intermediate phenotypes possibly of hybrid origin 121 (Banister, 1972; Vreven et al., 2019) that does not exclude their further specialization.

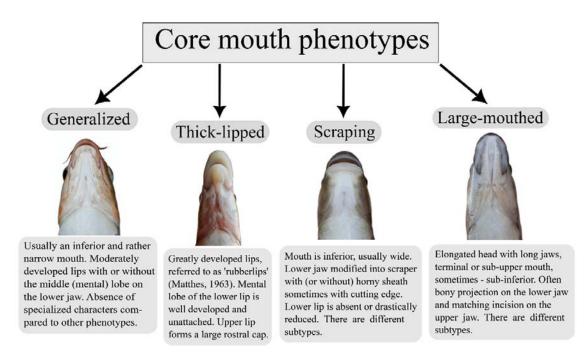


Figure 1. Pictorial scheme of core mouth phenotypes of the *Labeobarbus*. Photographexamples referred to the *Labeobarbus* from the rivers of the Ethiopian Highlands.

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126	This outstanding mouth polymorphism of the Labeobarbus likely stems from ancestral
127	inherited genetic templates since both parental lineages involved in the allopolyploidization
128	event (Yang et al., 2015; 2022) belong to extant lineages and represent different mouth types.
129	The contemporary representatives of the maternal lineage are characterized by generalized
130	and thick-lipped mouth polymorphism (tetraploid Torinae, 2n = 100 - Roberts &
131	Khaironizam, 2008; Walton et al., 2017; Coad, 2021) while paternal diploid lineage
132	(Cyprinion, 2 n = 50, Ünlü et al., 1997; Esmaeili & Piravar, 2006) is represented by scraping
133	mouth phenotype (Coad, 2021). Hence, mouth polymorphism of the Labeobarbus is
134	ancestrally heritable, likely preadaptive and might re-evolve.

135 The parallel diversifications of the *Labeobarbus* in the riverine environment of the 136 Ethiopian Highlands were recently discovered within the four geographically isolated 137 drainages - i) the Didessa River in Blue Nile basin; ii) the Sore River in White Nile basin; iii) 138 the Gojeb River in Omo-Turkana basin; and iv) the Genale River in Juba-Wabe-Shebelle 139 basin, Indian Ocean catchment (Mina et al., 1998; Golubtsov, 2010; Levin et al., 2019; 2020; 140 Golubtsov *et al.*, 2021). Each diversification includes from four to eight sympatric ecomorphs 141 divergent in mouth phenotype (Table 1) that presumably involves trophic adaptations. The 142 genetic studies showed that all four riverine radiations of *Labeobarbus* were independently 143 evolved from different ancestors although they are very closely-related based on mtDNA data 144 that suggests recent and rapid diversification (Levin et al., 2013; 2019; 2020). Moreover, 145 existing data suggests these radiations might be at various stages of the diversification (Levin 146 et al., 2020; 2023 - in press). Of special interest is that these radiations could repeatedly 147 emerge in the riverine environment that has been considered to be inappropriate for adaptive 148 radiation due to heterogeneous and unpredictable conditions for a long time (reviewed in 149 Burress et al., 2023).

150 A few studies on trophic ecology showed that sympatric riverine ecomorphs of the 151 Labeobarbus bearing different mouth phenotypes have partitioned trophic resources - in 152 particular, in the Genale River (Levin et al., 2019). Other riverine radiations of the 153 Labeobarbus were not studied comprehensively from the point of trophic divergence. The 154 recent but independent origin of similar diversifications in mouth phenotypes raises several 155 questions. Do all diversifications partition trophic resources according to homologous mouth 156 phenotypes? What is trophic specialization (if present) of various sympatric ecomorphs? 157 Whether riverine diversifications comprising different numbers of ecomorphs / mouth 158 (sub)phenotypes are at various stages of the process? If so, could patterns of the trophic 159 diversification be sorted from more primitive to advanced aiming reconstruction of the

trophic radiation? We try to answer these questions in a study of trophic diversification of
sympatric ecomorphs in each of four *Labeobarbus* riverine radiations using the stable isotope
composition and gut content.

Table 1. The composition of riverine sympatric ecomorphs of *Labeobarbus* spp. in
isolated riverine basins in the Ethiopian Highlands (summarized from Golubtsov, 2010;
Levin *et al.*, 2019; 2020; 2021; Mina *et al.*, 1998 and our unpublished data). The photographs
of the fish ecomorphs are given in Figure 2.

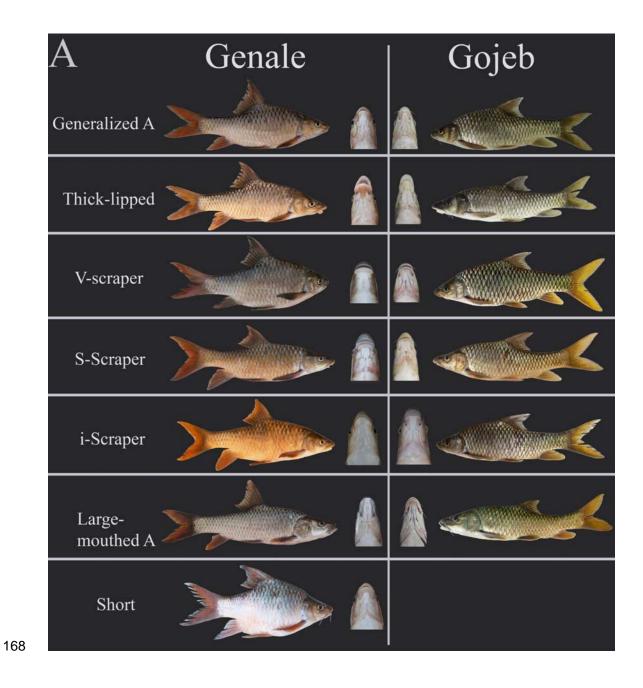
Species, river and	Ecomorphs	Source
basin		
Labeobarbus	1. Generalized A (common)	Golubtsov, 1993; Mina et al.,
gananensis complex.	2. Thick-lipped	1998; Dimmick et al., 2001;
Genale River, Juba-	3. V-scraper ¹ (<i>L. jubae</i>)	Levin et al., 2019; Golubtsov
Wabe-Shebelle basin,	4. S-scraper ²	et al., 2021
Indian Ocean	5. i-Scraper ³	
catchment.	6. Short ⁴	
	7. Large-mouthed A ⁵	
Labeobarbus	1. Generalized A	Golubtsov, 2010; Mina et al.,
intermedius complex,	2. Generalized B ⁶	1998; Levin et al., 2020; 2021;
L. zaphiri, L. beso.	3. Thick-lipped	original observations
Didessa River, Blue	4. Semi-thicklipped ⁷	
Nile basin, Atlantic	5. V-scraper (L. beso)	
Ocean catchment.	6. Large-mouthed A1 (<i>L</i> .	

