# Mesohabitat mosaic in lowland braided rivers: Short-term variability of macroinvertebrate metacommunities

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#### **ABSTRACT**

Braided rivers are among the most variable and dynamic riverine systems. Changes in these environments are sudden and frequent, driven by the high hydrological variability. They host high levels of local heterogeneity, with many different habitats in close proximity establishing a mosaic of patches. This provides the conditions for high levels of biodiversity, with strong community variability in particular among the different habitats at the stream-reach level. Nevertheless, these systems are still poorly studied and their complexity is often not taken into account in biomonitoring protocols. We applied mixed effects modelling, spatial ordination techniques and beta-diversity partitioning (into nestedness and turnover components) with the aim of improving the knowledge of braided rivers, investigating: i) the organization of macroinvertebrate communities among the different habitats of a river reach, and ii) the temporal variability of this organization (both among seasons and during summer). We predicted a differentiation of macroinvertebrate communities between distinct habitats within rivers, with this differentiation increasing during the low-flow period. We carried out our study in four braided rivers and streams of the Po River basin (Northern Italy) sampling three different kinds of mesohabitats (main channel, secondary channel and pool) in eight stations during seven campaigns from June 2015 to April 2016. We found a high variability of taxa richness, abundance and community structure among mesohabitats, with marginal ones accounting for the greater part of macroinvertebrate diversity. Secondary channels resulted as being the habitat hosting greater taxa diversity, with 10 exclusive taxa. Surprisingly the mesohabitat communities differed greatly during the seasonal phase, whereas their dissimilarity decreased during summer. This could be explained considering the summer flow reduction as a homogenizing force, leading to a general loss of the most sensitive taxa. However, the summer taxa turnover value resulted higher than nestedness, suggesting a strong environmental control on community organization, with taxa well adapted to the different conditions of mesohabitats and able to manage the effects of flow reduction. Our work represents a remarkable issue for biomonitoring protocols, highlighting the importance of taking into account the whole complexity of braided rivers for a more realistic evaluation of macroinvertebrate communities.

Key word: Braided rivers; macroinvertebrate metacommunity; mesohabitat; short-term variability; macroinvertebrate distribution.

## INTRODUCTION

Braided rivers (BRs) are defined as "systems composed by multiple channels, with bars and islands, often with poorly defined banks of non-cohesive sedimentary materials" (Tockner et al., 2006). These systems are spread worldwide and can be found in delta areas, where rivers enter lakes and oceans, or in floodplains in presence of low slope and sandy or gravel-filled substrates (Dodds, 2002). The main feature that shapes BRs is the extreme flow variability. The alternation of sudden and frequent flow changes, spanning flash floods and dry periods generates a mosaic of patches (embracing different degrees of lotic and lentic conditions), which undergoes rapid evolution (Gray and Harding, 2009). Events like the displacement of channels or disconnection of habitats can happen within short periods, spanning from a few weeks to a few hours. Van der Nat et al. (2003) estimated the turnover time of the different habitats in a BR system (Tagliamento, NE

Italy) reporting a high level of variation, with a total replacement of all the aquatic habitats of 82% during the period of study (2.5 years). Nevertheless, they reported that the relative proportion of the various habitats remained quite consistent. Based on these results, BRs can be conformed to the "shifting mosaic steady model" that identify systems where the habitat turnover is high but the proportions of habitats are constant (Tockner *et al.*, 2006; Gray and Harding, 2011).

Based on these attributes, BRs can be considered as very suitable systems for studying metacommunity dynamics. According to the metacommunity theory (Leibold *et al.*, 2004), environmental heterogeneity and *taxa* features (*e.g.*, dispersal ability and competition) determine the structure and evolution of metacommunities at different spatial scales (Siqueira *et al.*, 2012; McLaughlin *et al.*, 2013). In general, habitat heterogeneity has a positive effect on species richness (Poff and Ward, 1990; Garcia *et al.*, 2012; Astorga *et al.*, 2014), enhancing the niche availability and allowing the co-occurrence of *taxa* with dif-





