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        The scales of variability of stream fish assemblages
                        at tributary confluences
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            Running head: organization of stream fishes at tributary confluences
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#### Abstract

Tributary confluences play an important role in the dispersal of organisms, and consequently, in shaping regional scale diversity in stream networks. Despite their importance in dispersal processes, little is known about how ecological assemblages are organized in these habitats. We studied the scales of variability of stream fish assemblages over three seasons using a hierarchical sampling design, which incorporated three tributaries, three sites at the mouth of each tributary and using four sampling units at each site. We found strong scale dependent variability in species richness, composition and relative abundance. Most of the variation was accounted for by the interactive effect of season, between stream and between site effects, while habitat structure of the sampling units had a relatively minor role. Species richness showed a continuous decrease from the mainstem river in most cases, while species composition and relative abundance changed less consistently along the longitudinal profile. Consequently, we found that not only the junctions presented a strong filter on the species pool, but some species were filtered out if they passed this critical habitat bottleneck. Overall, our results suggest high variability in fish assemblages across multiple scales at tributary confluences. Environmental management should take a more critical care on the filtering role of tributary confluences in species dispersal, for better understanding patterns and processes in the branches of dendritic stream networks.


Keywords: stream networks; tributaries; ecotones; habitat bottlenecks; environmental filtering; dispersal

Introduction
A central task of community ecology is to disentangle how environmental filtering processes and biotic interactions select the assembly of local communities from the regional species pool (Vellend 2010). The act of environmental filtering tends to be the most prominent at the interface of those habitat patches which largely differ in their environmental conditions (ecotones). Characterizing how species disperse through these habitat bottlenecks and distribute themselves among patches is important, among others, for a more regional scale understanding of the organization of ecological communities (Risser 1995).

In dendritic stream networks tributary confluences (or junctions) provide one of the most characteristic examples of within stream ecotones (Ward and Wiens 2001). Here, abrupt changes often happen in width, depth, sediment characteristics and water quality along the longitudinal profile of the stream (Rice et al. 2008). Tributary confluences can also increase physical heterogeneity (e.g. in water and sediment characteristics) in the recipient river which have important consequences in channel and floodplain morphology (Benda et al. 2004; Rice et al. 2006). Confluences thus serve as key habitat structures, which may influence dispersal processes for stream organisms in both directions (i.e. from the tributary to the main channel and vica versa). However, despite their importance relative to their small spatial extent, ecological studies in stream networks concentrated mainly on understanding patterns and processes in the branches and not in the junctions (sensu Grant et al. 2007).

Studies which would specifically focus on the role of confluences considered mainly changes in the composition of the main river assemblages below tributary mouth sections (see e.g. Rice et al. 2001; Kiffney et al. 2006; Milesi and Melo 2014). Studies, which would have specifically considered the upstream filtering role of tributary mouths are especially rare (but see e.g. Grenouillet et al. 2004; Hitt and Angermeier 2008), and in most cases used only indirect evidence for the permeability vs. resistance function of the tributary-mainstem ecotone. For stream macroinvertebrates Beckmann et al. (2005) highlighted that not only tributaries influence the composition and assemblage structure of the main river, but that mainstem assemblages can intensively use the tributary mouth for habitat. For stream fish assemblages Thornbrugh and Gido (2010) showed an abrupt change in assemblage composition between mainstem river sites and tributary sample sites above confluences, followed by a gradual taxonomic change with increasing upstream distance. Overall, these studies suggest that confluences play a strong filter on the species pool of mainstem rivers and has an overarching role on assemblage organization of tributary streams.

Stream systems show extreme heterogeneity at a hierarchy of spatial scales from microhabitat to the catchment (Hildrew and Giller 1994; Poff 1997). Determining the scales at which stream organisms varies the most is essential for understanding the predictability of assemblages and providing implications for management (Heino et al. 2004, Erős and Schmera 2010; Ligeiro et al. 2010). Variability of stream assemblages can be especially high at tributary confluences due to the importance of these habitats (i.e. crossroad function) in the dispersal of organism in the dendritic network. Quantifying the importance of scale related variability at tributary confluences is thus important for understanding the variability of
assemblages upstream in the branches. For example, if one finds large between stream variability already at the tributary mouth section, this could (partly) explain the differences in the composition of assemblages among the branches, at least for those stream organisms, which dispersal is restricted exclusively to within stream movement. However, the relative role of different scales (e.g. between vs within stream effects, mesohabitat level heterogeneity within sites) is largely unknown on the assembly of stream organisms at tributary confluences.

In this study, we examine the filtering role of confluences on stream fish assemblages. Our goal was to determine the major scales of assemblage variation at tributary mouth sections. Specifically, we used a hierarchical (i.e. nested) sampling design to assess the role of season, stream identity (i.e. between stream differences), spatial position at the tributary mouth section (between reach differences), and habitat structure (between sampling unit or mesohabitat level differences) on the structure of fish assemblages using survey data from three tributary mouth sections of a lowland medium size river. Such studies have importance for a more complete understanding of the role of the junctions in the organization of assemblages in stream networks.