zaphiri)

7.	Large-mouthed	A2
(Labe	obarbus sp.1) ⁸	
8.	Large-mouthed	B ⁹
(Labe	obarbus sp.2)	

Labeobarbus	1. Generalized A	Golubtsov	, 2010; Levi	n <i>et al</i> .,
intermedius complex.	2. Thick-lipped	2020;	2021;	original
Gojeb River, Omo-	3. V-scraper	observatio	ons	
Turkana basin, inland	4. S-scraper			
drainage.	5. i-Scraper			
	6. Large-mouthed A			
		0.1.1.	2010 1 :	
Labeobarbus	- Generalized A	Golubtsov	2010; Levi	n <i>et al</i> .,
intermedius complex.	- Thick-lipped	2020; orig	inal observat	tions
Sore River, White Nile	- V-scraper			
	T (1 19			
basin, Atlantic Ocean	- Large-mouthed			

167 ^{1,2,3,4,5,6,7,8,9} see detailed description of the ecomorphs in the Supplementary materials (File S1).



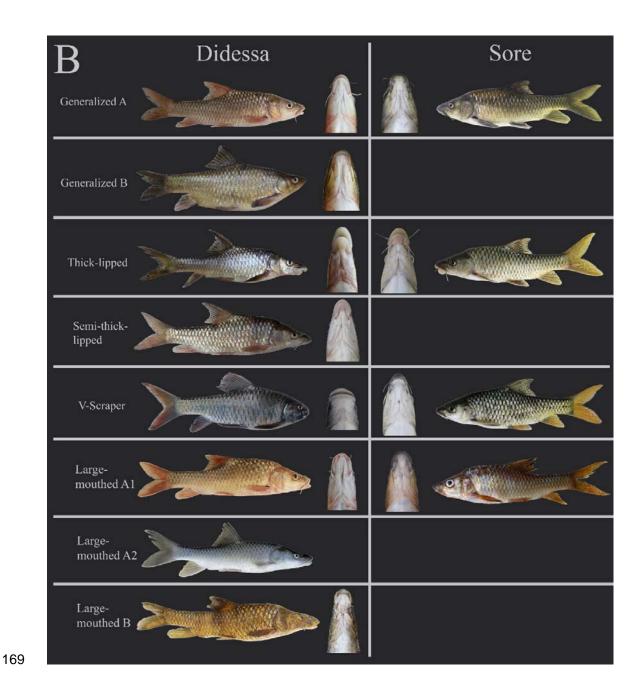
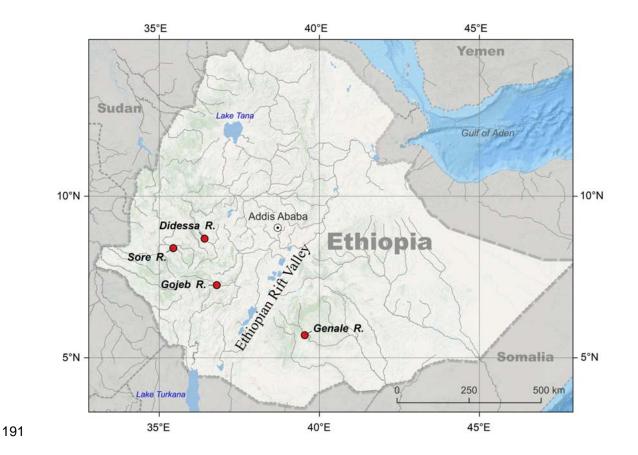


Figure 2. Appearance and mouth phenotypes of the ecomorphs of the *Labeobarbus* in different rivers of the Ethiopian Highlands: (A) the Genale and Gojeb Rivers; (B) the Didessa and Sore Rivers. An exceptionally rare representative of the Large-mouthed A2 ecomorph from the Didessa River is presented only from the lateral side.

175 2. Material and Methods

Ethical approval. All applicable international, national, and institutional guidelines for the
care and use of animals were strictly followed. All animal sample collection protocols
complied with the current laws of Russian Federation and Federal Democratic Republic
of Ethiopia.

180 Study sites and material studied. Samples were collected under the framework of the Joint 181 Ethiopian Russian Biological Expedition (JERBE) in four rivers draining the Ethiopian 182 Highlands and belonging to four major river basins (Fig. 2): (1) the Juba-Wabe-Shebelle 183 basin in the Indian Ocean catchment – the Genale $R. - 5.7025^{\circ} N 39.5446^{\circ} E$; (2) the Blue 184 Nile basin – the Didessa R., a tributary of the Blue Nile – 8.6921° N 36.4144° E; (3) the 185 Omo-Turkana enclosed basin – the Gojeb R., a tributary of the Omo R. – 7.2539° N 36.7943° 186 E; (4) the White Nile basin – the Sore R. \sim 35 km downstream City Metu – 8.3987° N, 187 35.4378° E. Fish were caught by gill and cast nets in February–March 2011 (Didessa), 188 March-April 2009, 2019 (Genale), February 2011 (Gojeb), April 2014 (Sore). Fish were 189 killed with an overdose of MS-222 anaesthetic and then photographed using a Canon EOS 190 50D camera (Canon Inc. Tokyo, Japan).



192 Figure 3. Map of sampling sites of the four riverine radiations of the Ethiopian *Labeobarbus*.

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Standard length (SL, mm) and gut length (GL, mm) were measured with a ruler. Fish were preserved first in 10% formalin and then transferred to 70 % ethanol. Gut length was studied in 461 individuals. All specimens were deposited at A.N. Severtsov Institute of Ecology and Evolution, the Russian Academy of Sciences, Moscow under provisional labels of JERBE. The detailed information on material studied (gut length, stable isotope composition and gut content) is given in Supporting Table S2.

Stable isotopes composition. For stable isotope (SI) analyses, white muscle tissue from the
dorsal side of the body under the dorsal fin was sampled from freshly collected specimens.
White muscle samples were dried at 60 °C. The samples were weighed using a Mettler

203 Toledo MX5 microbalance (Mettler Toledo, Columbus, OH, United States) with 2 µg 204 accuracy, and wrapped in tin capsules. The weight of the fish tissue samples varied from 250 205 to 500 µg. SI analysis was conducted at the Joint Usage Center of the A.N. Severtsov Institute 206 of Ecology and Evolution RAS, Moscow. Briefly, a Thermo Delta V Plus continuous-flow 207 IRMS was coupled to an elemental analyzer (Flash 1112) equipped with a Thermo No-Blank 208 device. The isotopic composition of N and C was expressed in the δ notation relative to the 209 international standards (atmospheric nitrogen and VPDB, respectively): δX (‰) = 210 $[(Rsample/Rstandard) - 1] \times 1000$, where R is the molar ratio of the heavier and lighter 211 isotopes. The samples were analyzed with a reference gas calibrated against the International 212 Atomic Energy Agency (IAEA) reference materials USGS 40 and USGS 41 (glutamic acid). 213 The measurement accuracy was ± 0.2 %. Along with the isotopic analysis, the nitrogen and 214 carbon content (as %) and C/N ratios were determined. In total, 400 white-muscle samples 215 were analyzed.