ferent requirements. The high dynamism of BRs generates a great heterogeneity, especially at the scale of river reach, with a wide range of different habitats, spanning from lotic to lentic conditions and with a time-variable level of connection. The degree of influence of dispersal dynamics and environmental forcing is strictly related to the connectivity of habitats, besides the dispersal ability of taxa (Padial et al., 2014). In riverine systems, the level of connectivity can change widely in time and among them BRs are one of the most dynamic and complex (Ward et al., 2002). All these conditions are the basis for the presence of biodiversity hot-spots, with high levels of diversity variation in particular among the different habitats at the stream-reach level. In fact, several authors pointed out high levels of lateral variation in taxa diversity and community structure for braided systems (e.g., Arscott et al., 2005; Gray and Harding, 2007, respectively in north-eastern Italy and New Zealand). Similar outputs were also recorded for primary producers in lowland rivers largely fed by groundwater (Bolpagni and Laini, 2016), suggesting the existence of complex metabolic gradients across habitats in hydro-systems.

The variation among habitats can be considered as a beta-diversity variation and therefore it can be ascribed to two different phenomena: nestedness and spatial turnover. Nestedness occurs when there is a non-random *taxa* loss, with the result that the poorer communities are a subset of the richer ones, while turnover is the result of *taxa* replacement (Baselga, 2010). Datry *et al.* (2016) highlighted that turnover is more related to environmental filtering, while nestedness is given by dispersal limitation. These two processes can assume differential importance in shaping local communities, in particular during low-flow periods, when connectivity among habitats is more variable.

Although these systems are widespread and considered as diversity hot-spots, for years they have been poorly studied (Gray and Harding, 2007), with a lack of knowledge, especially in how the different habitats in the river segment contribute to the total diversity and how these patterns change in time. This topic is particularly relevant considering that BRs are often located in areas heavily impacted by human activities, with all the possible consequences, like considerable water withdrawals, canalization and reduction or loss of lateral areas (Tockner et al., 2006; Gray and Harding 2011; Karaus et al., 2013). These phenomena lead to a trivialization of BRs, with the consequent reduction of habitat variability. Therefore, a good understanding of habitat heterogeneity contribution to the local diversity becomes a key point for biodiversity conservation.

The aims of this study are therefore i) to evaluate the seasonal structure and variation of benthic macroinvertebrate communities within the highly patchy environments of BRs and ii) to evaluate the short-term variability of these communities during the low-flow period. For this work, we focused on the mesohabitat sampling unit, demarcated according to the hydrodynamic characteristics in main channel, secondary channel and pool. Tickner *et al.* (2000) defined mesohabitats as "medium-scale habitats which arise through the interactions of hydrological and geomorphological forces". We hypothesize that: i) in general there is a differentiation of macroinvertebrate communities between different mesohabitats within rivers, and between considered seasons ii) during low-flow periods, with the increasing disconnection of mesohabitats there is an increase in community dissimilarity, with higher turnover in the less disconnected mesohabitats and higher nestedness in the more disconnected ones.

## **METHODS**

# Study area

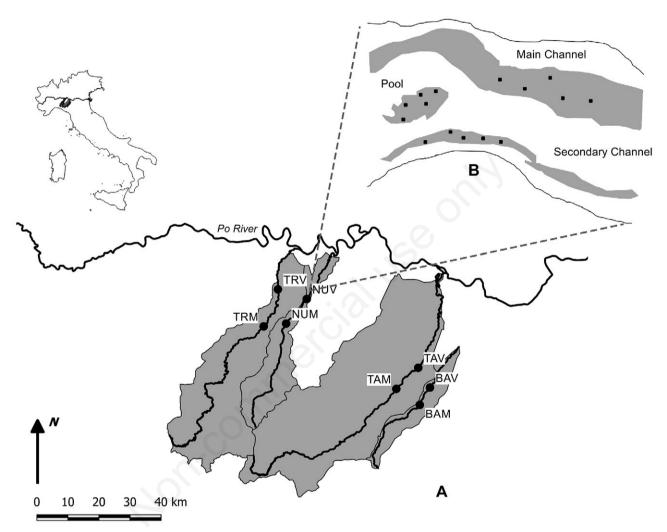
The study was performed in four braided watercourses (Trebbia River, Nure Stream, Taro River and Baganza Stream) of the Po River basin (Northern Italy, Fig. 1A). A description of the studied systems is reported in Tab.1. They are fed only by wet depositions and they present two high discharge periods (in autumn and spring) and a main low water period in summer (with a secondary additional one in winter). They are included in the Cfa (humid subtropical climate) and Csa (hot-summer Mediterranean climate) climatic regions. For each watercourse we selected two sampling stations in order to take into account the within river variability. Within each station, three model mesohabitats were further selected: main channel, secondary channel, and pool (Fig. 1B). For the first part of the study (seasonal phase, T6-T7) sampling was carried out in November 2015 and in April 2016 in the whole set of systems. For the second part (summer phase, T1-T5) the set of investigated systems was reduced to two (Trebbia and Taro rivers) and the sampling was carried out in five occasions in the period of low flow, from June to September 2015. The downsizing of sampled area was operated because i) Nure and Baganza streams completely dried up during the summer season and ii) to contain the sampling and processing effort.