## Materials and Methods

Study area
Our study streams were the 33.3 km long Hajagos (catchment area: $236.9 \mathrm{~km}^{2}$ ), the 57.1 km long Gerence ( $310.3 \mathrm{~km}^{2}$ ) and the 45.4 km long Sokoroi ( $349 \mathrm{~km}^{2}$ ) streams. These streams are the right side tributaries of the Marcal River, a 100.5 km long lowland river, which is situated in North-Western Hungary. At the middle and downstream part of the Marcal, the average wet with is $10-15 \mathrm{~m}$ and the mean annual discharge is $6.8 \mathrm{~m}^{3} \mathrm{~s}^{-1}$. The tributary mouths of the three studied streams are on average 20.3 km from each other along the Marcal. In their middle and downstream sections, the streams run on an agricultural landscape. They have of lowland character, with low flow, fine substrate (predominantly sand, silty sand, and fine and coarse gravel) and with no clear riffle-pool structure. At the most downstream (i.e. tributary mouth) section the mean channel width, velocity and depth varied between 2.89 and 5.19 m , 6.1 and $13.4 \mathrm{~cm} \mathrm{~s}^{-1}, 43.5$ and 88.1 cm , respectively (see Table 1 for details). Emergent (e.g. reed Phragmites australis, broadleaf and narrowleaf cattail Typha latifolia and Typha angustifolia, sweet-grass Glyceria sp.) and floating (yellow water-lily Nuphar lutea, duckweeds Lemna sp.) macrophytes can be abundant at some sites.

Fish and habitat sampling
We chose three sampling sites per stream. Their downstream end point situated at $0 \mathrm{~m}, 500 \mathrm{~m}$ and 1000 m from the tributary mouth (Fig. 1). Each site was 200 m long and were further divided into four 50 m long sampling units, to examine the effect of habitat structure on fish assemblage characteristics at a finer scale. We sampled fishes and characterized the habitat during three sampling periods in 2013: (1) spring (end of April, early May), (2) summer (early

July), and (3) autumn (early October). This sampling effort yielded a total of 108 samples (3 seasons $\times 3$ streams $\times 3$ sites per stream $\times 4$ habitat units per site).

We surveyed the 200 m long sites by wading, single pass electrofishing using a backpack electrofishing gear (IG200/2B, PDC, 50-100 Hz, 350-650 V, max. 10 kW ; Hans Grassl GmbH , Germany). The 50 m long sampling units were block netted before electrofishing, with nets of 2 mm mesh size anchored with chain to the bottom. Pulsating direct current with a frequency of $75-100 \mathrm{~Hz}$ and a voltage of $200-300 \mathrm{~V}$ was used. The 2 m long catcher rod had a ring shaped anode with a diameter of 40 cm equipped with a net (mesh size 6 mm ). The sampling crew consisted of 3-4 people: the electrofisher operator who effectively caught the fish and handled the machine, a netter who helped to catch escaping or unseen fish and one or two helpers who carried the buckets on the bank, took care of the fish and measured their standard length. The team electrofished the whole stream width while moving slowly upstream and trying to catch all fish in the study area. All fish were placed in large containers filled with water and released back to the stream after identification and length measurement.

Transect based measurements of habitat data were taken at each 50 m long unit after fish sampling. At each unit, four transects were placed perpendicular to the main axis of the stream at equal distances from each other. We measured wet width of the channel with a tape measure, depth and current velocity with a meter stick and a water velocity meter (FP101 Global Flow Probe, Global Water Instrumentation Inc., Gold River, CA, USA), respectively, at five equally spaced points along each transect. We also recorded aquatic vegetation type and substrate type (see Table 1) at every transect point. We calculated the mean values of the aforementioned environmental variables and the percentage occurrence of the categorical variables (i.e., substrate type and aquatic vegetation type) to characterise habitat features at the sample unit scale.

We also used fish monitoring data from the Marcal River for comparing similarities and differences in fish assemblages between the mainstem Marcal and the tributary fish assemblages. The river was sampled from a boat using the same electrofishing device and its accessories as was used for the tributary streams. To allow effective manuovering in this shallow and vegetated river, a small rubber boat (Yamaha 300S) was used. The crew comprised two persons: one for catching the fish with the hand-held anode $(2.5 \mathrm{~m}$ long pole with a net of 40 cm diameter, mesh size 6 mm ) and one for driving the boat. Two 400 m long reaches (altogether 800 m long sections) were electrofished in the vicinity (i.e. within 5 km ) of each tributary at our standard monitoring sites in this river. Continuous electrofishing was carried out, by dipping the anode into the water at approximately 2 m long intervals and pulling the anode toward the boat, while moving slowly downstream with the flow and controlling the boat by an oar if it was necessary. As for the tributary streams fish were released back to the river after identification and length measurement.

Statistical analysis

We used linear models in a nested design to assess the effect of (1) seasons, (2) tributary streams within seasons, (3) sites within tributary streams and seasons, and (4) habitat data nested within seasons and tributary streams on (1) species richness and (2) ecological dissimilarity of the assemblages compared to the main river. For dissimilarity index the Jaccard and the Bray-Curtis indices were used for species composition and relative abundance data (see Legendre and Legendre 1998), respectively.

We used principal component analysis on the variance-covariance matrix of the recorded habitat data (PCA) to characterize the physical structure of the habitat units, and used the first three component scores (hereafter habitat PC scores) to quantify habitat data in the nested model. The advantage of this procedure is that it reduces the number of the original habitat variables which usually are correlated with each other. The correlation between the habitat variables could be a nuisance factor in linear models, which phenomenon is known as multicollinearity (Graham 2003; Dormann et al. 2013). Here, the first three principal components of the PCA explained $24.4 \%, \mathrm{x} \%$ and $\mathrm{Y} \%$ of the variance, which justify their inclusion in the models. In PC1, habitat units with the strongest negative loadings were relatively deep and wide with relatively high velocity, silty-sand substrate and relatively dense submerged macrophyte vegetation. Units with strong positive loadings were relatively shallow with low velocity and relatively dense emergent vegetation (Table 5). In PC2 the habitat gradient was mainly characterized by differences in substrate composition, and especially sandy silt (negative loadings) and fine gravel (positive loadings). Although biologically less interpretable, abiotic differences between habitat units along PC3 were further refined by substrate composition and differences in vegetation structure (Table 6).