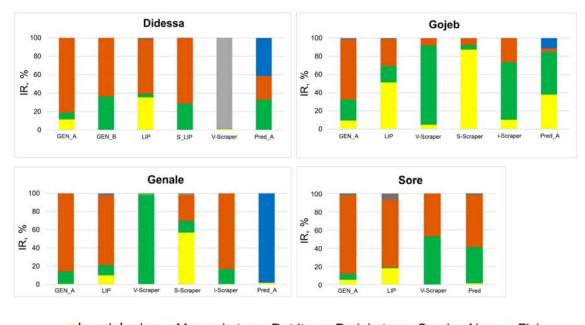
216 *Diet analysis.* Following the best practice for estimation of trophic niches the in tropic rivers 217 (e.g., Davis et al., 2012) we studied gut content in addition to SI composition. Gut content 218 was extracted from preserved specimens, dried on filter paper and weighed using a Pioneer 219 PX84/E balance with 0.0001 g accuracy. The diet particles were identified using Olympus 220 microscope $(100-1000 \times$ magnification) Motic DMW-143-N2GG CX41 and 221 stereomicroscope $(100-400 \times \text{magnification})$. The diet components were grouped into: (i) 222 detritus, (ii) invertebrates, (iii) macrophytes, (iv) periphyton, (v) filamentous algae, (vi) fish 223 (body remnants and scales), and (vii) mineral ground. The group 'Invertebrates' included 224 mainly the larvae and imago of amphibiotic insects and their fragments as well as imago of 225 aerial insects (Coleoptera, Hymenoptera); rarely Cladocera. The group 'Macrophytes' 226 included any fragments of helophytic and semi-aquatic plants - such as leaves, stems or 227 seeds. A composite measure of diet, an index of relative importance (IR) [Natarajan &

Jhingran, 1961; Popova & Reshetnikov, 2011], was used to assess the contribution of different components to the diet. The IR index was calculated as follows: IR = $(Fi \times Pi)/(\sum(Fi \times Pi)) \times 100\%$, where Fi = the frequency of occurrence of each food group, and *Pi* = its part by weight; the value of *i* itself changes from 1 to n (n = the part of food organisms in the food bolus). In total, 195 food boluses were analyzed for gut content (Supplementary Table S2).

234 Statistical analyses. Several R packages and functions were used for the statistical analyses 235 and plot construction. Basal descriptive statistics was obtained using the summarytools 236 library [Comtois, 2022]. Central tendencies in the text are presented as means and 1SD. The 237 Kruskal-Wallis test was applied for comparisons of the values of gut length GL and SI 238 composition using the function kruskalTest in each radiation (in library PMCMRplus) with 239 subsequent pairwise comparisons using post-hoc Dunn's all-pairs test with Bonferoni 240 adjustment - function kwAllPairsDunnTest [R Core Team, 2023]. The violin boxplots were 241 obtained using the ggplot2 library [Wickham, 2022]. The package SIBER v.2.1.6 [Jackson et 242 al., 2011] was used to assess the differences in the isotopic trophic niche features. The total 243 convex hull areas (TA), core trophic niche breadths, and sample size-corrected standard 244 ellipse area (SEAc) were estimated. The trophic overlap for 95% TA was estimated using 245 nicheROVER [Lysy et al., 2021], a method that is insensitive to the sample size and 246 incorporates statistical uncertainty using Bayesian approach [Swanson et al., 2015].

247 **3. Results**

3.1. Diet. Food spectra of *Labeobarbus* spp. were rather diverse. Feeding of sympatric
ecomorphs in all four rivers was divergent but at various degrees as estimated by index of
relative importance (Figure 4). Detailed description of the diet for each ecomorph is given in
Supplementary material S3.



Invertebrates Macrophytes Detritus Periphyton Sand Algae Fish

Figure 4. Food spectra (IR: the index of relative importance) of the sympatric ecomorphs of the *Labeobarbus* spp. from the Didessa, Gojeb, Genale, and Sore rivers. Abbreviations of the ecomorphs: GEN_A - Generalized A; GEN_B - Generalized B; LIP - Thick-lipped; S_LIP -Semi-thicklipped; Pred_A - Large-mouthed A; Pred_B - Large-mouthed B.

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258 Based on the food spectra, up to five trophic specializations can be detected in the 259 Didessa River: i) detritivore (Generalized A), ii) detritivore-macrophytophage (Generalized B 260 and Semi-thicklipped), iii) detritivore-invertivore (Lipped), iv) periphyton feeder (V-scraper), 261 and v) piscivore-omnivore (Large-mouthed A and B). Notably, detritus seems to be a core 262 food for many ecomorphs except for the V-Scraper and Large-mouthed. A sub-specialization 263 within the piscivory strategy is possible taking into account the drastic difference in mouth 264 structure (Figure 2B) and diet between the Large-mouthed ecomorphs. Seemingly, up to five 265 feeding strategies can be distinguished in the Gojeb River: i) detritivore (Generalized A), ii) 266 invertivore-detritivore (Lipped); iii) macrophytophagous (V-Scraper and i-Scraper), iv)