#### Physical and chemical variables

In order to check the difference between mesohabitats, for each sampling environmental data were collected with five random replicates (Fig. 1B). Flow velocity, water depth, temperature, conductivity and dissolved oxygen were recorded *in situ* by means of a current meter (FP101-FP102 Global Flow Probe) and a multi-parametric probe (HI 9828; Hanna Instruments). Water samples were collected for the determination of ammonium (NH<sub>4</sub><sup>+</sup>), nitrite (NO<sub>2</sub><sup>-</sup>), nitrate (NO<sub>3</sub><sup>-</sup>), soluble reactive phosphorous (SRP), dissolved silica (SiO<sub>2</sub>) and total dissolved inorganic

carbon (TCO<sub>2</sub>). Detection limits were 0.01 mg  $L^{-1}$  for NH<sub>4</sub><sup>+</sup>, NO<sub>3</sub><sup>-</sup>, SRP and SiO<sub>2</sub>, 0.005 mg  $L^{-1}$  for NO<sub>2</sub><sup>-</sup>, and 0.02 mM for TCO<sub>2</sub>. Precision ranged between  $\pm 3\%$  and

±5%. Chemical analyses were performed by means of spectrophotometric techniques, according to Valderrama (1977), Golterman *et al.* (1978), and APHA (2012).



**Fig. 1.** Map of the sampled area (A), with the studied basins coloured in grey and the sampling stations marked with black dots, and the sampling design (B) repeated in each of the sampling stations. Black squares represent the random sampling plots.

**Tab. 1.** Descriptive data of the investigated rivers/streams and stations.

River/stream	Qm (m <sup>3</sup> s <sup>-1</sup> )	L (km)	A (km²)	Station	Latitude	Longitude	Altitude (m asl)	W (m)	D (km)
Trebbia	22.0	120	1083	TRM TRV	44°51'11.05"N 44°58'18.51"N	9°32'11.75"E 9°35'32.54"E	166 97	298 440	13.9
Nure	15.0	77	458	NUM NUV	44°51'52.84"N 44°55'55.97"N	9°37'48.73"E 9°42'40.96"E	212 124	240 244	9.9
Taro	40.5	138	2051	TAM TAV	44°40'27.92''N 44°44'26.97''N	10° 4'23.03"E 10°10'6.14"E	151 95	530 445	10.6
Baganza	5.2	59	228	BAM BAV	44°37'54.67''N 44°40'54.20''N	10°10'10.33"E 10°12'36.90"E	308 213	106 183	6.4

Qm, mean annual flow; L, total length; A, basin area; W, width of the riverbed; D, linear distance between stations.

### **Macroinvertebrates**

In each mesohabitat, a  $\sim$ 50 m long stretch was sampled, choosing five random sampling points (Fig. 1B). Samples were collected using a surber net with frame area of 0.1 m<sup>2</sup> and mesh size of 500  $\mu$ m. The five replicates were cumulated for each mesohabitat. Samples were filtered and preserved in 70% ethanol for laboratory sorting, where the organisms were counted and identified to family or genus level, according to Tachet *et al.* (2010).