For a more complete understanding of confluence effects, we examined changes in species richness, and dissimilarity in species composition (Jaccard index) and relative abundance (Bray-Curtis index) from the mainstem Marcal River for each stream and in each season. Spearman correlation tests were used to test whether the values of these assemblage level variables showed significant decay with rank distance upstream from the mainstem at the level of habitat units ( $\mathrm{n}=12$ in each test).

Then, focusing exclusively on the tributaries, we used principal coordinate analysis (PCoA) to reveal patterns in species composition and relative abundance between seasons, streams and sites. Here, data from individual habitat units were pooled for the ease of visualization. The Jaccard and the Bray-Curtis indices were used for species composition and relative abundance, respectively. Finally, we used nested permutational analysis of variance (PERMANOVA) to identify the scales at which species composition and relative abundance of the tributary fish assemblages varied the most. As with the PCoA, the Jaccard and the Bray-Curtis indices were used for species composition and relative abundance, respectively.

All data analyses were performed in the R statistical environment ( R Core Team 2013).
PERMANOVA was conducted with 'adonis' function of the R package named vegan (Oksanen et al. 2013).

Results

Most species that occurred in the mainstem river appeared also in the tributaries (Table 2). In fact, only large bodied and/or strongly riverine species did not appear in the tributaries such as asp (Aspius aspius), barbel (Barbus barbus), European catfish (Silurus glanis), zingel (Zingel zingel). However, these species were rather rare in the mainstem river as well.

Species richness varied significantly at a variety of sampling scales in the tributaries (Table 3). Most of the variation was related to stream ( $36.4 \% ; \mathrm{P}<0.001$ ) and site level variability ( $19.3 \%$; $\mathrm{P}<0.001$ ), although seasonal differences also proved to be significant ( $7.5 \%$; $\mathrm{P}<0.001$ ). Species richness distributed relatively evenly amongstreams and sites in spring (Fig. 2), while it tended to decrease upstream in both summer and autumn. Species richness also varied among streams according to their upstream-downstream position in the mainstem in both summer and autumn. The most downstream Sokoroi stream had the highest species richness, while the most upstream Hajagos had the lowest. Interestingly, habitat structure (i.e. PC1-3) did not significantly influence species richness, although the overall contribution of the biologically interpretable most significant environmental gradients (i.e. PC1-3) were in fact higher than the effect of season (14.7\%).

Jaccard dissimilarity of species composition from the mainstem showed similar influence of sampling scales to species richness (Table 3). Most of the variation was related to stream ( $39.0 \% ; \mathrm{P}<0.001$ ) and site level variability ( $23.6 \% ; \mathrm{P}<0.001$ ), although seasonal differences $(3.2 \% ; \mathrm{P}<0.05)$ and the effect of habitat structure (i.e. $\mathrm{PC} 26.7 \% ; \mathrm{P}=0.026$ ) also proved to be significant. Dissimilarity from the mainstem was more even among streams and sites in spring (Fig. 3), while it tended to increase upstream among sites in the tributaries and among streams along their upstream-downstream position in the mainstem, both summer and autumn. For Bray-Curtis dissimilarity of relative abundance data seasonal changes proved to be the most influential ( $54.3 \% ; \mathrm{P}<0.001$ ). Clearly, dissimilarity from the mainstem increased from spring to autumn (Fig. 3), while the effect of hierarchical spatial levels proved to be low, albeit significant at the stream $(6.4 \% ; \mathrm{P}=0.003)$ and site $(12.4 \% ; \mathrm{P}<0.001)$ levels.

Species richness showed a continuous decrease from the mainstem river in most seasons and streams (Fig. 4; Table 4). However, species composition and relative abundance changed less consistently along the longitudinal profile. Dissimilarity in species composition generally increased from the mainstem in both summer and autumn, with the exception of the Sokoroi stream. On the contrary, dissimilarity in relative abundance showed a variety of patterns. For example, it decreased in the Gerence stream in both spring and summer, while it increased in the Sokoroi stream in autumn.

Species composition of the streams and sites within streams was determined by several species (Fig. 5). However, no clear pattern in assemblage composition was found regarding seasonal, between stream or between site differences. PERMANOVA analysis (Table 5) confirmed these findings and showed that relatively equal amount of variation of species composition was related to seasonal ( $14.6 \% ; \mathrm{P}<0.001$ ) and to between stream ( $22.4 \%$; $\mathrm{P}<0.001$ ) and between site ( $13.7 \% ; \mathrm{P}<0.001$ ) effects. Habitat structure also significantly influenced species richness (i.e. PC2 $6.5 \%$; $\mathrm{P}=0.018$ ), and the overall contribution of the
biologically interpretable most significant environmental gradients (i.e. PC1-3) was in fact higher $(15.8 \%)$ than the effect of season. Relative abundance data were mostly influenced by three dominant species (i.e. roach Rutilus rutilus, bitterling Rhodeus sericeus, pike Esox lucius; Fig. 5). Here, the greatest amount of variation in fish abundance data was related to the site scale ( $20.7 \% ; \mathrm{P}<0.001$ ), although seasonal ( $12.6 \% ; \mathrm{P}<0.001$ ) and stream level $(17.8 \%$; $\mathrm{P}<0.001$ ) effects, and the effect of individual environmental gradients ( $\mathrm{PC} 17.1 \% ; \mathrm{P}=0.007$; $\mathrm{PC} 26.4 \% ; \mathrm{P}=0.022 ; \mathrm{PC} 36.4 \% ; \mathrm{P}=0.021$ ) also proved to be significant (Table 5).