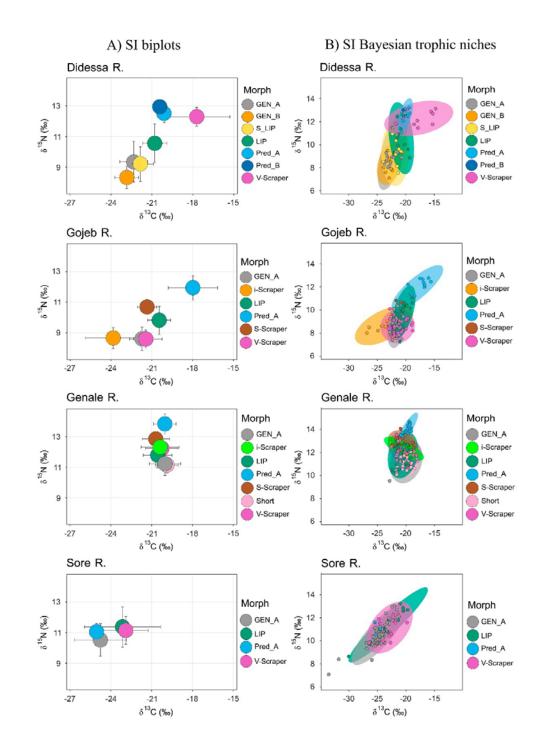
267 invertivore (S-scraper), and v) omnivore-piscivore (Large-mouthed A). Compared to the 268 Didessa, a detritus was a core food only for the Generalized A ecomorph. Large-mouthed 269 ecomorph is not predominantly piscivorous but rather omnivorous with a small portion of 270 fish food. In the Genale River, up to four feeding strategies can be detected -i) detritivore 271 (Generalized A, Lipped, and i-Scraper), ii) macrophytophagous (V-Scraper), iii) invertivore 272 (S-scraper), and iv) piscivore (Large-mouthed A). Remarkably, a detritus was a core food for 273 three ecomorphs in the Gojeb River, while the Large-mouthed ecomorph is a strongly 274 piscivorous specialist. In contrast, little diversification in food spectra was detected in the 275 Sore River. Briefly, two feeding strategies can be recognized: i) detritivore (Generalized A, 276 Lipped, and i-Scraper), and ii) macrophytophagous-detritivore (V-Scraper and Large-277 mouthed). Remarkably detritus was a core food for three of four ecomorphs although a 278 notable portion (almost 20 %) of benthic invertebrates was detected in the Lipped ecomorph. 279 Large-mouthed ecomorph is not a piscivorous specialist in the Sore River.

280 *3.2. Stable Isotope Composition.* Basic statistics for both $\delta^{15}N$ and $\delta^{13}C$ values and data on 281 the total area (TA), standard ellipse area (SEA), and corrected standard ellipse area (SEAc) 282 are given in Supplementary Materials S4. Detailed results are presented in subsequent 283 sections.

284 *3.2.1. Didessa Radiation.* Among seven ecomorphs, several were significantly divergent in SI 285 values (six pairwise comparisons in both δ^{15} N and δ^{13} C - Table S4). The largest values of 286 δ^{15} N were detected in two Large-mouthed (piscivory) ecomorphs (12.5±0.6 and 12.9‰) as 287 well as in highly specialized V-scraper *L. beso* (12.3±0.6‰). The V-scraper was also most 288 enriched in ¹³C (δ^{13} C -17.7±2.4%) (Figure 5). The lowest values of both δ^{15} N (8.33±0.7%) 289 and δ^{13} C (-22.8±0.9%) were in the Generalized B ecomorph. Other ecomorphs, Generalized 290 A, Thick-lipped, and Semi-thicklipped, had intermediate δ^{15} N and δ^{13} C values, although

Thick-lipped ecomorph had somewhat higher $\delta^{15}N$ values (10.6±1.3‰) compared to both 291 292 Generalized A and Semi-thicklipped $(8.3\pm1.3 \text{ and } 9.2\pm1.1\%)$, respectively). The maximum 293 difference in mean δ^{15} N values between ecomorphs in the Didessa was 4.61^{\overlines}. The highest difference between ecomorphs in the mean δ^{13} C values was 5.10%. The isotopic niches (as 294 295 assessed by the standard ellipses) were almost fully separated between Generalized B and V-296 Scraper (overlap <1%), between Generalized B and Large-mouthed A (overlap <1%), and 297 weakly overlapped between Semi-Thicklipped and V-scraper, Semi-Thicklipped and Large-298 mouthed A, and Generalized A and V-Scraper (overlap <6%, <10%, and <11%, 299 respectively) (see details in Supplementary File S4). The most overlapping trophic niches 300 were between Generalized A and B (57 and 90%) and between Generalized A and Semi-301 Thicklipped (77 and 79%) (Supplementary File S4).

302 3.2.2. Gojeb Radiation. Among six ecomorphs, some were significantly divergent from others in SI values (five and six pairwise comparisons in δ^{15} N and δ^{13} C values, respectively; 303 Table S4). The largest values of $\delta^{15}N$ and $\delta^{13}C$ were found in Large-mouthed (piscivory) 304 ecomorph (11.94±0.8‰ and -17.98 ±1.8‰, respectively) that was notably (sometimes 305 306 significantly - Figure 5, Table S4) higher than in all other sympatric ecomorphs. The minimal values of $\delta^{15}N$ were detected in the V-scraper (8.60±0.6‰) and Generalized (8.61±0.8‰) 307 ecomorphs while in the i-Scraper ecomorph the δ^{13} C value was at the minimum (-23.84±2.0 308 ‰). The total divergence between ecomorphs was lower in δ^{15} N but higher in δ^{13} C values 309 310 compared to the Didessa radiation. In particular, the highest difference between ecomorphs in the mean δ^{15} N values achieved 3.34‰, while the same for the mean δ^{13} C values was 5.86‰ 311 312 (Figure 5, Table S4). The isotopic niches were almost fully separated between Large-313 mouthed and V-Scraper (overlap <2%) and between Generalized and Large-mouthed (<4%) 314 while the most overlapping trophic niches were between Generalized and V-Scraper (overlap 315 70 and 91%) (Supplementary File S4).



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Figure. 5. Stable isotope biplots (A) showing mean values and 1 SD, and Bayesian ellipses showing trophic niche widths and overlaps (B) in sympatric ecomorphs of the *Labeobarbus* spp. from the Didessa, Gojeb, Genale, and Sore Rivers. Ellipses with 95% confidence intervals are based on standard ellipses corrected for small sample sizes (SEAc; isotopic niche metrics; SIBER package). Each point corresponds to the individual isotopic value.

Abbreviations of the ecomorphs: GEN_A - Generalized A; GEN_B - Generalized B; LIP -Thick-lipped; S_LIP - Semi-thicklipped; Pred_A - Large-mouthed A; Pred_B - Largemouthed B.

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326 3.2.3. Genale Radiation. Among seven ecomorphs, some were significantly divergent from 327 others in δ^{15} N values (eight pairwise comparisons), but not in δ^{13} C values (Table S4). The largest value of δ^{15} N was found in the Large-mouthed (piscivory) ecomorph (13.83±0.7 ‰) 328 (Figure 5, Table S4). The minimal values of $\delta^{15}N$ were detected in the Short and Generalized 329 330 ecomorphs (11.15±0.5‰ and 11.23±0.8‰, respectively). The range between largest and lowest mean δ^{13} C values was small, from -19.87±0.8‰ (Short) to -20.69±1.0‰ (S-Scraper). 331 The total divergence between ecomorphs was lower in $\delta^{15}N$ values than in both the Didessa 332 and Gojeb radiations. In particular, the highest difference between ecomorphs in mean $\delta^{15}N$ 333 values was 2.73‰, and only 0.82‰ in δ^{13} C values. In the isotopic niches the most divergent 334 335 ecomorph was the Large-Mouthed, which overlap with other ecomorphs ranged from 0.06% 336 (vs. Short) to 38 and 43% (vs. benthophagous S-Scraper). Among other comparisons a low 337 overlap was detected between S-Scraper and Short (7 and 8%) (Supplementary File S4).