## Data analysis

The difference between mesohabitats, in terms of physical variables, was assessed by means of mixed effects modelling, considering mesohabitat and time (sampling date) as fixed effects and station and site (namely the specific sampling location) as hierarchically organized random effects. A similar approach was followed also for testing the influence of mesohabitat and time on richness and abundance. The significance was checked by means of a likelihood-ratio test. The use of these models allows us to work with correlated and non-normally distributed data (McCulloch and Neuhaus, 2005), typical of nested and hierarchical designs. The effect of covariate was tested both for seasonal and summer data. Then the distribution of taxa between mesohabitats at station level was checked, by estimating the mesohabitat contribution to the total number of taxa. We did this by computing the percentage ratio for each station between the richness of each mesohabitat and the total richness of the station.

The organization of community structure in mesohabitats was explored with a non-Metric Multidimensional Scaling (nMDS), a spatial ordination technique that represents the set of objects along a predetermined number of axes maintaining the ordering relationships among them (Borcard *et al.*, 2011). As dissimilarity measure Bray-Curtis distance was used and the goodness of ordination was assessed with the stress measure.

To assess the nature of diversity variation between mesohabitats during the summer phase we performed a partition of beta-diversity, following Baselga (2010). This method produces three metrics: the total beta-diversity (the Sørensen Dissimilarity index), for all the possible pairwise comparisons, and its two additive components: nestedness, expressing the taxa loss between mesohabitats, and the turnover, expressing the taxa substitution. The values of nestedness and turnover were normalized by dividing them by the Sørensen dissimilarity value. We checked the effect of time for the Sørensen Dissimilarity index by means of linear mixed effect models and then we adjusted the p-values using the Bonferroni correction for multiple comparisons. We also applied mixed effects modelling to check the difference between beta-diversity components and their variation in time.

All analyses and graphs were performed with the statistical software R (R Core Team, 2016), with base version and with ggplot2 (Wickham, 2009), lme4 (Bates *et al.*, 2015), vegan (Oksanen *et al.*, 2016) and betapart (Baselga *et al.*, 2013) packages.

#### RESULTS

# Physical and chemical variables

Mean values of measured physical and chemical variables are reported in Tab. 2, according to season and mesohabitat. The distinction between mesohabitats was tested for physical variables considering the whole dataset (seasonal and summer data) and we found that they differ greatly for flow velocity (P<0.001) and water depth (P<0.001), while the others variables (temperature, conductivity and percentage of dissolved oxygen) varied significantly only in time (P<0.001) but not between mesohabitats.

#### **Macroinvertebrates**

A total of 74122 organisms, belonging to 94 taxa (75 families) was found globally. The sample with the highest taxa richness (34 taxa) was collected at the beginning of the summer period in the upstream pool of Trebbia River, while the one with the lowest (seven taxa) in the downstream pool of Nure Stream, during the November sampling campaign. The mean values of taxa richness and abundance were 18±5 and 837±743 for main channels, 22±5 and 796±546 for secondary channels and 17±6 and 426±552 for pools. The list of most abundant (A) and frequent (B) taxa is reported in Fig. 2. Chironomidae was both the most abundant and frequent taxon (abundance = 29.3%, frequency = 99.1%). Detection probabilities for the other most common taxa were unrelated to their abundance. Some taxa were found to be exclusive of one kind of mesohabitat: we found five exclusive taxa in the main channels (Heptagenia, Notonecta, Gordiidae, Besdolus and Brachyptera), six in the pools (Pseudocentroptilum, Pisidium, Dixidae, Hydrometra, Haplotaxidae and Protonemura) and 10 in the secondary channels (Hydridae, Blephariceridae, Dolichopodidae, Ephydridae, Rhagionidae, Valvata, Gerris, Helobdella, Nemoura and Lepidostomatidae).