## Discussion

In this study we examined the scales of variation of stream fishes at tributary confluences and found that the majority of assemblage variation could be explained by the joint effect of different hierarchical (spatial and temporal) scales. In general, seasonal, between stream and site level effects proved to be the most important in explaining assemblage structure, whereas physical attributes of the sample units ( 50 m long each) had a relatively minor role. These results on the importance of season, spatial positioning in the stream network (between stream effects) and positioning within a stream (between site effects) may suggest that dispersal processes had critical importance for fish assemblage organization at tributary confluences, and will be discussed below in more detail.

Most of the studies that considered the role of spatial processes (e.g. dispersal limitation, neutral effects) in stream networks used a metacommunity perspective and examined the relative role of environmental factors and spatial positioning of the sampling sites at large spatial extents (i.e. within and among different subcatchments) (e.g. Mykrä et al. 2007; Erős et al. 2012). The logical consensus of these studies is that the role of spatial effects decreases at smaller spatial extents (Heino et al. 2012; Muneepeerakul et al. 2008). However, it has been also shown that spatial processes like dispersal limitation can be very important at smaller extents too, for example within and between branches of the dendritic network (Fagan 2002; Erős and Schmera 2010). Our field study shows that between stream effects can be significant even among streams with relatively similar environmental and topographic characteristic and spatial positioning along the mainstem river, which, to our knowledge, has not been addressed in other studies.

Between stream effects were high for species richness and dissimilarity of species composition from the mainstem river (Table 3). Specifically, we found a reverse pattern between species richness and dissimilarity in species composition from the mainstem. Species richness was the highest in the Sokoroi and lowest in the Hajagos stream, while dissimilarity in species composition from the mainstem was the lowest in the Sokoroi and highest in the Hajagos, at least in summer and autumn. Lower species richness and higher dissimilarity from the mainstem in species composition suggest lower colonization potential of the Hajagos from the Marcal River. Confluences thus served as natural habitat bottlenecks for stream fishes in a stream specific manner. The results may suggest the role of stochastic effects in the colonization of tributary mouths by different species from the mainstem and/or suggest the
role of tributary spatial position in the mainstem river (see Osborne and Wiley 1992), even if the tributaries were relatively close to each other.

Temporal variability also had some legacy on between stream differences in richness and dissimilarity of species composition from the mainstem. These differences were relatively minor in spring and showed consistent pattern in summer and autumn. Higher water levels in spring could diminish between stream differences in habitat availability of the junctions for mainstem fishes, since alluvium bars (i.e. sand and gravel bars) at the mouth may hinder upstream migration at low water periods. Higher movement activity of fish during spring (spawning migrations) may also dampened differences among the tributaries. Dissimilarity from the mainstem in relative abundance data supports this argument on seasonality. Dissimilarity values were the lowest in spring and the highest in autumn. In fact, dissimilarity from the mainstem was most related to variation among seasons for relative abundance data Table 3), reflecting seasonal differences in habitat availability and/or in movement activity for the dominant species (Gorman 1986; Roberts and Hitt 2010).

At a lower hierarchical level, all assemblage attributes varied significantly between sites, which further indicates the importance of spatial processes for stream fishes at very small spatial extents (here within a 1 km long section). Species richness generally decreased upstream at least in summer and autumn (Fig. 2), while dissimilarity in species composition increased from the mainstem. To our knowledge no study examined differences in fish assemblages from the mainstem at such a small spatial extent, but these data and studies at larger spatial extents (Hitt and Angermeier 2008; Thornbrugh and Gido 2010) indicate that not only the junctions present a strong filter on the species pool, but that some species are filtered out even if they passed through this most critical habitat bottleneck. Dissimilarity in relative abundance from the mainstem was less related to variation between sites, which, compared with results on composition, indicates the effect of some dominant species on these results. Some habitat generalist species, like roach, bitterling and bleak (Alburnus alburnus), which are dominant both in the mainstem river and in the tributaries were abundant at all sites and could increase assemblage similarity at the site level, similarly to their effect at the stream level. Patterns in distance decay from the mainstem (Fig. 4) confirmed site level changes, and indicated the most consistent pattern between seasons and streams for species richness (i.e. decrease in richness upstream) and the least consistent patterns for relative abundance data.

Analyses which did not consider similarities and differences from the mainstem, but focused on assemblage variability in the tributaries exclusively (i.e. PCoA and PERMANOVA), showed that relatively equal amount of variation was related to season, and between stream and between site variation for both compositional and relative abundance data. These results suggest strong context dependency in fish assemblage organization at tributary confluences, which may harden generalizations across sites, streams and seasons (see Heino et al., 2012 for a larger scale study). The relatively high amount of variation explained (66.4-71.0\%) suggest that the joint consideration of seasonal effects and between and within stream positioning of the sampling sites is critically important for better understanding assemblage organization, beside the consideration of habitat structure.

For the lowest hierarchical level (i.e. sampling unit scale), we found that habitat structure had low importance relative to seasonal, between stream, and between site variation. Habitat structure is clearly the most important factor (group) for the organization of stream fishes (Matthews 1998; Jackson et al. 2001). It seems however that larger scale factors (e.g. permeability of the junctions, dispersal ability of species) can also significantly influence the species pool of lower hierarchical levels (i.e. species composition and relative abundance of fishes at the meso-habitat level) and in case of relatively homogenous habitat units, they can override the effect of habitat structure (see Grönroos and Heino 2012 for a study on stream macroinvertabrates). In this lowland system, the units had relatively similar physical characteristics regarding width, flow, depth and substrate, although the type and coverage of macrovegetation provided some heterogeneity, which could influence the distribution of fish to some extent.