338 *3.2.4. Sore Radiation.* Among four ecomorphs, a few were significantly divergent in δ^{13} C 339 values only (two pairwise comparisons - Generalized vs. Thick-Lipped and Generalized vs. 340 V-Scraper; Table S4). The highest difference between ecomorphs in δ^{15} N values was 0.85‰ 341 only, while in the mean δ^{13} C values it was 1.86‰. The isotopic niches greatly overlapped 342 (67...82%) in all comparisons (Supplementary File S4).

3.3. Gut Length. The relative gut length varied significantly within each radiation except for
the Sore River (Figure 6). In three of four radiations, there were significant differences
between some ecomorphs (Figure 6; Supplementary File S5). Longest guts (up to 548-692%)

SL) were detected among V-scrapers (in the Didessa, Genale, and Gojeb Rivers). The largemouthed ecomorphs showed shortest guts (highest values were up 194-228% SL) except for the Sore River, where it reached 251±75%. Other ecomorphs had intermediate gut length; sometimes they differed from each other like Generalized and Thick-lipped ecomorphs in the Gojeb River (Figure 6). In spite of diverse mouth phenotypes in the Sore River, the ecomorphs were not divergent from each other representing the middle length of the gut (means varied within 251-290% SL).

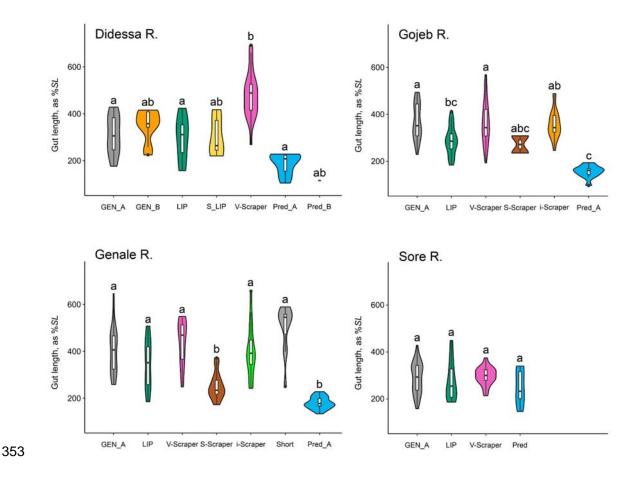


Figure 6. Violin plots of relative gut length distribution in sympatric ecomorphs from the four riverine radiations of the *Labeobarbus*. Min-max values (whiskers), 1st and 3rd quartiles (white vertical bars), median values (black horizontal bars), and outliers (black points) are indicated. Letters above the violin plots indicate significant differences between ecomorphs

(p < 0.05, Kruskal-Wallis test with Dunn's post hoc test). Abbreviations of the	358 (of the ecomorphs
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- 359 GEN_A Generalized A; GEN_B Generalized B; LIP Thick-lipped; S_LIP Semi-
- thicklipped; Pred_A Large-mouthed A; Pred_B Large-mouthed B.

361

362 4. Discussion

363 Our results show that parallel trophic diversifications occurred within the polyploid lineage, 364 the genus Labeobarbus. These diversifications differ in degrees of specialization within the 365 same set of core mouth phenotypes (generalized, thick-lipped, scraping, and large-mouthed), 366 indicating various stages of the process. We discuss the results obtained in the context of 367 trophic specializations of certain mouth phenotypes, the preadaptive nature of the ancestral 368 discrete mouth polymorphism, which likely originated via allopolyploidzation of the 369 Labeobarbus lineage, and attempt to reconstruct the process of trophic diversification by 370 analyzing four repeated cases discovered in the riverine environment of the Ethiopian 371 Highlands.

372 *4.1. Trophic resource partitioning and trophic specializations.*

373 Our study revealed that differences in diet between ecomorphs of the *Labeobarbus* in each 374 river were generally confirmed by SI analysis. For instance, the large-mouthed ecomorphs 375 which had piscivorous diet were also significantly enriched in δ^{15} N values (Figures 4 and 5). In contrast, the ecomorphs with detritivorous mode of feeding had lowest $\delta^{15}N$ values. The 376 periphython feeder that have been recorded in the Didessa River had a high level of $\delta^{15}N$ 377 378 values comparable with such in piscivorous ecomorphs but was characterized by enriched δ^{13} C values (Figure 5). The invertivorous ecomorphs whose diet was based on the larvae of 379 amphibiotic insects had intermediate levels of $\delta^{15}N$ values between detritivores and 380

piscivores. The data on gut length were less informative although the piscivorous ecomorphs had significantly shortened gut (Figure 6) that is in line with literature data for both *Labeobarbus* (Nagelkerke et al., 1994; Levin et al., 2019) and other fishes (e.g., Hugueny & Pouilly, 1999; Wagner et al., 2009). We summarized obtained data on the trophic specialization of sympatric ecomorphs from all rivers based on the diet, gut length, and SI composition in Table 2.

387

Table 2. Correlation between mouth phenotype and trophic specialization in the *Labeobarbus*spp. from four riverine radiations. The expected trophic niche is based on prediction from the
mouth phenotype and published data on trophic ecology*. Observed trophic niche - data

391 obtained in this study.

Mouth phenotype	Expected	River and observed trophic niches				
(core/subtype) trophic niche		Didessa	Gojeb	Genale	Sore	
1. Generalized A	Omnivore	Detritivore	Detritivore	Detritivore	Detritivore	
2. Generalized B	Omnivore	Detritivore- macrophyto phage	-	-	-	
3. Thick-lipped	Benthic invertivore	Detritivore- invertivore	Invertivore- detritivore	Detritivore	Detritivore	
4. Thick-lipped / Semi-thicklipped	Invertivore ?	Detritivore- macrophyto	-	-	-	

		phag			
5. Scraping / V-	Periphyton	Periphyton	Macrophyto	Macrophyto	Detritivore-
Scraper	feeder	feeder	phage	phage	macrophyto
					phage
6. Scraping / S-	Periphyton	-	Benthic	Benthic	-
Scraper	feeder		invertivore	invertivore	
7. Scraping / i-	Periphyton	-	Macrophyto	Detritivore	-
Scraper	feeder?		phage		
8. Large-mouthed A	Piscivore	Piscivore-	Omnivore-	Piscivore	Detritivore-
		omnivore	piscivore		macrophyto
					phage
9. Large-mouthed B	Piscivore	Piscivore?	-	-	-
No. trophic niches	-	5	5	4	2

392 * References used: Matthes, 1963; Nagelkerke et al., 1994; Golubtsov, 2010; Levin et al., 2019; Teshome et al.,
393 2023; Levin et al., in press.