The significance of mesohabitat and time for *taxa* richness and organism abundance was tested by means of several mixed effects models for seasonal and summer phases. Both *taxa* richness and abundance resulted related with mesohabitats, especially for seasonal data (P values 0.002 and 0.003 respectively), while for summer data these relations resulted weaker (P values 0.078 and 0.060). Time resulted significant only for the seasonal *taxa* richness (P= 0.026), with a variation between No-

vember and April, while no significant variation was found during summer nor for abundance. The effect of mesohabitat resulted clear also considering community composition patterns (Fig. 3 A,B): points corresponding to the three kinds of mesohabitats group into different areas of the nMDS plot, both for seasonal and summer

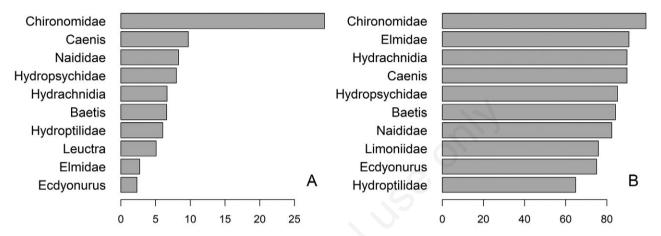


Fig. 2. Barplots of the first 10 taxa in terms of abundance percentage (A) and frequency percentage (B). Values are referred to the whole dataset.

**Tab. 2.** Physical and chemical variables for mesohabitats in each season.  $NH_4^+$ ,  $NO_2^-$  and SRP values are not shown because always lower than detection values. Autumn and Spring values are for the whole set of systems, whereas Summer values are referred only to Trebbia and Taro rivers.

		Autumn		Spring		Summer	mer	
		<b>Mean values</b>	SD	Mean values	SD	Mean values	SD	
Flow velocity (m s <sup>-1</sup> )	Main	0.47	0.24	0.46	0.07	0.42	0.16	
	Sec	0.31	0.36	0.14	0.07	0.22	0.19	
	Pool	0.08	0.15	0.00	0.01	0.07	0.13	
Water depth (cm)	Main	22.6	7.7	28.3	6.4	21.3	6.9	
	Sec	13.5	10.0	10.7	8.7	14.3	8.0	
	Pool	24.3	9.3	13.1	7.6	19.1	6.8	
Temperature (°C)	Main	13.0	1.2	14.1	1.3	23.5	2.0	
	Sec	13.8	1.5	14.9	2.2	23.7	3.8	
	Pool	13.3	1.3	15.4	1.6	24.6	1.9	
Conductivity (µS cm <sup>-1</sup> )	Main	250	29	253	24	328	45	
	Sec	287	81	272	35	383	143	
	Pool	268	38	274	27	375	74	
Dissolved oxygen (%)	Main	104.1	5.8	103.8	12.2	107.5	8.1	
	Sec	100.5	12.1	102.0	11.2	96.3	18.8	
	Pool	94.0	11.3	88.3	12.6	105.1	18.4	
NO <sub>3</sub> -	Main	0.29	0.08	0.16	0.08	0.23	0.14	
	Sec	0.36	0.22	0.17	0.09	0.49	0.64	
	Pool	0.31	0.07	0.21	0.06	0.44	0.67	
SiO <sub>2</sub> (mg L <sup>-1</sup> )	Main	1.36	0.27	1.24	0.36	2.62	0.62	
	Sec	1.37	0.32	1.32	0.38	3.00	0.80	
	Pool	1.38	0.34	1.39	0.23	3.10	0.38	
TCO <sub>2</sub> (mM)	Main	2.14	0.21	2.32	0.23	1.88	0.40	
	Sec	2.34	0.58	2.20	0.29	1.92	0.50	
	Pool	2.21	0.32	2.51	0.43	1.96	0.53	

Main, main channels; sec, secondary channels; pool, pools.