In conclusion, we found strong scale dependent variability in species richness, composition and relative abundance of stream fish assemblages at tributary confluences. Most of the variation was accounted for by the interactive effect of season, between stream and between site effects, while habitat structure of the sampling units (i.e. mesohabitat level structure) had a lower role. The results suggest that dispersal processes (e.g. permeability, dispersal limitation and movement activity of fishes) have critical importance on the assembly of stream fishes at very small spatial extents, which may strongly determine fish assemblages more upstream in the branches of the dendritic network. From an applied perspective revitalization projects often focus on enhancing instream habitat quality for the biota of streams (Lepori et al. 2005; Palmer et al. 2010). We suggest that environmental management of stream networks should take a more critical care on the filtering role of tributary confluences in species dispersal, for better understanding patterns and processes in the branches of dendritic stream networks.

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Table 1. The average habitat characteristics of the 200 m long sample sites in the Hajagos (H), Gerence (G) and Sokoroi (S) streams. Each mean value is based on the pooled mean data of seasonal (i.e. spring, summer, autumn) transect based surveys (see methods for details).

|  | Hajagos stream |  |  |  | Gerence stream |  |  |  |  |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  | H 1 | H 2 | H 3 | G 1 | G 2 | G 3 | Sokorói stream |  |  |
| Width (m) | $4.90(1.35)$ | $4.00(0.17)$ | $3.98(0.38)$ | $5.19(1.14)$ | $3.56(0.62)$ | $2.89(0.79)$ | $4.98(0.76)$ | $3.90(0.21)$ | $3.66(0.47)$ |
| Depth (cm) | $72.7(25.9)$ | $50.7(10.3)$ | $43.5(7.8)$ | $76.0(22.4)$ | $44.8(18.0)$ | $47.3(19.8)$ | $88.1(10.1)$ | $71.9(10.1)$ | $54.6(17.0)$ |
| Velocity (cm s ${ }^{-1}$ ) | $8.1(4.9)$ | $7.7(7.7)$ | $10.4(8.8)$ | $6.1(4.9$ | $13.4(12.6)$ | $9.7(6.2)$ | $6.6(4.4)$ | $9.8(4.6)$ | $7.5(6.1)$ |
| Silty sand (\%) | $49.1(23.5)$ | $38.7(23.4)$ | $28.1(7.0)$ | $46.3(18.4)$ | $47.6(28.2)$ | $81.9(13.4)$ | $81.3(23.5)$ | $77.1(28.0)$ | $56.3(16,4)$ |
| Sand (\%) | $0.4(1.4)$ | - | - | - | $1.1(1.7)$ | $3.2(9.6)$ | $2.1(4.5)$ | $4.6(5.4)$ | $8.3(12.3)$ |
| Fine gravel (\%) | $30.1(16.7)$ | $16.1(10.8)$ | $25.2(13.2)$ | $35.6(15.5)$ | $25.9(16.3)$ | $12.4(11.9)$ | $8.3(12.3)$ | $18.3(23.0)$ | $30.8(21.7)$ |
| Coarse gravel (\%) | $8.9(10.3)$ | $45.2(15.8)$ | $46.7(10.9)$ | $11.4(17.9)$ | $25.4(25.2)$ | $2.5(3.2)$ | $8.3(12.3)$ | - | $4.6(7.8)$ |
| Stone (\%) | $9.6(12.9)$ | - | - | $5.4(14.4)$ | - | - | - | - | - |
| Rock (\%) | $1.9(5.8)$ | - | - | $1.3(3.1)$ | - | - | - | - |  |
| Emergent plant (\%) | $52.3(40.1)$ | $61.9(40.8)$ | $67.7(35.8)$ | $23.7(21.6)$ | $27.5(26.5)$ | $53.1(22.2)$ | $15.6(9.2)$ | $26.6(10.9)$ | $69.2(24.0)$ |
| Submerged plant (\%) | $24.9(31.0)$ | $22.9(34.1)$ | $17.1(26.7)$ | $2.1(5.8)$ | $15.2(18.2)$ | $9.8(14.8)$ | $14.0(6.8)$ | $16.7(11.2)$ | $6.5(7.5)$ |
| Floating leaved plant (\%) | $4.4(6.0)$ | $8.5(15.9)$ | $4.3(8.2)$ | $20.6(35.6)$ | $0.1(0.3)$ | - | $17.1(27.8)$ | $3.1(6.1)$ | - |
| Filamentous algae (\%) | - | - | - | - | $0.2(0.4)$ | - | $4.6(6.9)$ | $13.3(20.6)$ | $2.2(3.6)$ |

Table 2. The fish species, species codes and the relative abundance of species collected in the mainstem Marcal River (M) and the Sokoroi (S), Gerence (G) and Hajagos (H) tributary streams.