394

Results of this study suggest five main trophic strategies among sympatric ecomorphs of the *Labeobarbus*: i) detritivory, ii) macrophytophagy, iii) invertivorous benthophagy, iv) periphyton feeding, and v) piscivory. Apart those, some ecomorphs had mixed modes of feeding - e.g., detritivory-macrophytophagy, detritivory-invertivory, and piscivory-omnivory (also with inclusion of a noted portion of detritus). Their more flexible

400 trophic specialization might be an adaptive strategy for living in mountain rivers with 401 unstable hydrological regimes (Jepsen & Winemiller, 2002). Although the core of mouth 402 phenotypes (generalized, thick-lipped, scraping, and large-mouthed) was the same in four 403 riverine radiations of the *Labeobarbus*, some phenotypes display more diversified sets of 404 ecologically relevant mouth subtypes. Scraping mouth phenotype was represented by three 405 subtypes, which occupy four different trophic niches (periphyton feeding, detritophagy, 406 invertivorous benthophagy, and macrophytophagy). It is an outstanding example of trophic 407 diversification of scraping mouth phenotype that previously was considered as adapted for 408 feeding via scraping the algal periphyton from stones and rocks (Matthes, 1963; see also 409 Vreven et al., 2016). Recent studies uncovered similar patterns of diversifications within 410 other phylogenetically distant cyprinid lineages bearing the scraping mouth phenotype - the 411 Garra and Schyzopygopsis, in particular (Komarova et al., 2021; 2022; Levin et al., 2021a). 412 Contrary to the Labeobarbus radiations that likely initiated from the generalized ancestor as 413 most ubiquitous throughout the generic range, the highly specialized scraper lineages (Garra 414 and Schyzopygopsis) could give a trophic radiation outside the ancestral narrow 415 specialization.

416 The Large-mouthed phenotype of the Labeobarbus demonstrated further sub-417 specialization within the adaptive zone of the piscivory in the Didessa River while in other 418 rivers this phenotype had only one feeding strategy that varied from obligate piscivory 419 (Genale) via piscivory-omnivory (Gojeb) to non-piscivory mode of feeding (Sore). The sub-420 specialization of the Large-mouthed phenotype in the riverine environment is similar in some 421 extent to the more diversified set of specializations discovered in Lake Tana where up to 422 seven sympatric large-mouthed species/ecomorphs were described (Nagelkerke et al., 1994), 423 whose piscivory specialization subdivided in benthic and pelagic zones (de Graaf et al., 424 2010).

425 A large incongruence between mouth phenotype and expected trophic niche is noted 426 not only for the trophic specialists. The Generalized ecomorph, including Ethiopian species 427 and populations, was considered omnivorous (Matthes, 1963; Teshome et al., 2023), but 428 according to our data it is detritivorous in all studied rivers (Levin et al., 2019; 2023 - in 429 press; this study). At the same time further diversification within Generalized mouth 430 phenotype (A and B subtypes) have been found. It is the emergence of an ecomorph with 431 horseshoe shape of the lower jaw without mental lobe and deep body in the Didessa River (Generalized B – Figure 2B). This ecomorph had lowest both $\delta^{15}N$ and $\delta^{13}C$ values among 432 433 other sympatric ecomorphs occupying a niche of detritivore-macrophytophage (Figures 4-5).

Noticeably, a detritus was the main food not only for generalists but often for thicklipped phenotype and also for all ecomorphs from the Sore River regardless their mouth phenotypes. We consider detritus as the most available and permanent trophic resource in riverine ecosystems of the Ethiopian Highlands with an unstable hydrological regime. In fact, it is a core food for the Generalized ecomorph and an important food resource for other ecomorphs.

440 Apart from the core mouth phenotypes, the intermediate ones were also found. For 441 instance, i-Scraper phenotype is intermediate between the Scraping and Generalized. Its 442 hybrid origin in the Genale River (by S-scraper and Generalized) was confirmed by genetic 443 data (Levin et al., 2019). Isotopic niche of i-Scraper from the Genale was between parental 444 species (Figure 5) but the gut content was similar to sympatric Generalized ecomorph. 445 Supposedly, i-Scraper intermediate phenotype might have its own trophic sub-niche. Another 446 example of intermediate mouth phenotype that seemingly occupies its own sub-niche refers 447 to the Semi-thicklipped ecomorph (see Figures 4-5) in the Didessa River. Obtained results are 448 consistent with the syngameon hypothesis, proposing that hybridization between members of

449 the radiation can promote further niche expansion and diversification (see Seehausen, 2004; 450 Frei et al., 2022). Intermediate mouth phenotypes were reported among the Labeobarbus – 451 between scraping 'Varicorhinus'-like and generalized phenotypes by Banister (1972) and 452 Nagelkerke and Sibbing (1996) as well as between scraping and thick-lipped mouth 453 phenotypes by Vreven et al. (2019). In the last case the intermediate phenotype resembled the 454 generalized one (Generalized B in our study). Authors revealed a hybrid nature of this 455 phenotype by genetics (Vreven et al., 2019) but there was no data on trophic ecology or 456 specialization. Actually, the recruiting of novel trophic niches via hybridization is an 457 intriguing issue that is yet weakly studied (but see experimental study of Selz and Seehausen, 458 2019).

In summary, the *Labeobarbus* trophic radiations involve a mixture of pelagic predator (piscivore) and a wide assortment of benthic-oriented specialists. The benthicpelagic habitat axis is most frequent in cyprinid and cichlid adaptive radiations in both lakes and rivers (Nagelkerke et al., 1994; Cooper et al. 2010; Levin et al., 2019; Burress et al., 2023).