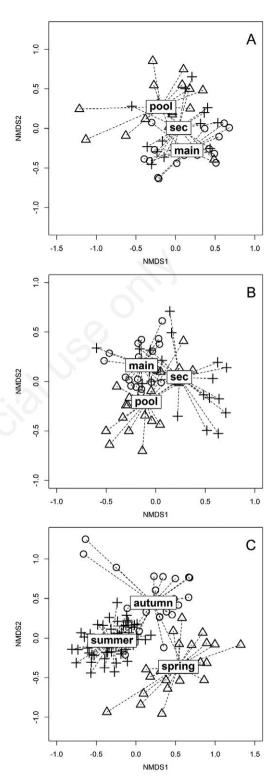
communities. Moreover, comparing the two graphs, it can be seen that the segregation between mesohabitats resulted slightly greater during the seasonal phase (November and April) than for summer. We also found a variability of communities in time (Fig. 3C), with a clear segregation of autumn, spring and summer data in three different clusters.

The mesohabitats contribution to the *taxa* richness at station level resulted significantly different, either seasonally (P=0.002) or during the summer (P=0.045). The greater contribution was the one given by marginal mesohabitats and by the secondary channels in particular, while the importance of main channels resulted limited (Fig. 4).

Variation during the summer phase of the Sørensen Dissimilarity index and of the beta-diversity partition for the pairwise comparisons between mesohabitats are reported in Fig. 5 A,B. The dissimilarity values (Sørensen Dissimilarity index) showed similar trends in all comparisons, with a decrease in June and July (T1:T5 in Fig. 5A) and a new increase at the beginning of autumn (T6, T7). Nevertheless, this trend resulted significant only considering the comparison between main channels and pools (P=0.027). The two components of beta-diversity resulted significantly different for all the comparison (P<0.001), with higher values for *taxa* turnover. No significant temporal trends resulted from the analysis.

#### DISCUSSION AND CONCLUSIONS

This study highlights a strong variability of macroinvertebrate assemblages in BRs, with different mesohabitats hosting different communities. These findings are consistent with Gray and Harding (2009), Zilli and Marchese (2011), Karaus et al. (2013) and Starr et al. (2014), that reported significant levels of variation of taxa richness and abundance among mesohabitats inside river reaches. Arscott et al. (2005) found greater diversity in macroinvertebrate communities of backwaters areas of Tagliamento River, while Gray and Harding (2009) pointed out spring creeks, spring sources and ponds and Zilli and Marchese (2011) isolated lakes as mesohabitats hosting greater diversity in New Zealand rivers and in Panamá River floodplain, respectively. By contrast, in our systems secondary channels resulted as being the mesohabitat hosting greater taxa diversity, both for seasonal and summer data, while in the other works these mesohabitats resulted in those with low diversity compared to the other ones. This higher diversity could be explained considering that secondary channels were characterized by intermediate levels of hydrological disturb (mean discharge, depth and water velocity), were often located near the margin of riverbeds and presented a higher heterogeneity of microhabitats (cobbles, gravel, clay, algal mats and roots). These features have been shown to be critical



**Fig. 3.** nMDS ordination outputs for seasonal (A), summer (B) and all (C), data. For seasonal and summer data (A and B) the mesohabitat segregation is shown, while for the all data graph (C) the temporal (between seasons) segregation of communities is shown. Seasonal nMDS stress, 0.176; Summer nMDS stress, 0.157; all data nMDS stress, 0.175; main, main channels; sec, secondary channels; pool, pools.

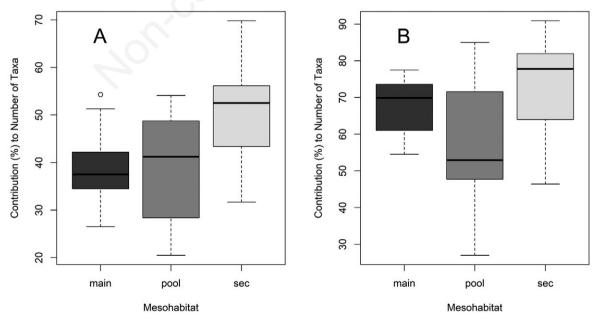
in enhancing the within site richness (Downes *et al.*, 2000). Additionally, they show greater shadowing levels that significantly modulate the colonization patterns of primary producers (algae, vascular macrophytes; data not shown), influencing the availability of resources for macroinvertebrates.