| Species name | Species code | $\begin{gathered} \text { Relab } \\ \text { (M) (\%) } \end{gathered}$ | $\begin{gathered} \hline \text { Relab (S) } \\ (\%) \\ \hline \end{gathered}$ | Relab (G) (\%) | Relab (H) (\%) |
| :---: | :---: | :---: | :---: | :---: | :---: |
| Abramis brama (Linnaeus, 1758) | abrbra | 0.094 | 0.247 | - | 0.191 |
| Alburnus alburnus (Linnaeus, 1758) | albalb | 40.508 | 5.890 | 2.476 | 4.498 |
| Ameiurus melas (Rafinesque, 1820) | amemel | 0.399 | 0.038 | - | 0.383 |
| Aspius aspius (Linnaeus, 1758) | aspasp | 0.141 | - | - | - |
| Ballerus ballerus (Linnaeus, 1758) | balbal | 0.047 | 1.102 | 0.667 | - |
| Barbatula barbatula (Linnaeus, 1758) | ortbar | 0.023 | - | 0.334 | - |
| Barbus barbus (Linnaeus, 1758) | barbar | 0.141 | - | - | - |
| Blicca bjoerkna (Linnaeus, 1758) | blibjo | 1.739 | 2.451 | 7.322 | 12.344 |
| Carassius gibelio (Bloch, 1782) | cargib | 0.352 | 0.418 | - | 2.967 |
| Chondrostoma nasus (Linnaeus, 1758) | chonas | 0.094 | - | 0.018 | - |
| Cobitis elongatoides (Băcescu \& Maier, 1969) | cobelo | 0.235 | 1.045 | 0.474 | 0.287 |
| Cyprinus carpio (Linnaeus, 1758) | cypcar | 0.023 | 0.019 | - | - |
| Esox lucius (Linnaeus, 1758) | esoluc | 2.867 | 4.275 | 1.247 | 7.464 |
| Gobio obtusirostris (Valenciennes, 1842) | gobgob | - | - | 0.246 | - |
| Gymnocephalus cernuus (Linnaeus, 1758) | gymcer | - | - | 0.018 | - |
| Lepomis gibbosus (Linnaeus, 1758) | lepgib | 0.258 | 0.038 | 1.826 | 0.096 |
| Leuciscus idus (Linnaeus, 1758) | leuidu | 1.386 | 3.040 | 0.158 | 1.627 |
| Leuciscus leuciscus (Linnaeus, 1758) | leuleu | 0.752 | 0.817 | 1.212 | 0.670 |
| Misgurnus fossilis (Linnaeus, 1758) | misfos | - | 0.247 | 0.228 | 2.584 |
| Neogobius fluviatilis (Pallas, 1814) | neoflu | 0.728 | 0.171 | 1.159 | 0.957 |
| Neogobius melanostomus (Pallas, 1814) | neomel | 1.269 | 0.095 | 1.054 | 1.053 |
| Perca fluviatilis (Linnaeus, 1758) | perflu | 1.621 | 3.648 | 1.018 | 0.191 |
| Phoxinus phoxinus (Linnaeus, 1758) | phopho | - | - | 0.035 | - |
| Proterorhinus semilunaris (Pallas, 1814) | prosem | 1.692 | 2.812 | 0.544 | 3.923 |
| Pseudorasbora parva (Temminck \& Schlegel, 1842) | psepar | - | 0.399 | 0.105 | 2.105 |
| Rhodeus sericeus (Pallas, 1776) | rhoser | 11.302 | 46.114 | 58.472 | 25.359 |
| Rutilus rutilus (Linnaeus, 1758) | rutrut | 31.790 | 23.580 | 18.262 | 28.134 |
| Scardinius erythrophthalmus (Linnaeus, 1758) | scaery | 0.705 | 0.836 | 0.070 | 0.957 |
| Silurus glanis (Linnaeus, 1758) | silgla | 0.211 | - | - | - |
| Squalius cephalus (Linnaeus, 1758) | squcep | 1.081 | 2.641 | 3.020 | 4.115 |
| Tinca tinca (Linnaeus, 1758) | tintin | 0.023 | 0.038 | - | 0.096 |
| Vimba vimba (Linnaeus, 1758) | vimvim | 0.470 | 0.038 | 0.035 | - |
| Zingel zingel (Linnaeus, 1758) | zinzin | 0.047 | - | - | - |

Table 3. Summary results of the linear models (nested design) for species richness, and dissimilarity of species composition and relative abundance of fishes from the mainstem Marcal River.

| Source of variation | d.f. | MS | $F$ | $P$ | \% variance |
| :--- | ---: | ---: | ---: | ---: | ---: |
| Species richness |  |  |  |  |  |
| Season | 2 | 42.19 | 10.62 | $<0.001$ | 7.5 |
| Stream (within season) | 6 | 68.30 | 17.19 | $<0.001$ | 36.4 |
| Site (within season and stream) | 9 | 24.19 | 6.09 | $<0.001$ | 19.3 |
| Habitat PC1 (within season and stream) | 9 | 6.85 | 1.72 | 0.102 | 5.5 |
| Habitat PC2 (within season and stream) | 9 | 7.13 | 1.79 | 0.087 | 5.7 |
| Habitat PC3 (within season and stream) | 9 | 4.33 | 1.09 | 0.384 | 3.5 |
| Residuals | 63 | 3.97 |  |  | 22.2 |
| Dissimilarity in species composition |  |  |  |  |  |
| Season | 2 | 0.03 | 5.00 | 0.010 | 3.2 |
| Stream (within season) | 6 | 0.12 | 20.26 | $<0.001$ | 39.0 |
| Site (within season and stream) | 9 | 0.05 | 8.19 | $<0.001$ | 23.6 |
| Habitat PC1 (within season and stream) | 9 | 0.01 | 1.95 | 0.061 | 5.6 |
| Habitat PC2 (within season and stream) | 9 | 0.01 | 2.31 | 0.026 | 6.7 |
| Habitat PC3 (within season and stream) | 9 | 0.00 | 0.60 | 0.792 | 1.7 |
| Residuals | 63 | 0.01 |  |  | 20.2 |
| Dissimilarity in relative abundance |  |  |  |  |  |
| Season | 2 | 1.36 | 95.77 | $<0.001$ | 54.3 |
| Stream (within season) | 6 | 0.05 | 3.76 | 0.003 | 6.4 |
| Site (within season and stream) | 9 | 0.07 | 4.88 | $<0.001$ | 12.4 |
| Habitat PC1 (within season and stream) | 9 | 0.02 | 1.56 | 0.147 | 4.0 |
| Habitat PC2 (within season and stream) | 9 | 0.01 | 0.65 | 0.750 | 1.7 |
| Habitat PC3 (within season and stream) | 9 | 0.02 | 1.30 | 0.253 | 3.3 |
| Residuals | 63 | 0.01 |  |  | 17.9 |