464 *4.2. Decoupled form and function – preadaptive phenotypes and their functionalization.*

465 Our results show that the same mouth phenotypes in the *Labeobarbus* may occupy different 466 trophic niches. A mismatch between expected and observed trophic specializations may have 467 different nature: i) specialized phenotype has unspecialized (generalized or omnivorous) 468 feeding mode that is known as Liem's paradox (Liem, 1980; Robinson et al., 1998), and ii) 469 specialized phenotype has unexpected (biased) trophic specialization possibly due to 470 insufficient knowledge on trophic specializations in certain lineages. We focus in this study 471 on the first phenomenon. Liem's paradox was established for cichlid fishes (e.g., Liem, 1980; 472 Sturmbauer et al., 1992; Wagner et al., 2009; Binning et al., 2009; Torres-Dowdall & Meyer,

473 2021) but it was also detected in other taxonomic groups (e.g., in cyprinids – Lammens et al., 474 1991; Levin et al., 2021a; Komarova et al., 2022). The mismatch between form and function 475 might be i) a temporary phenomenon for proper specialist explained by plasticity of its diet in 476 some circumstances (ecological release that may be a base for evolutionary re-specialization 477 - see above-mentioned example of highly specialized scraping periphyton feeders of the 478 genus Garra that could re-specialize in other trophic specializations – Komarova et al., 2022) 479 or ii) heterochronous decouple of form and function when the emerging phenotype is not yet 480 functionalized, i.e. phenotype is preadaptive (expectation on its trophic adaptation based on 481 phenotypic features) but not vet involved in trophic resource partitioning. This is exactly the 482 case of the mouth diversity in the Sore River, where detritus was a main food for all mouth 483 phenotypes including the Large-mouthed, i.e. mouth divergence was uncorrelated with the 484 use of food resources. In other words, mouth polymorphism in the Sore River is ecologically 485 almost non-functional compared to other riverine diversifications. This contradicts the 486 'ecological theory' of adaptive radiation in neglecting the 'habitat first rule' in particular (see 487 Schluter, 2000) but fits the flexible stem hypothesis (Wund et al., 2008; Gibert, 2017) or its 488 variant known as the 'plasticity-first' evolution (Levis & Pfennig, 2016; 2019). According to 489 this hypothesis, an adaptive phenotypic plasticity in an ancestral population could precede 490 adaptation to a new environment through the process of genetic assimilation. Remarkably, the 491 case of decoupled form and function in the Labeobarbus is shared to some extent with such 492 in riverine adaptive radiations in the pike cichlids, genus *Crenicichla* (Burress et al., 2023) 493 that may be a general feature for young radiations in riverine environments.

As already mentioned, the repeated mouth polymorphisms of the *Labeobarbus* may be predetermined due to its polyploid origin (Yang et al., 2022). Its maternal lineage is characterized by generalized / thick-lipped mouth polymorphism widely distributed among Torinae in the genera *Tor* and *Neolissochilus* (Hoang et al., 2015; Walton et al., 2017) – so

498 the generalized/thick-lipped polymorphism is persistent within a Torinae lineage. Paternal 499 lineage referred to contemporary genus *Cyprinion* is represented by scraping mouth 500 phenotype (Coad, 2021). Hence, mouth polymorphism of the *Labeobarbus* is ancestrally 501 heritable and re-evolved under particular ecological circumstances. Given this, the thick-502 lipped and scraping mouthed phenotypes of the *Labeobarbus* in the Sore River are seemingly 503 'preadaptive' upon emergence *de-novo*.

504 Modular-assembling genome of the Labeobarbus consisted of the different 505 subgenomes (Yang et al., 2022) that is able to produce discrete mouth phenotypes inherent to 506 parental lineages is formally still within 'flexible stem hypothesis'. However, from another 507 hand, it might be also in frame of 'transporter process' (Schluter & Conte, 2009; Marques et 508 al., 2019; Martin & Richards, 2019) meaning that the adaptive alleles and genetic 509 architectures differentiating each species within a rapid radiation are older than the radiation 510 itself. The oldest paleorecords of the Labeobarbus in East Africa are dated by Late Miocene 511 (Stewart & Murray, 2017). Some trophic specialists, for instance, with scraping phenotype 512 are remarkably older than the recently emerged repeated radiations in the Ethiopian 513 Highlands under consideration. Their age is dated as Pleistocene according to molecular 514 clocks (Beshera et al., 2016). It suggests that similar mouth diversification might evolved 515 repeatedly before the diversifications we observe in this study. Therefore, the genomes of the 516 modern Labeobarbus might be rather experienced, i.e. may have the standing genetic 517 variation contributing to the ecological speciation and diversification. Regardless which is the 518 certain prerequisite for adaptive radiation of the Labeobarbus - 'flexible stem' or 'transporter 519 process' (or their combination) – it might significantly facilitate the diversification.

520 *4.3.* Possible scenario of evolution of trophic radiation: from incipient to matured

521 Our results demonstrate that four riverine assemblages of the *Labeobarbus* in the Ethiopian 522 Highlands being similar in the core mouth phenotypes and direction of trophic diversification 523 are at various stages of the evolution. The number of mouth phenotypes together with 524 subtypes and their ecologic functionality in the *Labeobarbus* varied significantly from river 525 to river. We tried to range the cases from simplest to most matured based on the obtained 526 results – diversity of ecomorphs/mouth phenotypes and their functionality (as estimated by 527 the gut content and isotopic niche) within each radiation (Table 3; Figures 4-5).

Table 3. Ranking the diversifications by number of ecological niches and other aspects of trophic differentiation. Number of points in rows 3 and 4 corresponds to the number of significant differences in pairwise comparisons between sympatric ecomorphs (taken from Supplementary File S4).

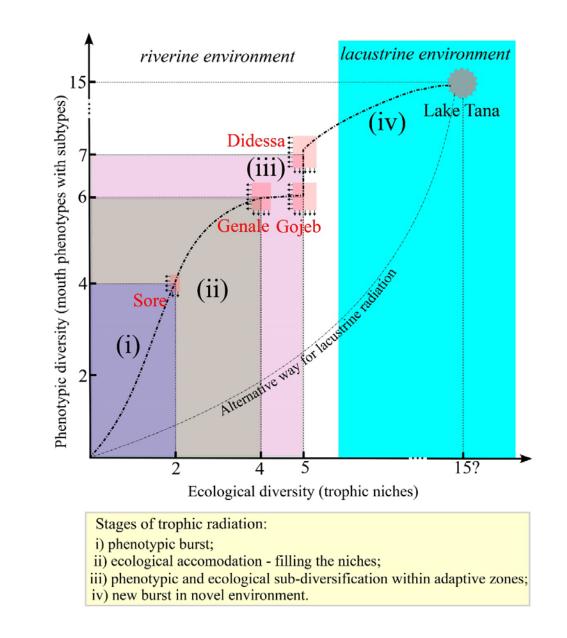
Nos.	Parameters	Sore	Genale	Gojeb	Didessa
1	Nos. mouth phenotypes	4	6	6	7 ¹
2	Nos. trophic niches ²	2	4	5	5
3	Nos. dif. in δ^{15} N	0	8	5	6
4	Nos. dif. in $\delta^{13}C$	2	0	6	6
5	Nos. trophic levels ³	0.36	1.18	1.45	2.0

¹Eight ecomorphs were detected in the Didessa but only seven were included in the study due to absence of data on the exceptionally rare Large-mouthed ecomorph. ² Number of trophic niches is taken from Table 2. ³ One trophic level is arbitrary taken as 2.3‰ between mean δ^{15} N values according to McCutchan et al. (2003) and Vanderklift & Ponsard (2003): the number of trophic levels was counted by dividing the difference between minimal and maximal mean δ^{15} N values within a pool of sympatric ecomorphs by 2.3.