Several factors have been proposed as main drivers for BRs macroinvertebrate community differentiation: conductivity and percentage of sand (Zilli and Marchese, 2011), flow velocity (Arscott *et al.*, 2005), nature of substrate (Beisel *et al.*, 1998). In our work, we considered the mesohabitat category as a proxy of physical environment differentiation: in particular, we found that main channels, secondary channels and pools mainly differed in flow velocity and water depth. Given the high significance of mesohabitats for macroinvertebrate, we hypothesized a strong physical control of communities, with a selection of *taxa* based on their habitat needs, also suggested by the presence of unique *taxa* for different mesohabitats.

A temporal trend that arose from our results is the evident difference in the importance of mesohabitats between seasonal and summer phases, supported by both mixed effects modelling and nMDS ordination. These findings are in contrast with Starr *et al.* (2014) and Arscott *et al.* (2003), who reported an increase of compositional heterogeneity coming from the increasing isolation of sampling sites, and from flood homogenization respectively. García-Roger *et al.* (2011) instead found similar results, with a mesohabitat (riffles and pools in their

study) differentiation smaller during the dry season for a decrease of mesohabitat heterogeneity. The greater differentiation of mesohabitat communities observed in the present study during seasonal samplings (November and April) could be the result of a major connectivity that allows organisms to actively choose the best living place, according to their necessities. This generates a high environmental control on the community from the moment that there are no dispersal limitations and the choice of the most suitable environmental features drives the community. On the other hand, during summer the disconnection increases, hampering the dispersion of organisms. Unlike our initial hypothesis, in this phase we observe a temporal trend of community dissimilarity reduction between mesohabitats, coupled with a prevalence of taxa turnover on taxa loss. These phenomena could be due to a general loss of the more sensitive and specialized taxa, which leads to the homogenization of communities. Nevertheless, the turnover remains greater than nestedness, suggesting the presence of taxa well adapted to the different conditions of mesohabitats and able to manage with the effects of flow reduction.

This work provides significant insights, also into the biomonitoring procedures. Hence, for BRs the Italian legislation limits the range of application of the standard biomonitoring methods to the main channel. This choice, based on our data, could lead to collect samples unrepresentative of the real communities, with a loss of 20% of families, and to obtain incorrect evaluations. In BRs the

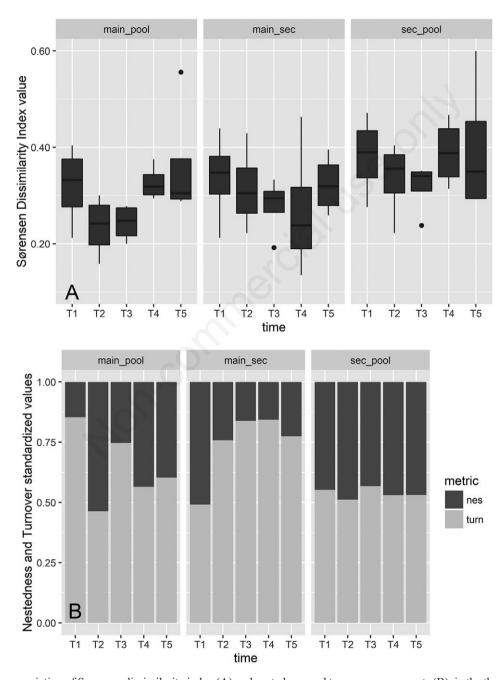


**Fig. 4.** Mesohabitat contribution to the total number of *taxa* for seasonal (A) and summer data (B). Main, main channels; sec, secondary channels; pool, pools.

distribution of benthic macroinvertebrates exhibits high levels of heterogeneity and therefore the ecological status cannot be evaluated considering exclusively the main channels, but it should be assessed considering the river ecosystem as a whole, including marginal water bodies (secondary channels and pools) that are common in these environments.

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**Fig. 5.** Short-term variation of Sørensen dissimilarity index (A) and nestedness and turnover components (B), in the three pairwise confrontations between mesohabitats (main\_pool, main channel-pool; main\_sec, main channel-secondary channel: sec\_pool, secondary channel-pool). Nestedness and turnover values were standardized by dividing by the Sørensen index. T1:T5, sampling campaigns.

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