Table 4. Spearman correlation tests between species richness and rank distance from the junction, dissimilarity in species composition (Jaccard index) and rank distance from the junction, and relative abundance (Bray-Curtis index) and rank distance from the junction for each stream and season ( $\mathrm{n}=12$ in each test).

| Species richness |  |  |  |  |  |  |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  | Spring |  | Summer |  | Autumn |  |
|  | $\mathrm{R}^{2}$ | $P$ | $\mathrm{R}^{2}$ | $P$ | $\mathrm{R}^{2}$ | $P$ |
| Sokoroi | -0.79 | 0.002 | -0.52 | 0.080 | -0.61 | 0.037 |
| Gerence | -0.69 | 0.013 | -0.79 | 0.002 | -0.76 | 0.004 |
| Hajagos | -0.12 | 0.704 | -0.54 | 0.067 | -0.89 | <0.001 |
| Species composition |  |  |  |  |  |  |
|  | Spring |  | Summer |  | Autumn |  |
|  | $\mathrm{R}^{2}$ | $P$ | $\mathrm{R}^{2}$ | P | $\mathrm{R}^{2}$ | $P$ |
| Sokoroi | -0.45 | 0.138 | 0.09 | 0.770 | 0.34 | 0.276 |
| Gerence | 0.30 | 0.338 | 0.80 | 0.002 | 0.83 | $<0.001$ |
| Hajagos | 0.29 | 0.367 | 0.69 | 0.013 | 0.80 | 0.002 |
|  |  |  |  |  |  |  |
| Relative abundance |  |  |  |  |  |  |
|  | Spring |  | Summer |  | Autumn |  |
|  | $\mathrm{R}^{2}$ | $P$ | $\mathrm{R}^{2}$ | $P$ | $\mathrm{R}^{2}$ | $P$ |
| Sokoroi | -0.60 | 0.039 | -0.37 | 0.236 | 0.85 | $<0.001$ |
| Gerence | -0.71 | 0.009 | -0.71 | 0.009 | 0.04 | 0.914 |
| Hajagos | -0.04 | 0.914 | 0.25 | 0.430 | 0.03 | 0.931 |

Table 5. Summary results of the PERMANOVA analyses for species composition and relative abundance data.

| Source of variation | d.f. | MS | $F$ | $P$ | \% variance |
| :--- | ---: | :--- | ---: | ---: | ---: |
| Dissimilarity in species composition |  |  |  |  |  |
| Season | 2 | 1.90 | 13.70 | $<0.001$ | 14.6 |
| Stream (within season) | 6 | 0.97 | 6.99 | $<0.001$ | 22.4 |
| Site (within season and stream) | 9 | 0.39 | 2.85 | $<0.001$ | 13.7 |
| Habitat PC1 (within season and stream) | 9 | 0.16 | 1.16 | 0.154 | 5.6 |
| Habitat PC2 (within season and stream) | 9 | 0.19 | 1.35 | 0.018 | 6.5 |
| Habitat PC3 (within season and stream) | 9 | 0.11 | 0.78 | 0.938 | 3.7 |
| Residuals | 63 | 0.14 |  |  | 33.6 |
| Dissimilarity in relative abundance |  |  |  |  |  |
| Season | 2 | 1.47 | 13.75 | $<0.001$ | 12.6 |
| Stream (within season) | 6 | 0.69 | 6.44 | $<0.001$ | 17.8 |
| Site (within season and stream) | 9 | 0.54 | 5.01 | $<0.001$ | 20.7 |
| Habitat PC1 (within season and stream) | 9 | 0.18 | 1.71 | 0.007 | 7.1 |
| Habitat PC2 (within season and stream) | 9 | 0.17 | 1.55 | 0.022 | 6.4 |
| Habitat PC3 (within season and stream) | 9 | 0.17 | 1.55 | 0.021 | 6.4 |
| Residuals | 63 | 0.11 |  |  | 29.0 |

## Captions to figures

Fig. 1 Scheme of the study design using three tributary streams (Sokoroi, Gerence, Hajagos) and three sites in each stream in the Marcal River system, North Western, Hungary. The sites were situated 0,500 and 1000 m from the tributary mouth. Numbers indicate individual habitat units ( 50 m long each)

Fig. 2 Mean species richness ( $\pm$ S.D.) of the sampling units ( 50 m long each) at three different sites in the Sokoroi, Gerence and Hajagos streams of the Marcal River in spring, summer and autumn

Fig. 3 Mean dissimilarity ( $\pm$ S.D.) of the sampling units ( 50 m long each) from the mainstem Marcal River for (a) species composition (Jaccard index) and (b) relative abundance (BrayCurtis index) data of fishes at three different sites in the Sokoroi, Gerence and Hajagos streams in spring, summer and autumn

Fig. 4 Changes in (a) species richness, and dissimilarity in (b) species composition (Jaccard index) and (c) relative abundance (Bray-Curtis index) data from the mainstem Marcal River at the level of the sampling units ( 50 m long each) in the Sokoroi, Gerence and Hajagos streams in spring, summer and autumn