The trophic diversifications could be ranged from the most undifferentiated in the Sore River
via the Genale and Gojeb Rivers to the most diversified in the Didessa River (Table 3, Figure
7). Next stages of the trophic diversification can be provisionally outlined (see scheme on
Figure 7):

- 542 (1) Burst of mouth polymorphism. A set of core mouth phenotypes quickly emerges but 543 their ecological functionalization is immature (preadaptive phenotypes). 544 Morphological phenotypes are trophically irrelevant or there is a weak phenotype-545 trophic correlation (Liem's paradox). At this stage phenotypic diversity strongly 546 dominates over the ecological diversity - see the case of the Sore River (four 547 phenotypes vs. two trophic niches).
- 548 (2) Filling the trophic niches (ecological accommodation). Preadaptive phenotypes began
 549 to be functionalized by occupation of the trophic niches best matched to the mouth
 550 type. This process might be named as ecological accommodation of phenotypic trait –
 551 analogously to term 'phenotypic accommodation' sensu West-Eberhard (2003) The
 552 numbers of phenotypes and trophic niches are almost equal. Probably it is a very
 553 rapidly realizing process else it is hard to explain what are reasons to sustain non554 functional mouth polymorphism.
- 555 (3) Continued diversification within the core mouth phenotypes with corresponding 556 increase of the ecological diversity. Upon filling the most evident trophic niches, a 557 further sub-diversification occurs in some radiations. It might go by various scenarios 558 - via ecological sub-specialization of certain phenotypes or via hybridization and 559 functionalization of intermediate phenotypes. At matured stages of the diversification, 560 the biotic interactions may serve as an additional ecological opportunity ('diversity 561 begets diversity' – Martin & Richards, 2019). Significant increase of the populations 562 size during morpho-ecological diversification (due to utilization of various trophic

563	resources in a river) makes the piscivory strategy more reliable that may result in sub-
564	diversification within this adaptive zone - like in the Didessa River (see also
565	cascading speciation - Broderson et al., 2017; Bracewell et al., 2018).
566	(4) A new diversification burst provoked by colonization of the novel environment might
567	occur as exemplified from the lacustrine burst of the Labeobarbus in Lake Tana
568	(Ethiopia) resulted in evolution of 15-16 species/ecomorphs (Nagelkerke et al., 1994;
569	Mina et al., 1996; Sibbing, Nagelkerke, 1998; Nagelkerke et al., 2015; Beshera and
570	Harris, 2023). It is still unknown whether Lake Tana radiation evolved from a single
571	generalized ancestor or stemmed from already diversified set of riverine radiation.



573

Figure 7. Schematic reconstruction of process of morpho-ecological diversification of the *Labeobarbus* in the Ethiopian Highlands. The number of little arrows at each box correspond
to the numbers of mouth phenotypes (horizontal arrows) and numbers of trophic niches
(vertical arrows).

578

579 The most immature diversification has been detected in the Sore River where a form580 (mouth phenotype) and its ecological function were largely decoupled. The ecologically non-

581 functional or non-matured mouth polymorphism in the Sore River might be explained by 582 several reasons. First, it might be due to a very young age of this radiation (at stage of 583 incipient diversification) that is supported by low genetic diversity and absence of mtDNA 584 haplotype sorting (Levin et al., 2020). Another reason refers to insufficient ecological 585 opportunities in the river to realize the existing potential (mouth polymorphism). 586 Heterogeneous and unpredictable environment in the rivers (flow regimes, turbidity, depth, 587 water chemistry, temperature, food availability, etc.) may weaken the divergent selection 588 necessary to drive specialization, instead favoring opportunistic ecological roles such as 589 detritivory-omnivory (Jepsen and Winemiller, 2002; Burress et al., 2023). But the Sore River 590 is a home for another bright adaptive radiation of hillstream cyprinid fish of the genus Garra 591 that is represented by six genetically, morphologically and trophically divergent young 592 species (Golubtsov et al., 2012; Levin et al., 2021a; Komarova et al., 2022). Possible 593 overlapping in the trophic specializations (predator, periphyton feeding, benthic invertivore) 594 with more advantageous cyprinid radiation in shared environment can acutely suppress or 595 constrain the diversification process in the Sore population of the Labeobarbus resulting in 596 freezing the non-functional mouth polymorphism stage for a while. Simultaneously, other 597 studied riverine populations of the *Labeobarbus* have represented more advanced trophic 598 diversification stages in absence of other sympatric radiations (as far as we know) and 599 characterized by matured or ecologically functional mouth phenotypes.

600

Conclusions

Using the polyploid African barbs of the genus *Labeobarbus* as model, we showed the
parallel morpho-ecological diversifications being similar in this phenomenon with other
prominent examples of adaptive radiation among vertebrate animals like cichlids (Seehausen,
2015; Torres-Dowdall & Meyer, 2021; Burress et al., 2023) and Caribbean *Anolis* (Losos,

605 2009; Stroud & Losos, 2020). The most striking difference from above-mentioned examples 606 is the sharply preadaptive nature of inherited mouth polymorphism as a starter of radiation. 607 Likely, the diversification of the *Labeobarbus* starts from the onset of genomic pre-existing 608 templates of mouth polymorphism. The emerged mouth phenotypes are preadaptive but not 609 (or weakly) functional at the incipient stage of trophic diversification demonstrating 610 decoupled form and function. Enhancement of ecological functions of preadaptive 611 phenotypes (ecological accommodation) increases upon filling the trophic niches and result 612 in maturation of trophic specialization that is often continued by further sub-diversification of 613 mouth types and sub-specialization within previously occupied adaptive zones. The 614 preadaptive mouth polymorphism can be considered a key innovation of the Labeobarbus 615 that makes the bridge between phenotypic diversification and ecological opportunities 616 remarkably shorter that might accelerate diversification rates. It apparently explains why this 617 lineage overcomes the obstacles in unstable riverine environments unfavored for adaptive 618 radiations by other fish lineages.

619

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634	Data availability. Raw isotopic data are available as Supplementary data.
635	Supplementary data. Supplementary data to this article can be found online.
636	Competing interests. The authors declare that they have no competing interests.
637	Authors' contributions. B.A.L. and A.S.G. conceived and designed the research, conducted
638	field surveys and collected samples; A.S.K. and A.V.T. performed the stable isotope samples
639	preparation and their analysis; A.S.K. performed diet analysis; All authors wrote the
640	manuscript. All authors read and approved the final version of the manuscript.
641	
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