Fig. 5 Principal Coordinate Analyses (PCoA) summarizing the variation across sites in three tributaries of the Marcal River for (a) species composition and (c) relative abundance data, and their associated species loadings ( $b$ and d). For species code abbreviations see Table 2

## Appendix

Table 6. Pearson correlation matrix of the abiotic variables and the first three principal component axes ( $\mathrm{PC} 1, \mathrm{PC} 2, \mathrm{PC} 3$ ). Lower and upper diagonals show correlations and their significance values, respectively.

|  | PC1 | PC2 | PC3 | Width (m) | Depth (cm) | Velocity ( $\mathrm{cm} \mathrm{s}^{-1}$ ) | Silty sand (\%) | Sand (\%) | Fine gravel (\%) | Coarse gravel (\%) | Stone and rock (\%) | Emergent plant (\%) | Submerged plant (\%) | $\begin{gathered} \hline \text { Floating leaved } \\ \text { plant (\%) } \\ \hline \end{gathered}$ | Filamentous algae (\%) |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| PC1 | - | 1.000 | 1.000 | $<0.001$ | $<0.001$ | $<0.001$ | $<0.001$ | 0.001 | 0.151 | $<0.001$ | 0.054 | <0.001 | $<0.001$ | 0.137 | 0.424 |
| PC2 | $<0.001$ | - | 1.000 | $<0.001$ | 0.002 | 0.091 | <0.001 | 0.027 | <0.001 | 0.001 | $<0.001$ | 0.447 | 0.483 | $<0.001$ | 0.006 |
| PC3 | $<0.001$ | <0.001 | - | 0.331 | $<0.001$ | $<0.001$ | $<0.001$ | $<0.001$ | 0.194 | $<0.001$ | 0.827 | 0.59 | $<0.001$ | 0.007 | $<0.001$ |
| Width (m) | -0.603 | 0.55 | 0.095 | - | $<0.001$ | 0.281 | 0.342 | 0.003 | 0.011 | 0.279 | 0.002 | $<0.001$ | 0.038 | 0.541 | 0.627 |
| Depth (cm) | -0.794 | 0.297 | 0.334 | 0.732 | - | 0.085 | 0.016 | 0.05 | 0.789 | $<0.001$ | 0.002 | $<0.001$ | 0.001 | 0.661 | 0.162 |
| Velocity ( $\mathrm{cm} \mathrm{s-1)}$ | -0.614 | -0.164 | -0.489 | 0.105 | 0.167 | - | 0.238 | 0.014 | 0.646 | 0.807 | 0.586 | $<0.001$ | $<0.001$ | 0.003 | 0.054 |
| Silty sand (\%) | -0.4 | -0.76 | 0.404 | -0.092 | 0.231 | 0.115 | - | 0.709 | $<0.001$ | $<0.001$ | 0.013 | 0.02 | 0.509 | 0.035 | 0.177 |
| Sand (\%) | 0.306 | -0.213 | 0.394 | -0.282 | -0.189 | -0.236 | -0.036 | - | 0.284 | 0.168 | 0.558 | 0.317 | 0.264 | 0.33 | $<0.001$ |
| Fine gravel (\%) | 0.139 | 0.731 | 0.126 | 0.243 | 0.026 | -0.045 | -0.587 | -0.104 | - | 0.579 | 0.913 | 0.548 | 0.059 | 0.005 | 0.007 |
| Coarse gravel (\%) | 0.375 | 0.305 | -0.73 | -0.105 | -0.367 | -0.024 | -0.703 | -0.134 | -0.054 | - | 0.725 | 0.03 | 0.389 | 0.362 | 0.208 |
| Stone and rock (\%) | -0.186 | 0.355 | -0.021 | 0.293 | 0.297 | -0.053 | -0.237 | -0.057 | 0.011 | -0.034 | - | 0.966 | 0.617 | 0.514 | 0.44 |
| Emergent plant (\%) | 0.792 | -0.074 | -0.052 | -0.347 | -0.567 | -0.501 | -0.223 | 0.097 | 0.058 | 0.209 | 0.004 | - | $<0.001$ | 0.145 | 0.209 |
| Submerged plant (\%) | -0.67 | -0.068 | -0.417 | 0.2 | 0.316 | 0.609 | 0.064 | -0.108 | -0.182 | 0.084 | 0.049 | -0.568 | - | 0.038 | 0.72 |
| Floating leaved plant (\%) | 0.144 | 0.484 | 0.258 | 0.06 | 0.043 | -0.288 | -0.203 | -0.095 | 0.266 | 0.089 | -0.063 | -0.141 | -0.2 | - | 0.072 |
| Filamentous algae (\%) | 0.078 | 0.263 | 0.535 | -0.047 | 0.135 | $-0.186$ | -0.131 | 0.346 | 0.26 | -0.122 | -0.075 | -0.122 | -0.035 | 0.174 | - |

Distance from the mouth

|  | 1000 m | 500 m | 0 m |
| :---: | :---: | :---: | :---: |
|  | $\stackrel{\sim}{\text { ® }}$ | $\stackrel{\text { n }}{\text { ¢ }}$ | $\stackrel{0}{\square}$ |
|  | $\omega$ | N | $\rightarrow$ |
| so6ere\% | बजलग |  |  |


| site $1{ }^{\frac{1}{\frac{2}{3}}}$ | 1 |
| :---: | :---: |
| site $2 \frac{\frac{1}{2}}{\frac{2}{3}}$ | $\frac{1}{2}$ |
| site $3 \frac{2}{\frac{2}{3}}$ | 圱 |










